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Research Article

Prostacyclin (PGI₂) is a key mediator of pulmonary vasodilation in the perinatal period and its synthesis in the pulmonary vasculature increases markedly during late gestation due to enhanced expression of the rate-limiting enzyme cyclooxygenase-1 (COX-1). The hormone estrogen may play a role in COX-1 upregulation since fetal estrogen levels rise dramatically during late gestation and estrogen enhances PGI₂ synthesis in nonpulmonary vascular cells. We therefore studied the direct effects of estrogen on COX-1 expression in ovine fetal pulmonary artery endothelial cells (PAEC). Exposure to estradiol-17beta (E2beta, 10⁽⁻⁾10 to 10⁽⁻⁾6 M) caused a dose-related increase in COX-1 mRNA expression that was evident after 48 h and maximal at 10⁽⁻⁾8 M (fourfold increase). COX-1 mRNA stability was unchanged, suggesting that the upregulation is mediated at the level of transcription. E2beta treatment (10⁽⁻⁾8 M for 48 h) also caused a threefold increase in COX-1 protein expression and a threefold increase in PGI₂ synthesis stimulated by bradykinin, the calcium ionophore A23187, or arachidonic acid. The estrogen receptor (ER) antagonist ICI 182,780 fully reversed the effects of the hormone on COX-1 protein expression and on arachidonic acid-stimulated PGI₂ synthesis, and ER expression was evident in the PAEC by immunoblot analysis. These findings indicate that physiologic levels of estrogen cause upregulation of COX-1 expression and PGI₂ synthesis in fetal PAEC via activation of PAEC ER. This [...]

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Estrogen Upregulates Cyclooxygenase-1 Gene Expression in Ovine Fetal Pulmonary Artery Endothelium

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Abstract

Prostacyclin (PGI₂) is a key mediator of pulmonary vasodilation in the perinatal period and its synthesis in the pulmonary vasculature increases markedly during late gestation due to enhanced expression of the rate-limiting enzyme cyclooxygenase-1 (COX-1). The hormone estrogen may play a role in COX-1 upregulation since fetal estrogen levels rise dramatically during late gestation and estrogen enhances PGI₂ synthesis in nonpulmonary vascular cells. We therefore studied the direct effects of estrogen on COX-1 expression in ovine fetal pulmonary artery endothelial cells (PAEC). Exposure to estradiol-17β (E₂β, 10⁻¹⁰ to 10⁻⁶ M) caused a dose-related increase in COX-1 mRNA expression that was evident after 48 h and maximal at 10⁻⁸ M (fourfold increase). COX-1 mRNA stability was unchanged, suggesting that the upregulation is mediated at the level of transcription. E₂β treatment (10⁻⁸ M for 48 h) also caused a threefold increase in COX-1 protein expression and a threefold increase in PGI₂ synthesis stimulated by bradykinin, the calcium ionophore A23187, or arachidonic acid. The estrogen receptor (ER) antagonist ICI 162,780 fully reversed the effects of the hormone on COX-1 protein expression and on arachidonic acid-stimulated PGI₂ synthesis, and ER expression was evident in the PAEC by immunoblot analysis. These findings indicate that physiologic levels of estrogen cause upregulation of COX-1 expression and PGI₂ synthesis in fetal PAEC via activation of PAEC ER. This process may play a critical role in optimizing the capacity for PGI₂-mediated pulmonary vasodilation at birth, and it may also be involved in estrogen responsiveness in other vascular beds. (*J. Clin. Invest.* 1998; 102:176–183.) Key words: estrogen receptor • immunoblotting • polymerase chain reaction • prostacyclin • pulmonary circulation

Introduction

Prostacyclin (PGI₂)¹ and other vasodilator prostaglandins are important mediators of pulmonary vascular and parenchymal function in the perinatal period. PGI₂ infusion causes pulmo-

nary vasodilation in the fetus and newborn, and the inhibition of endogenous PGI₂ synthesis leads to pulmonary vasoconstriction and marked attenuation of the fall in pulmonary vascular resistance at birth (1, 2). There is also evidence that PGI₂ modulates vascular cell growth in the pulmonary circulation (3). In addition, endogenous prostaglandins are important stimulators of surfactant synthesis and cell differentiation in the developing lung (4). PGI₂ is the primary prostaglandin produced in the developing pulmonary vasculature, where the main site of synthesis is the endothelium (5).

Studies in several species have shown that PGI₂ synthesis in the whole lung increases dramatically during late gestation (6–8). In experiments with lambs, we have previously demonstrated that PGI₂ synthesis in intrapulmonary arteries rises markedly during late fetal life, and that this is due to a developmental increase in the expression of the rate-limiting enzyme for PGI₂ synthesis, cyclooxygenase (COX) (5). In particular, we have shown that there is a maturation-related increase in the expression of the type 1 isoform of COX, or COX-1 (5).

The mechanism underlying the developmental increase in COX-1 abundance in the fetal pulmonary vasculature is unknown. Upregulation during late fetal life suggests that it is due to a factor which has increasing activity in the lung during late gestation. One potential factor is the hormone estrogen, which increases markedly in abundance in fetal plasma during this period due to rising production by the placenta (9–11). Previous studies have shown that prolonged estrogen exposure enhances PGI₂ production in nonpulmonary vascular cells (12, 13). However, it is not known whether estrogen modulates COX-1 expression, or whether the hormone alters PGI₂ synthesis in fetal pulmonary vascular cells.

To better understand the basic mechanisms regulating COX-1 expression in the fetal pulmonary circulation, the present investigation was designed to determine the direct effects of estrogen on COX-1 in fetal pulmonary artery endothelial cells (PAEC). Experiments were performed in early passage, cultured ovine fetal PAEC, which we have used previously in studies of oxygen modulation of COX-1 expression (14). Based on the observation that pulmonary arterial COX-1 expression increases in sheep during late gestation while fetal plasma estrogen levels are rising (5, 9–11), and the finding that prolonged estrogen exposure enhances PGI₂ synthesis in certain nonpulmonary vascular cells (12, 13), we hypothesized that estrogen upregulates COX-1 expression in ovine fetal PAEC. In addition to testing this hypothesis, studies were performed to answer the following questions: (a) Is COX-1 expression modified by estrogen at physiologic concentrations? (b) What is the mechanism(s) underlying the effects of estrogen on COX-1? (c) Are there resulting changes in PAEC PGI₂

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1. Abbreviations used in this paper: COX, cyclooxygenase; E₂β, estradiol-17β; MDH, malate dehydrogenase; PAEC, pulmonary artery endothelial cell; PGI₂, prostacyclin; RT, reverse transcription.

synthesis? And (d) what is the role of estrogen receptors (ER) in this process?

Methods

Cell culture and treatment. PAEC were obtained from mixed breed fetal lambs at 125–135 d gestation, with term being 144 ± 4 d, using methods that we have previously described (14). The PAEC were propagated in RPMI-1640 medium containing 10% iron-supplemented calf serum, 10% lamb serum, 1% L-glutamine, 1% antibiotic-antimycotic mixture, 0.15% nystatin, 0.15% gentamycin, and 0.10% tylosin, in a humidified incubator with 5% CO₂ in air at 37°C. The identity of the cells was confirmed by phenotype (cobblestone appearance and contact inhibition), by immunofluorescence studies with antibody to factor VIII-related antigen, and by examination of acetylated low density lipoprotein uptake. The cells were studied at passage 4–6.

Near-confluent PAEC were placed in phenol red-free, serum-free media for 12 h to remove the effects of the estrogen-like activity of phenol red and serum-derived estrogen. The cells were then placed in phenol red-free media containing 20% charcoal-stripped serum. The charcoal stripping removes estrogen metabolites and other steroid hormones (15). The cells were treated for up to 96 h with either control media or media containing varying concentrations of estradiol-17 β (E₂ β) ranging from 10^{-10} to 10^{-6} M. Culture media was replaced every 48 h and estrogen treatment was repeated every 24 h.

Reverse transcription-polymerase chain reaction (RT-PCR) assay. A semiquantitative RT-PCR assay was established to evaluate COX-1 mRNA abundance in PAEC because the mRNA was not detectable in PAEC by Northern analysis of poly A(+) RNA. Total cellular RNA was obtained from control and estrogen-treated PAEC grown in 75-cm² flasks by a single extraction method with an acid guanidinium thiocyanate-phenol-chloroform mixture (16). RT was performed by methods previously reported using 5 μ g of total RNA (16). Briefly, cDNA synthesis was carried out using 200-U Maloney murine leukemia virus reverse transcriptase, 5 μ M oligo-(dT), 1 mM deoxyribonucleoside triphosphate (dNTP), and 3 mM Mg²⁺ in a volume of 20 μ l. In selected tubes the reverse transcriptase was omitted to control for amplification from contaminating cDNA or genomic DNA. The temperature profile was: (a) annealing at room temperature (25°C) for 5 min, (b) extension at 42°C for 45 min, and (c) termination at 99°C for 5 min.

PCR was performed on the resulting RT product using specific oligonucleotide primers for sheep COX-1 (17). The sequence of the sense primer was 5'-ATGAGTACCGCAAGAGGTTTGG-3' and that of the antisense primer was 5'-ACGTGGAAGGAGACAT-AGG-3'. The PCR reactions contained 1.5 mM Mg²⁺, 1- μ M primers, 200- μ M dNTPs, reaction buffer, and 5 μ l cDNA in a final volume of 50 μ l. To minimize nonspecific amplification, a "hot start" procedure was used in which the PCR tubes were placed in a thermal cycler (480; Perkin-Elmer Cetus Corp., Emeryville, CA) prewarmed to 94°C. After 2 min, each tube was opened sequentially and 2.5 U (in 0.25 μ l) Taq DNA polymerase was added. The PCR temperature profile consisted of 30 cycles of 94°C for 45 s (denaturation), 57°C for 45 s (annealing), and 72°C for 1 min (extension) followed by an additional 5 min final extension at 72°C. The primer location, primer concentration, Mg²⁺ concentration, and annealing temperature were optimized to produce the greatest amount of a single PCR product.

The PCR products were size fractionated by agarose gel electrophoresis. The identity of the PCR products was confirmed, and they were quantitated by transferring the DNA to nylon filters, probing with a ³²P end-labeled internal oligonucleotide primer specific for sheep COX-1, and performing densitometric analysis on the resulting autoradiographs. PCR product identity was also confirmed by direct double-stranded sequencing. To control for the RT step and RNA stability, RT-PCR was also done for the housekeeping gene malate dehydrogenase (MDH) using published oligonucleotide primer se-

quences (18). The PCR temperature profile for MDH was identical to that described above for COX-1.

Preliminary experiments were performed to determine the relationship between the quantity of total RNA subjected to RT-PCR and the amount of PCR product generated. Linear regression analysis showed high correlations between densitometry values for RT-PCR products and the quantity of total RNA used for COX-1 RT-PCR ($r = 0.95-0.99$, $n = 3$ experiments) and for MDH RT-PCR ($r = 0.96-0.99$, $n=3$). We have previously used RT-PCR assays done in this semiquantitative manner in studies of pulmonary endothelial nitric oxide synthase expression; Northern analyses were performed in parallel with RT-PCR assays, and identical results were obtained with the two techniques (16, 19). To evaluate COX-1 mRNA stability, additional RT-PCR experiments were done using cells treated with 25 μ g/ml actinomycin D for varying time periods up to 2 h.

Immunoblot analysis. To quantitate the levels of COX-1 protein expression in control and E₂ β -treated cells, immunoblot analysis was performed using methods that generally followed those we have previously reported (14). PAEC cells were harvested in ice-cold PBS, pelleted, resuspended in 50 mM Tris buffer (pH 7.4) containing 16 mM CHAPS, 100 mM NaCl, 0.5 mM EDTA, 0.02 mM EGTA, 0.4 mM β -mercaptoethanol, 1.6 mM dithiothreitol, and 2 μ g/ml each of soybean trypsin inhibitor, limabean trypsin inhibitor, antipain, and leupeptin, and ultrasonically disrupted (Branson Ultrasonics, Chicago, IL). The protein contents of the preparations were determined, SDS-polyacrylamide gel electrophoresis was performed on equivalent protein samples with 10% acrylamide, and the proteins were electrophoretically transferred to PVDF membranes. The membranes were blocked overnight in buffer containing 137 mM NaCl and 20 mM Tris (pH 7.5) with 0.5% Tween-20 and 5% dried milk, and were incubated with 1:100 COX-1 antiserum (Oxford Biomedical Research, Inc., Oxford, MI) for 2 h at room temperature. After incubation with primary antiserum, the membranes were washed with the 137-mM NaCl buffer with Tween-20 at 0.2% and dried milk at 0.2% and incubated for 1 h with a 1:10,000 dilution of anti-rabbit Ig antibody horseradish peroxidase conjugate raised in donkey. The membranes were washed in the 137-mM NaCl buffer with Tween-20, and the band for COX-1 was visualized by chemiluminescence (ECL Western Blotting Analysis System, Amersham Corp., Arlington Heights, IL) and quantitated by densitometry. Purified COX-1 protein (Cayman Chemical Co., Ann Arbor, MI) was used as a positive control. Similar techniques, including the use of a positive control, were used to evaluate COX-2 protein expression. To determine the role of estrogen receptor (ER) activation in the effects of the hormone on COX protein expression, additional studies were performed in cells treated with the ER antagonist ICI 182,780 (10^{-5} M) (20).

To determine if ER protein is expressed in the ovine fetal PAEC, immunoblot analysis was also performed using 2 μ g/ml of the mouse monoclonal antibody AER 320 directed against amino acids 495–595 in the estrogen-binding domain of human ER α (Neomarkers, Inc., Fremont, CA). Since the ligand-binding domains of the classical ER α isoform and the newly described ER β isoform are highly homologous (21, 22), the antisera may recognize either receptor subtype.

Incubations for prostaglandin synthesis. Control and estrogen-treated PAEC grown in 24-well plates were preincubated for 15 min in a humidified incubator at 37°C with 500 μ l of phenol red-free RPMI-1640 media added per well. The preincubation media was replaced with fresh RPMI-1640 media and 30–90-min incubations were performed. At the end of the incubation, the media was placed into ice-cold tubes containing 100 μ g of acetylsalicylic acid and stored at -20°C until the time of assay for PGI₂. We have previously demonstrated that under these conditions the PGI₂ measured is newly synthesized (14).

In experiments designed to determine the reaction in the PGI₂ synthetic cascade that is modified by estrogen exposure, selected wells were incubated in phenol red-free RPMI-1640 media alone, indicative of basal (nonstimulated) synthesis, and others were treated with agents that activate the synthetic pathway at various steps. Incu-

bations were performed in the presence of bradykinin to assess PGI₂ synthesis stimulated by receptor-mediated mobilization of arachidonic acid from phospholipids (23). Incubations with the calcium ionophore A23187 were performed to evaluate PGI₂ synthesis stimulated by an increase in cytosolic-free calcium, which activates arachidonic acid mobilization by a nonreceptor-mediated process (23). In addition, PGI₂ synthesis stimulated by exogenous arachidonic acid was also measured to determine if estrogen-mediated changes in synthesis are related to changes in COX activity (23). Preliminary experiments revealed that maximal stimulation of PGI₂ synthesis is obtained with bradykinin, A23187, and arachidonic acid at 10⁻⁵ M. Therefore, this concentration was used in all ensuing experiments. Preliminary experiments also showed that basal and stimulated PGI₂ synthesis rose linearly with time between 30–90 min of cell incubation. As a result, 60-min incubations were used in all ensuing studies. Additional experiments were performed comparing arachidonic acid-stimulated PGI₂ synthesis in control PAEC, estrogen-treated PAEC (10⁻⁸ M for 48 h), and PAEC treated with both estrogen and ICI 182,780 (10⁻⁵ M). ICI 182,780 alone had no effect on PGI₂ synthesis in control cells. In all experiments, *n* = 4–6 for each determination, and findings were replicated in three independent experiments.

Prostaglandin assays. Samples of incubation media were assayed for the stable metabolite of PGI₂, 6-keto-PGF_{1α}, by radioimmunoassay as previously reported (14). Briefly, the assay procedure used duplicate aliquots of standard (0–1,000 pg) or samples placed into a solution containing 0.1 M phosphate-buffered saline plus 0.1% polyvinylpyrrolidone (1:1). Antiserum (0.1 ml; 1:4,000 titer) and [³H]6-keto-PGF_{1α} (0.1 ml; 12,000 dpm) were added, and the tubes were incubated at 4°C for 12–18 h. Bound and free ligand were separated with dextran-coated charcoal, and bound ligand was counted by liquid scintillation spectrometry. The unknown quantities of PGI₂ were determined from the standard curves generated.

Statistical analysis. Analysis of variance with Newman-Keuls post-hoc testing was used to compare mean values between more than two groups. Nonparametric analysis of variance was used when indicated. Single comparisons between two groups were performed with non-paired Student's *t* tests. Significance was accepted at the 0.05 level of probability. All results are expressed as mean ± SEM.

Results

COX-1 mRNA expression. The effect of varying concentrations of E₂β on COX-1 mRNA expression in fetal PAEC is shown in Fig. 1. The cells were treated with control media or media with 10⁻¹⁰ to 10⁻⁶ M E₂β for 48 h. Single PCR products were obtained for COX-1 at the expected size of 355 bp (Fig. 1 A). The representative Southern blot reveals a dose-dependent increase in COX-1 mRNA abundance as determined by RT-PCR in response to E₂β treatment. PCR was also performed for MDH to control for the RT step, yielding a single PCR product at the expected size of 369 bp. There was no change in MDH mRNA abundance with E₂β treatment. Quantitative densitometry for three independent experiments confirmed these results (Fig. 1 B). There was a concentration-related increase in steady-state COX-1 mRNA levels, with a maximal increase of 4.4-fold at 10⁻⁸ M E₂β.

The time course for E₂β modulation of COX-1 mRNA expression in fetal PAEC is shown in Fig. 2. In the representative Southern blot (Fig. 2 A), COX-1 mRNA abundance as determined by RT-PCR was increased following both 48 and 96 h of treatment with 10⁻⁸ M E₂β compared with controls. E₂β had no effect on MDH mRNA abundance. Quantitative densitometry for three independent experiments confirmed these results, showing a 4.4-fold increase in COX-1 expression at 48 h and a persistent increase of 4.2-fold at 96 h (Fig. 2 B). Before

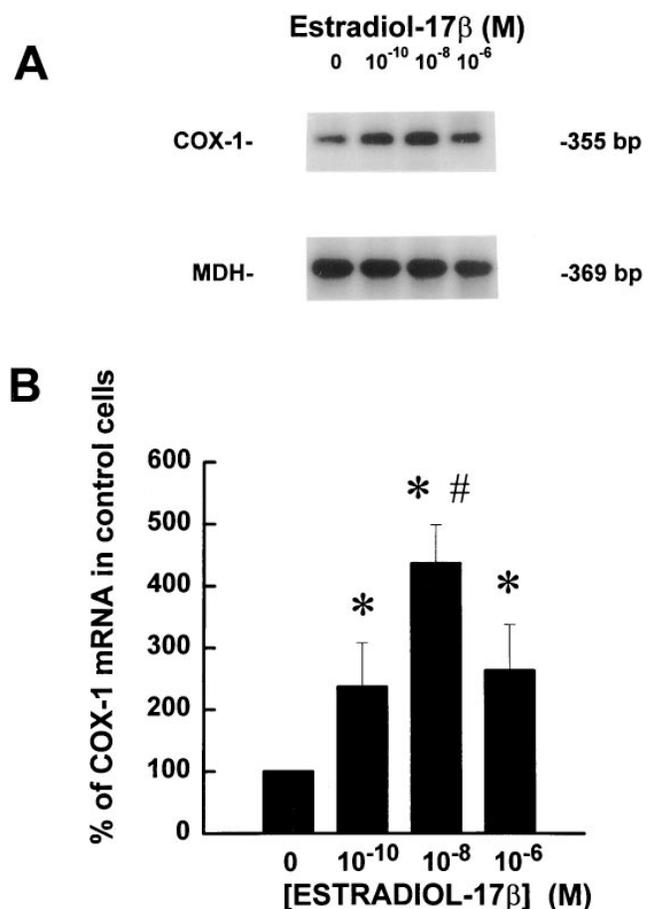


Figure 1. (A) Southern analyses of RT-PCR products for COX-1 (top) and malate dehydrogenase (MDH, bottom) in fetal PAEC exposed to varying concentrations of E₂β for 48 h. Band sizes were: COX-1, 355 bp; MDH, 369 bp. (B) Summary data for three independent experiments. COX-1 densitometry values corrected for MDH are expressed as a percentage in control cells (mean ± SEM). **P* < 0.05 vs control, #*P* < 0.05 vs 10⁻¹⁰ M E₂β.

48 h there was no significant change in COX-1 mRNA abundance with exposure to E₂β (data not shown).

The effects of E₂β on COX-1 mRNA stability are depicted in Fig. 3. RT-PCR was performed on cells treated with control media or media with 10⁻⁸ M E₂β for 48 h, followed by 25 μg/ml actinomycin D for varying durations up to 2 h. The representative study shown reveals no difference in COX-1 mRNA degradation in control and E₂β-treated cells. In four independent experiments COX-1 mRNA half-life was similar in control and E₂β-treated cells, being 2.0 ± 0.2 and 2.1 ± 0.5 h (mean ± SEM), respectively.

COX protein expression. The effects of E₂β on COX-1 protein abundance are shown in Fig. 4. In the representative immunoblot shown (Fig. 4 A), COX-1 was increased in cells treated with 10⁻⁸ M E₂β for 48 h, and this effect was completely reversed by ER antagonism with ICI 182,780. The cumulative findings of three independent experiments confirmed these results (Fig. 4 B), revealing a 2.7-fold increase in COX-1 protein with E₂β treatment that was fully negated by ICI 182,780 (10⁻⁵ M). COX-2 protein was not detected in any study groups (data not shown).

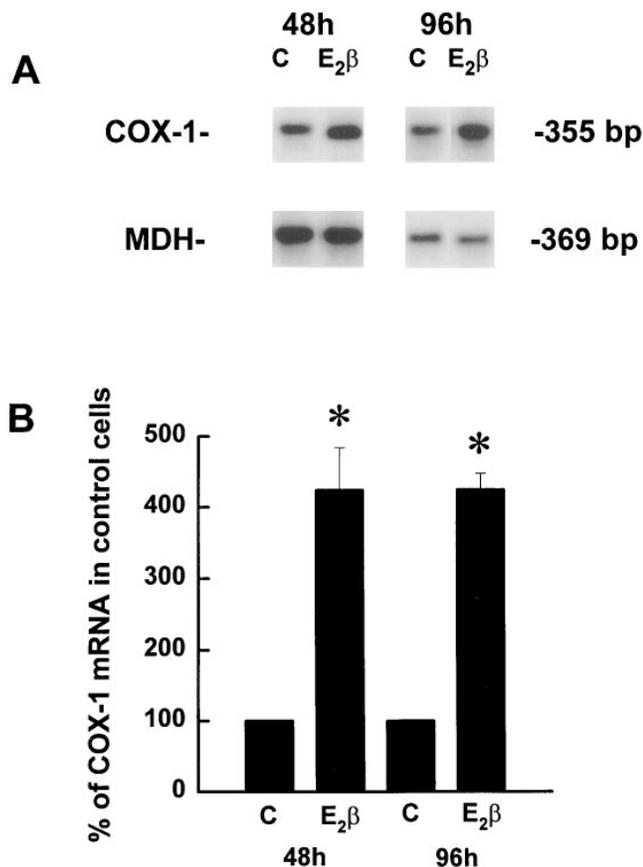


Figure 2. Time course of the effect of estrogen on COX-1 mRNA abundance in fetal PAEC. Cells were exposed to control media (C) or media containing 10^{-8} M E₂β. (A) Southern analysis of RT-PCR products for COX-1 (top) and MDH (bottom) in control and E₂β-treated cells. Band sizes were: COX-1, 355 bp; MDH, 369 bp. (B) Summary data for three independent experiments. COX-1 densitometry values corrected for MDH are expressed as a percentage in control cells (mean ± SEM). * $P < 0.05$ vs control.

PGI₂ synthesis. The effect of E₂β on basal and bradykinin-stimulated PGI₂ synthesis in fetal PAEC is shown in Fig. 5 A. PGI₂ synthesis was measured in control and E₂β-treated cells (10^{-8} M for 48 h) during 60-min incubations. In control cells, basal PGI₂ synthesis was 10.6 ± 1.4 pg/well, and synthesis was stimulated 2.4-fold by bradykinin. Prolonged E₂β treatment caused a 64% increase in basal PGI₂ synthesis, and a 2.9-fold increase in bradykinin-stimulated synthesis. The effect of E₂β on A23187- and arachidonic acid-stimulated PGI₂ synthesis is depicted in Fig. 5 B. In parallel with the estrogen-mediated increases in basal and bradykinin-stimulated production, synthesis with A23187 was increased by 2.7-fold and synthesis with arachidonic acid was increased by 2.5-fold after exposure to 10^{-8} M E₂β for 48 h.

Role of estrogen receptors. To further determine the role of ER in the response to E₂β, arachidonic acid-stimulated PGI₂ synthesis was compared in control cells, cells treated with 10^{-8} M E₂β for 48 h, and cells treated with E₂β plus the ER antagonist ICI 182,780 (10^{-5} M). Similar to the findings shown in Fig. 5 B, estrogen caused a 2.3-fold increase in arachidonic

acid-stimulated PGI₂ synthesis compared to control conditions (Fig. 6). This effect was completely reversed by ER antagonism. To determine if ER protein is expressed in the ovine fetal PAEC, immunoblot analysis was performed. Immunoblots revealed signal for ER protein at 67 kD (Fig. 6, inset). Similar observations were made in three independent experiments.

Discussion

In the present study, we have evaluated the effects of E₂β on COX-1 expression in cultured ovine fetal PAEC, enabling us to examine the direct effects of the hormone on the fetal pulmonary endothelium without potential secondary effects due to changes in cardiac or systemic vascular function (24, 25). We have demonstrated that E₂β causes a marked upregulation in COX-1 mRNA expression in the ovine fetal pulmonary artery endothelium. To our knowledge, this is the first demonstration of a direct effect of estrogen on COX-1 expression in any cell type.

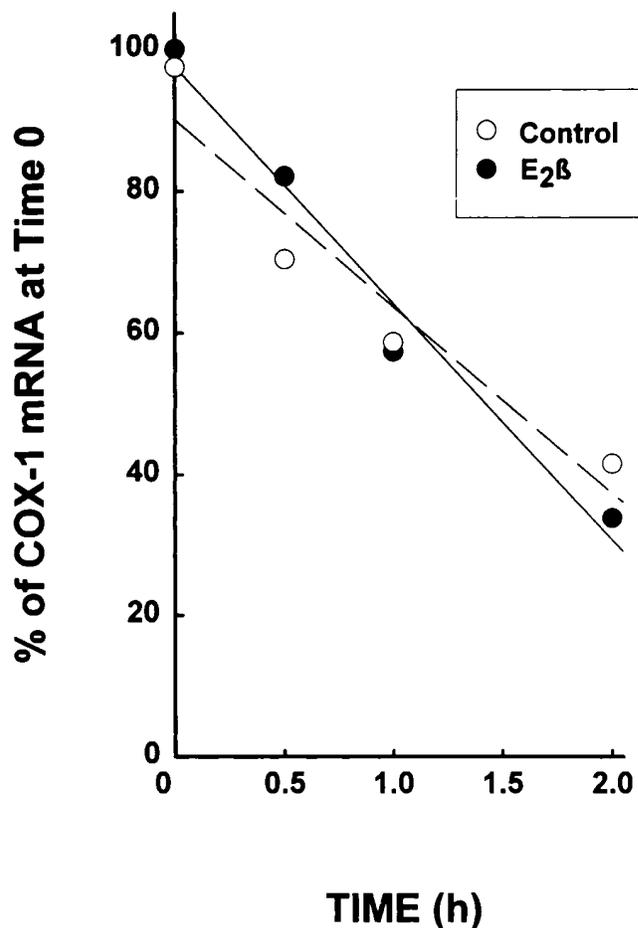


Figure 3. Effect of estrogen on COX-1 mRNA stability. Cells were treated with control media (open symbol, dashed line) or media containing 10^{-8} M E₂β for 48 h (closed symbol, solid line), followed by 25 μg/ml actinomycin D for varying time periods up to 2 h. COX-1 mRNA abundance, assessed by RT-PCR, is expressed as a percent at time 0 of actinomycin D. Linear regression analysis yielded $r = -0.95$ for control cells and $r = -0.99$ for E₂β-treated cells. Similar findings were obtained in four independent experiments.

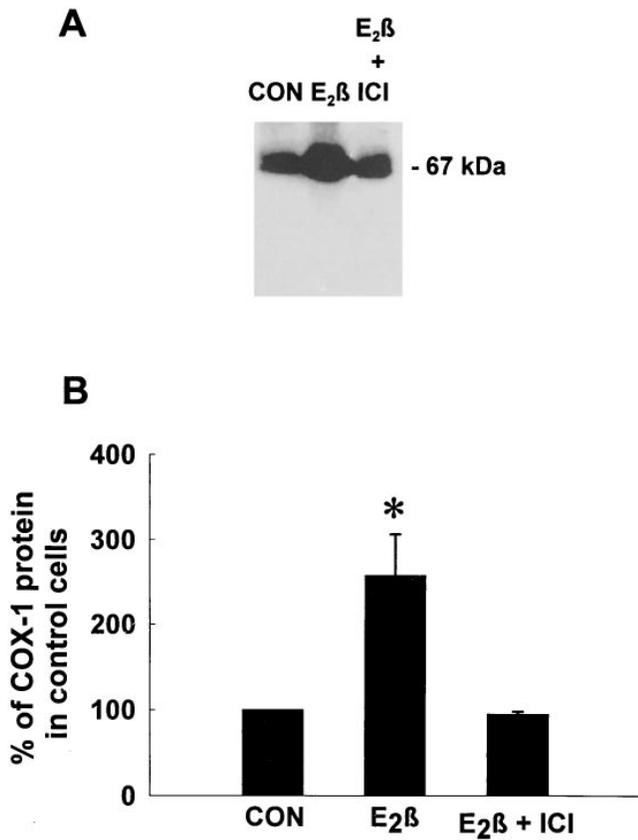


Figure 4. Effect of estrogen on COX-1 protein abundance evaluated by immunoblot analysis. Cells were exposed to control media (CON), media containing 10^{-8} M E₂β, or media containing E₂β plus ICI 182,780 (10^{-5} M) for 48 h. COX-1 protein was detected at 70 kD. The immunoblots shown are representative of three independent experiments. Summary data for quantitative densitometry for the three experiments is given in *B*. Mean \pm SEM values are depicted for protein abundance expressed as a percentage in control cells. * $P < 0.05$ vs control.

We have also shown that the effect of estrogen on COX-1 mRNA occurs at a threshold concentration of 10^{-10} M, and that there is maximal upregulation by more than fourfold at 10^{-8} M. This indicates that the effect of the hormone occurs at levels that are achieved in the fetal plasma in a variety of species during late gestation (9–11). For example, plasma levels of unconjugated estradiol in fetal sheep increase fivefold from 80 to 140 d gestation (term = 144 d), reaching concentrations in the range of 10^{-9} M (9). In addition, we have shown that the effect of E₂β on COX-1 mRNA is evident by 48 h and persists for at least 96 h. This time course is consistent with prior studies examining the direct effect of chronic estrogen exposure on PGI₂ synthesis in nonpulmonary vascular cells. In both piglet aortic endothelium and rat aortic smooth muscle cells, PGI₂ synthesis is enhanced after 2–3 d of exposure to 10^{-9} M E₂β (12, 13). This suggests that COX-1 expression is regulated by prolonged exposure to physiologic concentrations of the hormone in a variety of vascular cells in addition to the fetal pulmonary endothelium. Furthermore, we have demonstrated in the fetal PAEC that COX-1 mRNA stability is not altered by prolonged E₂β exposure, suggesting that the chronic effects of

the hormone on COX-1 are mediated at the level of gene transcription. These findings contrast with those of Myers et al., who demonstrated enhanced PGI₂ synthesis in rat aortic endothelium after only 60 min exposure to E₂β that was not due to changes in COX or PGI₂ synthase abundance (26). Cumulatively, these observations suggest that there may be both chronic and acute effects of the hormone on the PGI₂ synthetic cascade that are related to different mechanisms. In addition to the experiments examining COX-1 mRNA abundance, we performed studies evaluating the effect of E₂β on COX-1 protein expression. After exposure to 10^{-8} M E₂β for 48 h, COX-1 protein was increased by 2.7-fold, paralleling the rise in COX-1 mRNA. In contrast to the findings for COX-1, COX-2 protein was not detected in either control or E₂β-treated cells, revealing that the effects of E₂β on COX-1 are unique to that isoform.

Along with the studies of COX-1 expression, we determined whether there are resulting changes in endothelial cell PGI₂ synthesis. After treatment with 10^{-8} M E₂β for 48 h,

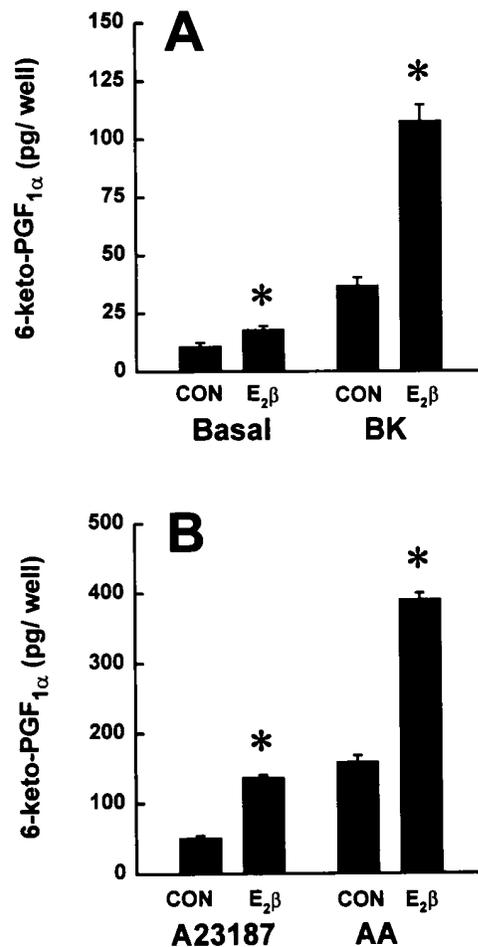


Figure 5. Effect of estrogen on basal and stimulated PGI₂ synthesis in fetal PAEC. Cells were exposed to control media (CON) or media containing 10^{-8} M E₂β for 48 h and then incubated for 60 min in the absence of exogenous stimulation (Basal) or in the presence of bradykinin (BK) (A), or in the presence of A23187 or arachidonic acid (AA) (B), each at 10^{-5} M. PGI₂ (6-keto-PGF_{1α}) synthesis was measured by RIA. Values are mean \pm SEM, $n = 4-6$. * $P < 0.05$ vs control.

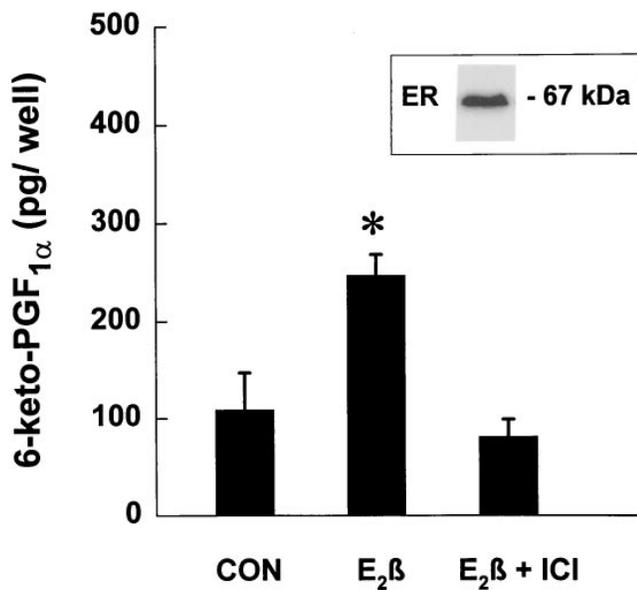


Figure 6. Role of estrogen receptors in the effect of E₂β on COX activity. Cells were exposed to control media (CON), media containing 10⁻⁸ M E₂β, or media containing E₂β plus ICI 182,780 (10⁻⁵ M) for 48 h and then incubated for 60 min in the presence of 10⁻⁵ M arachidonic acid. PGI₂ (6-keto-PGF_{1α}) synthesis was measured by RIA. Values are mean ± SEM, n = 4–6. *P < 0.05 vs control. (Inset) Immunoblot analysis for ER protein in ovine fetal PAEC. Signal for ER protein was evident at 67 kDa. Results are representative of three independent experiments.

basal PGI₂ production was increased by 64%, and synthesis stimulated with bradykinin and A23187 were increased 2.9- and 2.7-fold, respectively. In a similar manner, synthesis in the presence of excess exogenous arachidonic acid was augmented 2.5-fold after estrogen treatment, indicating that the effect of estrogen on PGI₂ production is not mediated by changes in the capacity for arachidonic acid mobilization. Instead, these results reflect an enhancement in COX enzymatic activity, and this is in agreement with the observation that estrogen augments COX-1 mRNA and protein expression.

In the present investigation, we also determined whether ER are involved in the effect of the hormone on COX-1 protein expression and PGI₂ synthesis. The ER antagonist ICI 182,780 fully inhibited the estrogen-mediated increases in both COX-1 protein expression and arachidonic acid-stimulated PGI₂ synthesis indicative of the level of COX activity, revealing that ER activation is necessary for these effects. In addition, ER protein expression was demonstrated by immunoblot analysis. These findings indicate that functional ER are expressed in the fetal PAEC, and that they play a key role in the regulation of COX-1. However, it is not yet clear which ER subtype is involved. The ERα and ERβ subtypes, which are known to function as transcription factors, are highly homologous, particularly in the DNA-binding domain (95–96%) and in the COOH-terminal hormone-binding domain (55–58%) (21, 22). In addition, both ERα and ERβ are inhibited by ICI compounds (20, 21), and the immunoblot analysis may recognize either ERα or ERβ. Thus, the observed effects of estrogen on PAEC COX-1 expression may be mediated by either ERα, ERβ, or both receptor subtypes, or by a yet unknown

ER subtype. Studies of ER subtype in adult rat lung reveal the expression of both ERα and ERβ, with the latter subtype predominating (27), but the pulmonary cell specificity of ER subtype expression is yet to be determined. Further studies are now warranted to distinguish the role of ERα and ERβ in COX-1 regulation in the pulmonary endothelium.

The present studies suggest that the principal effects of prolonged ER activation on COX-1 expression in the fetal PAEC occur at the level of gene transcription. Although the 5' flanking region of COX-1 does not contain a canonical TATA box typical of transcriptionally regulated genes, there are putative transcription factor-binding sites that may be highly relevant to estrogen-mediated regulation of COX-1 expression. There are four potential polyomavirus enhancer activator 3, or PEA3, promoter elements (28, 29). PEA3, which belongs to the Ets family of transcription factors, is principally upregulated by the MAP kinase signal transduction pathway (30). Since estrogen is a potent activator of the MAP kinase pathway in a variety of cell types (31, 32), it is plausible that the hormone may upregulate COX-1 in the PAEC through MAP kinase and PEA3. In addition, the 5' flanking region of COX-1 contains four copies of the estrogen response element (ERE) half-palindromic motif, GGTC A, or its reverse complement, within 2,100 bp of the translation start codon (28, 29). Many estrogen-regulated promoters do not contain perfect palindromic EREs, but include one or more GGTC A motifs, which are sometimes widely spaced (33–35). As such, the four ERE half-palindromes should also be considered as potential regulatory elements involved in the modulation of COX-1 expression by the hormone. An investigation of the transcriptional transactivation of COX-1 by estrogen is now indicated to further elucidate the underlying mechanisms.

A degree of caution may be warranted in the direct extrapolation of the present findings in the cultured cells to processes in the intact fetal lung. However, the use of the cultured cells allows us to evaluate the direct effects of single factors such as estrogen on PAEC phenotype, thereby avoiding potential secondary changes resulting from known cardiac and systemic effects of the hormone (24, 25). In addition, we have previously demonstrated in parallel studies of intrapulmonary arterial segments and early passage PAEC from newborn lambs that the effects of oxygen on COX-1 expression are conserved in culture (14). Furthermore, more recent studies of COX-1 expression in ovine fetal and newborn intrapulmonary arteries and early passage PAEC have shown that the developmental increase in COX-1 expression is also conserved in the cultured cells (Shaul, P.W., and T.S. Brannon, manuscript submitted for publication). Thus, the ovine fetal PAEC provide an excellent model for the study of the regulation of COX-1 gene expression, and the observed effect of estrogen on COX-1 expression and PGI₂ synthesis in the cultured cells may indeed reflect events occurring in the intact fetal lung.

Keeping these potential limitations in mind, there are important physiological implications of estrogen-mediated regulation of pulmonary endothelial COX-1 expression in both the adult and the fetus. In an adult rat model of chronic hypoxic pulmonary hypertension, Rabinovitch et al. found that the disease is much milder in female versus male animals (36). The present observations suggest that this may be due to enhanced COX-1 gene expression related to the effects of estrogen on the pulmonary endothelium in the adult females. In the fetus, estrogen-mediated effects on COX-1 expression may at least

partially explain the developmental increase in pulmonary COX-1 expression that occurs during late gestation, at a time when fetal plasma estrogen levels are increasing dramatically due to enhanced placental production of the hormone (5, 9). We have now demonstrated that physiologic concentrations of E₂ upregulate COX-1 expression in the fetal pulmonary artery endothelium. This process may be critically involved in optimizing the capacity for PGI₂-mediated pulmonary vasodilation at birth.

There are also potential pathophysiologic implications of the present findings, particularly in the fetal lung. In a model of intrauterine infection induced with group B streptococcus in fetal rhesus monkeys, the normal rise in plasma estrogen which occurs before birth is absent (37). In addition, estrogen levels are greatly reduced in the cord blood of postmature human newborns (38). These observations suggest that in pregnancies complicated by placental dysfunction such as that associated with intrauterine infection or postmaturity, diminished estrogen synthesis by the placenta may lead to diminished fetal pulmonary COX-1 expression and impaired pulmonary PGI₂ synthesis, thereby contributing to the pathogenesis of persistent pulmonary hypertension of the newborn.

The observed hormonal regulation of COX-1 in the PAEC also has important implications on estrogen-mediated effects in other vascular beds. Studies have shown that PGI₂ synthesis increases dramatically in the uterine circulation during pregnancy and that this is not due to increased availability of the substrate arachidonic acid (39). This may instead be related to effects of estrogen on COX-1 gene expression since Moonen et al. determined that there is significantly greater COX enzyme in pregnant versus nonpregnant human myometrium (40). In the coronary circulation, the risk of vascular disease in premenopausal women is very low compared to men, it increases markedly after menopause, and it diminishes again with estrogen replacement therapy after menopause (41, 42). These findings may also be related to estrogen-mediated effects on prostaglandin production because PGI₂ synthesis is significantly decreased in isolated arteries from postmenopausal versus premenopausal women (43), and PGI₂ is critically important as a vasodilator and an inhibitor of thrombosis and platelet aggregation (44). Thus, estrogen-mediated upregulation of COX-1 may play a major role in the function of not only the developing pulmonary circulation, but also the uterine and coronary circulation. Further studies of estrogen-mediated regulation of COX-1 gene expression in fetal PAEC will continue to advance our knowledge of the role of this hormone in the pulmonary circulation and in other estrogen-responsive vascular beds.

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