The prostaglandin F synthase activity of the human aldose reductase AKR1B1 brings new lenses to look at pathologic conditions

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PROSTAGLANDINS ARE IMPORTANT REGULATORS OF FEMALE REPRODUCTIVE FUNCTIONS TO WHICH ALDOSREductase AKR1B1 (Bresson et al., 2011) were highly functional PGFS. Prostaglandins (PGs) are notorious mediators of pain and inflammation and also associated with proliferation of cancer cells. While the same mediators are recognized as primary regulators of female reproductive function (ovulation, uterine receptivity, implantation, and parturition) they also contribute to occurrence of endometrial carcinomas, menorrhagia, dysmenorrhea, endometriosis, and premature labor. Among the different PGs, PGE2 and PGF2α are the main prostanoids produced in the human endometrium (Smith and Kelly, 1988; Sales and Jabbour, 2003a). PGs are synthesized from arachidonic acid (AA) and converted to PGG2 and PGH2 by PGH synthases (PGHS), also known as cyclooxygenases (COX). There are two documented isoforms of PGHS in human, the constitutive COX-1 and the inducible COX-2 encoded by two distinct genes (Smith et al., 1996). PGH2 produced by COXs is the common precursor of all PGs generated by specific terminal synthases such as PGE synthase for PGE2α and PGE synthase for PGE2. The human terminal synthases responsible for more than four decades (Srivastava et al., 2005). AKR1B1 enzyme which has been associated with complications of diabetes for more than four decades (Srivastava et al., 2005). AKR1B1 is considered as the rate limiting enzyme of the polyol pathway responsible for the conversion of glucose into sorbitol. The association of AKR1B1 with cardiovascular risks, neuropathies, and nephropathies characteristic of diabetes is receiving increased acceptance (Alexiou et al., 2009). However, the contribution of the polyol pathway to the same complications is less convincing (Del Corso et al., 2008). Indeed, while many factors including inflammation induce increased expression and activity of AKR1B1, this enzyme has very little affinity for glucose at physiological concentrations and a detoxification action involving reduction of lipid peroxides has been proposed (Srivastava et al., 2005). While searching for the enzyme responsible for the production of PGF2α in the endometrium, we made the serendipitous and unexpected discovery that AKR1B5 (Madore et al., 2003) and later the human aldose reductase AKR1B1 (Bresson et al., 2011) were highly functional PGFS. Prostaglandins (PGs) are notorious mediators of pain and inflammation and also associated with proliferation of cancer cells. While the same mediators are recognized as primary regulators of female reproductive function (ovulation, uterine receptivity, implantation, and parturition) they also contribute to occurrence of endometrial carcinomas, menorrhagia, dysmenorrhea, endometriosis, and premature labor. 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While the same mediators are recognized as primary regulators of female reproductive function (ovulation, uterine receptivity, implantation, and parturition) they also contribute to occurrence of endometrial carcinomas, menorrhagia, dysmenorrhea, endometriosis, and premature labor. Among the different PGs, PGE2 and PGF2α are the main prostanoids produced in the human endometrium (Smith and Kelly, 1988; Sales and Jabbour, 2003a). PGs are synthesized from arachidonic acid (AA) and converted to PGG2 and PGH2 by PGH synthases (PGHS), also known as cyclooxygenases (COX). There are two documented isoforms of PGHS in human, the constitutive COX-1 and the inducible COX-2 encoded by two distinct genes (Smith et al., 1996). PGH2 produced by COXs is the common precursor of all PGs generated by specific terminal synthases such as PGE synthase for PGE2α and PGE synthase for PGE2. The human terminal synthases responsible
for the biosynthesis of PGE2 (Park et al., 2006), PGD2 (Zhou et al., 2010), and PGJ2 (Wu and Liou, 2005) are well characterized, but only little is known for the enzyme responsible for PGF2α production. Before our identification of the PGF synthase activity of aldo-ketoreductases (AKR) of the 1B family (Madore et al., 2003; Bresson et al., 2011) only one AKR of the 1C family, AKR1C3 was recognized as a functional PGF synthase in the human (Suzuki-Yamamoto et al., 1999).

PGF2α is a biologically active prostanoïd belonging to the eicosanoid family of bioactive lipids (Narumiya and FitzGerald, 2001). Its biosynthesis occurs via different pathways involving reduction of PGH2 by a 9,11-endoperoxide reductase (Watanabe, 2002). Several PGFS have been identified in animals (Madore et al., 2003), but until recently, AKR1C3 was the only isoform currently identified in human (Komoto et al., 2006). In the bovine endometrium, using a cell-free system, we have demonstrated a strong PGFS activity of purified recombinant AKR1B5 recently renamed as bos taurus AKR1B1 (Gene ID: 317748) as well as circumstantial association between its pattern of expression and PGF2α production. This represents a new putative function for this enzyme previously known for its 20α-HSD and glucose metabolism activities (Madore et al., 2003). The human and bovine AKR1B1 (EC:1.1.1.21) both belong to the AKR1B family and share 86% identity or homology. The human AKR1B1 (Gene ID: 231) also known as the aldose reductase or ALDR1 belongs to the AKR superfamily composed of 140 members divided into 15 families (Jin and Penning, 2007). AKR1B1 (EC:1.1.1.21) is primarily known as the rate limiting enzyme for conversion of glucose to sorbitol in the polyol pathway, but recent studies revealed its ability to convert a wide array of substrates including aldehydes generated during lipid peroxidation and their glutathione (GSH) conjugates, phospholipids, atherogenic lipids, and steroids (Srivastava et al., 2005).

After identifying the bovine AKR1B1 as a potential PGFS (Madore et al., 2003), we have shown circumstantial association with PGF2α production in human endometrial cell lines (Chapdelaine et al., 2006) and in decidualized stromal cells (Kang et al., 2006). Later, the PGFS activity of mouse AKR1B3 and AKR1B7 as well as the human AKR1B1 were tested and confirmed using purified recombinant proteins in a cell-free system in vitro (Kabututu et al., 2009). Using a similar approach complemented with gain and loss of function in endometrial cell lines, we have demonstrated that the AKR1B1 was a functional PGFs in the human endometrium (Bresson et al., 2011). In the present study, we have investigated further the regulation of PGF2α production by different stimulators and inhibitors as well as the promoter cis-elements responsible for IL-1β response in human endometrial cell lines. Finally, we have studied the association between AKR1B1 and PGF2α production in representative tissues where PGF2α was found to exert physiological action in vivo. We have evaluated the potential of different AKR1B1 inhibitors (ARI) to alter PGs production in living cells and in fresh endometrial tissues under normal and high glucose conditions.

**MATERIALS AND METHODS**

**REAGENTS**

RPMI 1640 culture medium without phenol, Superscript II reverse transcriptase, TRIzol, lipofectamine 2000, and pCR3.1 vectors were purchased from Invitrogen (Life technologies, ON, Canada). T AQ DNA polymerase and buffer used for polymerase chain reaction were from NEB (New England Biolabs, ON, Canada). Ribomax polymerase kit for siRNAs was purchased from Promega (Madison, WI, USA). Qiagroup quick gel extraction kit and TA cloning pDrive vector were from Qiagen (Mississauga, ON, Canada). All oligonucleotide primers were chemically synthesized by Integrated DNA Technologies (IDT) (Coralville, IA, USA). [α-32P]dCTP radioactivity was bought from PerkinElmer Life Sciences (Markham, ON, Canada). Bright Star-Plus nylon membrane and UltraHyb solution were purchased from Ambion Inc. (Austin, TX, USA). The rabbit COXs antibodies were kindly provided by Dr S Kargman (Merck, QC, Canada), rabbit anti-Phospho-cPLA2 (Ser505) was from Santa Cruz biotechnology (Santa Cruz, CA, USA) and AKR1C3 was from Abcam Inc. (Cambridge, MA, USA). Goat anti-rabbit horse-radish peroxidase-conjugated IgG was bought from Jackson ImmunoResearch Laboratories (West Grove, PA, USA). The Western Lightning™Chemiluminescence Reagent Plus was purchased from PerkinElmer (Woodbridge, ON, Canada). Arachidonic acid (AA), and AL-8810 were from Cayman Chemicals (Ann Arbor, MI, USA) and recombinant human IL-1β was purchased from Research and Diagnostic Systems (Minneapolis, MN, USA). Aldose reductase inhibitors (ARIs), [ponalrestat (Statil), Alrestatin, and EBPC] were from Tocris Bioscience (Ellisville, MO, USA).

![FIGURE 1 | Prostaglandin F2α and E2 biosynthetic pathways. In order to generate PGF2α, AKR1B1 must team up with other constitutive and inducible enzymes involved in the production of PGH2, the common precursor of all PGs. Among those, we have found that cPLA2 liberating AA from membrane phospholipids, COX-2, AKR1B1, and mPGES-1 are inducible by the cytokine IL-1β. AKR1B1 apparently works with both COX-1 and COX-2 producing PGH2 to generate PGF2α whereas mPGES-1 preferentially associates with COX-2 to generate PGE2. Both epithelial and stromal cells exhibit PGF2α responsiveness and stimulation or inhibition of the FP receptor alters the production of both PGE2 and PGF2α through action at several steps of the PG biosynthesis cascade. We also present in this figure the PG transporters PGT also known as solute carrier SLCOA1 and MRP4 the multidrug resistance protein 4 or ABCC4.](image-url)
CULTURED CELL MODELS

Human endometrial stromal (HIESC-2) and epithelial (HIEEC-22) cell lines were immortalized using SV40 large T antigen (Chapdelaine et al., 2006) and shown to exhibit most characteristics of freshly isolated endometrial cells (Kang et al., 2004, 2006). Bovine endometrial epithelial cells were immortalized using a similar protocol and also presented most characteristics of freshly isolated cells (Krishnaswamy et al., 2009, 2010; Lacroix-Pepin et al., 2011). Human prostate smooth muscle cells, human artery smooth muscle cells, human umbilical vein endothelial cells (Huvec), and human cardiomyocytes cells were from ScienCell Research Laboratories (Carlsbad, CA, USA).

NORTHERN BLOT

Northern blot analysis was performed as follows: total RNA (∼20 μg) was extracted with Trizol (Invitrogen, Life technologies, ON, Canada) from cultured endometrial cells, loaded on a 1.2% formaldehyde-agarose gel and electrophoresed at 100 V in 1× MOPS buffer. After electrophoresis, RNA was transferred overnight onto a nylon membrane in 10× saline-sodium citrate (SSC). The AKR1B1, mPGES-1, COX-1, and COX-2 cDNA probes were produced by digestion with restriction enzymes of different plasmids containing the corresponding cDNAs previously generated in our laboratory, each releasing a ∼500 bp fragment. The resulting probes were labeled with [α-32P(dCTP (3000 Ci/mmol) using the Ready-To-Go DNA labeling Kit (GE Healthcare Life Science, QC, Canada). Prehybridization (2–4 h) and hybridization (overnight) were done at 45°C using UltraHyb solution (Ambion Inc., Invitrogen, Life technologies, ON, Canada). The blots were then washed at 65°C twice for 15 min in 0.5× SSC and exposed for 24–48 h at −80°C on BioMax films PerkinElmer (Woodbridge, ON, Canada) to quantify the hybridization signal intensity. Finally, 18S ribosomal RNA stained with ethidium bromide was used to confirm uniform loading of RNA samples and a 32P-labeled oligonucleotide specific to 18S ribosomal RNA was used as an internal standard for Northern blot analysis.

WESTERN BLOT

Western blot analysis was performed as we described (Chapdelaine et al., 2006). Briefly, total proteins (∼20 μg) extracted from culture cells were loaded in each lane and electrophoresis done on 10% SDS-PAGE followed by electrotransfer onto nitrocellulose membrane (Bio-Rad laboratories, Mississauga, ON, Canada). The
FIGURE 3 | Time course of gene expression and PGs production in human endometrial stromal and epithelial cells following stimulation with IL-1β. Human endometrial stromal (HIESC-2) and epithelial (HIEEC-2) cells were grown to confluence and treated with IL-1β for different periods up to 24 h. Northern Blots analysis (A, B) was used for detection of AKR1B1, COX-1/-2, mPGES-1 mRNAs expression. The production of PGs was evaluated as a function of time in stromal (C) and epithelial (D) endometrial cells where stimulation of PGF2α precedes that of PGE2 (P < 0.05).

primary antibodies used for the present study were the rabbit AKR1B1 (dilution 1/1000), COX-1 or COX-2 (dilution 1/3000), cPLA2 (1/1000) and goat AKR1C3 (1/500) anti-serums. As an internal control, the β-actin monoclonal antibody (1/5000, Sigma, Mississauga, ON, Canada) was used. The goat anti-rabbit IgG conjugated with horse radish peroxidase (HRP), rabbit anti-goat IgG HRP or goat anti-mouse IgG HRP were used as secondary antibodies. The chemiluminescence was analyzed with autoradiography films at optimal times of exposure following treatment of the membranes with the Western Lightning™Chemiluminescence Reagent Plus (PerkinElmer Canada).

ANALYSIS OF AKR1B1 PROMOTER ACTIVITY
A nested PCR strategy was developed to generate a long ~4.5 kb promoter to study the regulation of the AKR1B1 gene (AF032455). The promoter PCR fragments were amplified with PFU turbo DNA polymerase (Stratagene, Agilent technologies, USA) and cloned directly by fusion (Clontech, TAKARA Biocompany) in a pGL3 basic vector (Promega) containing the firefly luciferase gene. Constructs of AKR1B1 promoter of seven different lengths were generated using restriction enzymes targeting unique sequences in the long promoter. A short portion of the AKR1B1 promoter corresponding to the multiple stress response region (MSRR) −1710 to −1064 to the translation initiation ATG position was amplified by PCR, cloned in a TA cloning vector (pDrive, Qiagen) for sequencing and linked in a pGL3 vector (Promega) to the minimal SV40 promoter. From this, additional constructs were produced by deletion of the cis-elements corresponding to TONE and AP1 or ARE1 and 2. Stromal (HIESC-2) and epithelial (HIEEC-22) cells were co-transfected with the lipofectamine 2000 transfection agent (Invitrogen) and the different plasmid constructs possessing the firefly luciferase reporter and a plasmid containing the synthetic Renilla luciferase gene. After transfection, cells were treated for 24 h with IL-1β (1ng/ml), and the promoter activity was expressed as the ratio of firefly luciferase to the renilla luciferase using the Dual-Luciferase Reporter (DLR™) Assay System (Promega, Madison, WI, USA) using a Luminoskan Ascent luminometer (ThermoElectron Corporation, Milford, MA, USA).
FIGURE 4 | Analysis of AKR1B1 promoter activity involved in IL-1β response in human endometrial cells. (A) Schematic representation of different AKR1B1 promoter constructs and corresponding activity following treatment of transfected stromal and epithelial cells treated with IL-1β. (B) Nucleotide sequence of the AKR1B1 promoter MSRR region (−1710 to −1064) showing the organization of the important cis-elements TonE (NFκB), AP1 and anti-oxidant response elements 1 and 2 (ARE1 and ARE2). (C) Relative activity of MSRR cis-elements of the AKR1B1 promoter following stimulation with IL-1β of transfected stromal endometrial cells.

CELL TREATMENTS
HIESC-2 (passages 15–22) and HIEEC-22 (passages 15–22) were cultured in RPMI 1640 without phenol red, containing 50 IU penicillin-streptomycin supplemented with 10% whole FBS during growth and 10% dextran-coated charcoal extracted FBS once cells have reached confluence and for transfection or treatments. Knockdown transfection of cells with AKR1B1 specific siRNA was performed with lipofectamine 2000 for 4 h in culture medium without antibiotic. Unless specified, otherwise cells were treated for 24 h with IL-1β (1 ng/ml) or other agents at specified concentrations in RPMI 1640 medium without serum. For Western blot analysis (described above), the cells were grown in 24-well plates.
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Figure 5 | Association between induction of AKR1B1 and PGF2α production following cytokine stimulations in cells from different human tissues. Cells from different human tissues were grown to confluence and treated with IL-1β or IL-1α 1 ng/ml or TNFα at 10 ng/ml for 24 h. The culture medium was recovered for assay of PGF2α by EIA and protein from attached cells recovered for Western blot analysis of COX-2 and AKR1B1. (A) Prostatic smooth muscle cells, (B) human artery umbilical smooth muscle cells (HUASMC), (C) human cardiomyocytes, and (D) Human Umbilical Endothelial cells (HUVEC).

whereas for Northern blots analysis (described above), cells were in 6-well plates. At the end of the treatment period, the culture medium was recovered and stored at −20°C until evaluation of PGF2α or PGE2 production by EIA whereas protein and RNA were extracted directly from cell monolayer at the bottom of 24-well and 6-well plates, respectively.

Endometrial explants culture
Endometrial tissue was obtained either from biopsies collected with an endometrial curette (Pipelle) and obtained from women aged between 25 and 50 years. Informed consent for donation of anonymous endometrial samples was obtained before tissue collection. The research protocol was approved by the Ethics Committee on Human Research of Centre Hospitalier Universitaire de Québec. Endometrial explants were prepared from fresh biopsies as described previously (Bresson et al., 2011). Endometrial explants were stimulated or not with IL-1β in the presence or absence of the aldose reductase inhibitor ponalrestat (100 μM) under normal (5 mM) or high glucose (25 mM) for 24 h.
FIGURE 6  | Influence of different aldose reductase inhibitors on PGF2α and PGE2 production in human and bovine endometrial cells. Human endometrial stromal [HIESC-2 (A,B)] and human [HIEEC-22 (C)] or bovine epithelial [bEEL (D)] cells were grown to confluence and treated with 1 ng/ml IL-1β (A–C) or 10μM arachidonic acid (AA, D) for 24 h in presence and absence of increasing concentrations of indicated aldose reductase inhibitors. The culture medium was recovered and PGE2 and PGF2α were assayed by EIA. (A) Alrestatin (B–D) ponalrestat. Results are the mean ± SEM of three experiments run in triplicate.

MEASUREMENT OF SORBITOL
We have developed a LC/MS/MS procedure for quantitative measurement of sorbitol generated in briefly, the culture medium was removed for estimation of PGE2 and PGF2α and sorbitol was extracted from explants by mixing with lysis buffer (200 μl), methanol (600 μl), chloroform (200 μl), and water (500 μl). The mixture was centrifuged for 5 min at room temperature and 100 μl of the upper phase was evaporated with a turbo-vap at 45°C for 30 min. The pellet was resuspended with 200 μl of Pyridine 10%: acetic acid acetic acid 10% in acetonitrile and brought at 70°C for 45 min. After drying samples were diluted in 100μl methanol/water (1:1) containing 5 mM ammonium acetate. The samples were spiked with Sorbitol-d8 (Omicron Biochemicals Inc., IN, USA) also used for establishment of a standard curve and analyzed on an Applied Biosystems API-5000 LC/MS/MS. The procedure was optimized to minimize interference with sugars and alcohols such as glucose, fructose, galactose, myoinositol, and galactitol.

PGE2 AND PGF2α EIA
PGE2 and PGF2α were assayed by competitive EIA using acetylcholinesterase-linked PG tracers (Cayman, Ann Arbor, MI, USA) as described previously (Asselin et al., 1996).

DATA ANALYSIS
Data were analyzed by One or Two way ANOVA with Bonferroni as post hoc test with 95% confidence intervals using GraphPad Prism 5 program. For data with less than three columns a two-tailed unpaired t test with 95% confidence intervals was performed using GraphPad Prism 5 program. Data are presented as the mean ± SEM. Each experiment was repeated three times unless specified otherwise. Values were considered statistically significant for P < 0.05.

RESULTS
While glucose and oxidized lipids are passive stable substrates for AKR1B1, generation of PGF2α requires the generation of the unstable PGH2 precursor within the producing cell (Figure 1). Figure 2 illustrates that upon stimulation by IL-1β, the rate limiting enzymes cPLA2 releasing AA from membranes phospholipids, COX-2 generating the PGH2 precursor and AKR1B1 are all increased in both endometrial stromal (A) and epithelial (B) cells. The production of PGF2α is also increased following IL-1β stimulation, a response blocked by the COX-2 selective inhibitor NS-398 (C and D). As was found previously (Pelletier et al., 1999; Bresson et al., 2011), AKR1C3 is present only in epithelial cells.
with AKR1B1 preceding COX-2 preceding mPGES-1. Accordingly, PGF2α release precedes that of PGE2 in both stromal and epithelial cells ($P < 0.05$).

We have cloned the AKR1B1 promoter (~4.5 kb) and generated eight constructs to identify potential response elements sensitive to IL-1β stimulation (Figure 4). IL-1β stimulates the AKR1B1 promoter activity from -583up in both endometrial stromal and epithelial cells (Figure 4A). A MSRR containing two putative antioxidant response elements (ARE) adjacent to T onE and AP1 previously identified for the mouse AKR1B3 (Nishimaka and Yabe-Nishihara, 2005) was also identified for the human AKR1B1 gene at −1710 to −1064 (Figure 4B). The human MSRR region was analyzed in a vector containing a minimal promoter SV40 to identify putative cis-elements of the AKR1B1 gene promoter activity involved in IL-1β response (Figure 4C). The AREs ARE1 and ARE2 appear particularly efficient to confer IL-1β response and may explain the quick activation of AKR1B1 and PGF2α production observed in Figure 3.

The PGF synthase activity of AKR1B1 and its stimulation by IL-1β and TNFα was investigated in cultured cells isolated from different human tissues (Figure 5). In addition to endometrial cells where we made our initial discovery, we have found that IL-1β and to a lesser extent TNFα stimulated AKR1B1, COX-2, and PGF2α production in prostatic smooth muscle cells (A), arterial smooth muscle cells (B), cardiomyocytes (C), and umbilical endothelial cells (HUVEC D). In the last two models, the observed increase in PGF2α production appeared to result from increased COX-2 expression coupled with already high preexisting levels of AKR1B1.

We have compared the inhibition potential of three aldose reductase inhibitors; alrestatin, EBPC (not shown) and ponalrestat on human endometrial cells and tested the latter on bovine endometrial epithelial cells (Figure 6). Alrestatin did not have any effect on PGF2α production (Figure 6A). EBPC inhibited PGF2α production with an apparent EC50 around 10 μM (not shown) whereas ponalrestat was the most potent with an EC50 of 1 μM (Figure 6B). Ponalrestat also inhibited PGF2α in human (Figure 6C) and bovine (Figure 6D) endometrial epithelial cells albeit with reduced efficiency exhibiting an EC50 of 50 μM. Interestingly, ARIs not only inhibited PGF2α but also PGE2 in all cell models.

The potential non-selective inhibition of several components of PG biosynthesis by ARIs was investigated further (Figure 7). When AKR1B1 expression was knocked down by siRNA, similar observation of concurrent inhibition of PGF2α and PGE2 was observed (Figure 7A) suggesting potential interactions between the two biosynthetic pathways. The possibility that AKR1B1 could exert positive regulation of PGE2 production through PGF2α and its FP receptor was tested (Figure 7B). Interestingly, when AKR1B1 and PGF2α production are stimulated with IL-1β, PGF2α, and the concomitant increase of PGE2 is reduced in presence of the FP receptor antagonist AL-8810 suggesting that PGF2α was able to stimulate its own and PGE2 production. The latter hypothesis was tested using the PGF2α analog bimatoprost which stimulated PGE2 release in a dose dependent manner (Figure 7C).

Finally, the relative affinity and potential competition between the PGFS and polyol pathway was tested under close to in vivo and is not induced by IL-1β suggesting minimal contribution to PGF2α production under these conditions.

The time course of PG biosynthetic enzymes activation following IL-1β stimulation was investigated at the mRNA level and associated with PGF2α release (Figure 3). Note that there is basal expression of AKR1B1 gene under non-stimulated conditions and that there is serial transcription of biosynthetic enzyme genes
FIGURE 8 | Influence of glucose concentration on the relative production of sorbitol and PGF2α in fresh explants from human endometrium. Endometrial explants were prepared from fresh endometrial biopsies, cultured for 24 h, and stimulated with IL-1β (10 ng/ml) in presence or absence of Statil (ponalrestat, 100 μM) under normal (LG) 5 mg/ml or High (HG) 25 mg/ml glucose concentrations. PGF2α production was measured by EIA. Results from two different experiments are illustrated.

conditions (Figure 8). Fresh endometrial explants were treated with IL-1β in presence and absence of ponalrestat under normal and high glucose conditions. Interestingly, IL-1β previously demonstrated to increase AKR1B1 expression increased PGF2α or sorbitol production depending on normal or high glucose conditions respectively. This demonstrates that PGF2α and sorbitol are formed competitively depending on the relative availability of AKR1B1 substrates.

DISCUSSION

Prostaglandins are important regulators of female reproductive function and contribute to gynecological disorders. Normal menstruation depend on an equilibrium between vasoconstrictors such as PGF2α (Lockwood and Schatz, 1996; Sales and Jabbour, 2003a) and vasodilators such as PGE2 or nitric oxide (NO) (Tschugguel et al., 1999). Excessive production of contracting prostaglandins create an ischemia-reperfusion response causing painful menstruation or dysmenorrhea (Okazaki et al., 2005) whereas increased vasodilatation leads to abundant menstrual bleeding (Sales and Jabbour, 2003b). NSAIDs represent the most important and widely used drugs on the market and they are all efficient to treat menstrual disorders at some level. However these drugs act at an early step of biosynthesis common to all PGs and not only the isotype responsible for the pathological response (Figure 9). Because of its notorius role on inflammation and pain, the biosynthetic pathway leading to PGE2 has been studied extensively, but that of PGF2α is poorly documented. The data presented in this manuscript address the conditions regulating the newly identified PGFS activity of AKR1B1. In the human endometrium, it has been reported that production of PGF2α is higher in late secretory and menstrual periods of the menstrual cycle (Downie et al., 1974). Accordingly, both AKR1B1 and AKR1C3 enzymes are present in the endometrium throughout the menstrual cycle. By contrast with AKR1B1 expressed in both stromal and glandular epithelial cells and modulated in association with endometrial PGF2α production, AKR1C3 expression is constant and completely absent in stromal cells as was reported previously (Pelletier et al., 1999; Zakharov et al., 2010). The absence of the only currently accepted human PGFS AKR1C3 in stromal cells was surprising because we and others have shown that human endometrial stromal cells produce high levels of PGF2α (Huang et al., 1998; Chapdelaine et al., 2006; Kang et al., 2006). Because of a similar finding in the bovine endometrium and our identification of AKR1B5 as a functional PGFS in that system (Madore et al., 2003), we hypothesized that the human AKR1B1, could also express PGFS activity in the human endometrium. Our initial studies showing the association between AKR1B1 expression and PGF2α production was confirmed unequivocally in the human endometrium using gain and loss of function approaches (Bresson et al., 2011). The PGF synthase activity of AKR1B1 and other members of this family is now acknowledged by leaders in the field of prostaglandin synthases (Nagata et al., 2011; Smith et al., 2011; Watanabe, 2011).

In the present study we highlight the necessary co-activation of genes of the PG biosynthesis pathway to provide the PGH2 precursor substrate to AKR1B1 in order to generate PGF2α. This is in clear contrast with detoxification or aldose reductase activities of AKR1B1 for which substrates are provided by extracellular sources (glucose) or as byproducts of intracellular reactions (oxidized lipids). Endometrial cells express constitutively GLUT2 and GLUT3 and do not require insulin for glucose entry, as such; the endometrium belongs to the category of tissues most adversely affected by poorly controlled glucose levels occurring during insulin resistance or diabetes. Figures 2 and 3 illustrate that genes and proteins of the biosynthetic pathway are increased in a time and dose dependent manner to explain increased PGF2α production following IL-1β stimulation.

In Figure 3 and then in Figures 6 and 7 we observe that PGE2 production closely follows that of PGF2α under both stimulatory and inhibitory conditions. We first considered these results
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**FIGURE 9** Regulation of PGs of the relative PGE2 and PGF2α release in endometrial cells. Following activation of endometrial cells with the cytokine IL-1β, several enzymes leading to the production of PGF2α and PGE2 exhibit increased expression and/or activity. Treatment of cells with NSAIDs will inhibit the generation of PGH2 at the level of COX-1 and COX-2, thus blocking the release of both PGF2α and PGE2 without an alteration of the relative production of the two PGs. Inhibition of AKR1B1 activity with ARIs will inhibit preferentially PGF2α release thus creating an imbalance in favor of PGE2. However, decreased stimulation of the FP receptor should down-regulate the production of PGE2. In presence of high glucose concentration, the stimulation of AKR1B1 may be exacerbated, but glucose should inhibit competitively PGF2α release in which case reduced FP stimulation should also reduce PGE2 production. However, when glucose concentration falls back to normal, high AKR1B1 levels would favor PGF2α release and ischemic responses until PGE2 is able to compensate through EP2 and bring back a silent condition where the balance between PGF2α and PGE2 is re-established.

as artifact and resulting from defective experimental design. However, when we observed that inhibition of FP receptor prevented the increase in PGE2 and that FP receptor stimulation alone was sufficient to stimulate PGE2 production, we proposed that there was a positive feedback loop between PGF2α release and production of PGE2. A link between FP activation and increased PG production could have been anticipated from previous observations in a human endometrial model (Jabbour et al., 2005; Sales et al., 2008) and from our own observations in bovine endometrial cells (Krishnaswamy et al., 2010) given that FP and the oxytocin receptor (OT) share the same signal transduction systems. However, this is the first evidence of a specific link between FP activation and PGE2 production within a homogeneous cell system with potential physiological and pathological relevance. If we hypothesize that PGE2 acting on EP2 receptors and PGF2α acting through FP work as a dyad with opposite actions in the reproductive system, just like the thromboxane/prostacyclin dyad in the vascular system, then the interactions between the two systems are not only probable but also necessary.

Prostaglandins work like a micro endocrine system sensing altered conditions and reestablish locally, optimal conditions for tightly regulated events such as ovulation or recognition of pregnancy in the reproductive system or hemostasis in the vascular system. In this respect, PGF2α exhibits vasoconstrictive and prothrombotic responses comparable to thromboxane with the additional feature that the PGF2α molecule is chemically stable and remains active until catabolism by prostaglandin dehydrogenase, usually through a single passage in the lung. By contrast PGE2/EP2 trigger a vasodilator response with anti thrombotic action similar to prostacyclin but again PGE2 is much more stable than PGI2. Thus it is important to tightly regulate the relative production and release of PGF2α and PGE2 and in this respect; our identification of a feedback loop between PGF2α and PGE2 within endometrial cells has important implications for the observed complications associated with over-expression of AKR1B1. We have initiated the characterization of the AKR1B1 promoter and identified a MSRR containing two putative AREs adjacent to T onE and AP1. The AREs ARE1 and ARE2 appear to be involved in IL-1β response, which supports the local sensing functions of PGs to which AKR1B1 is now a contributing component. The MSRR portion of the AKR1B1 promoter will have to be investigated further under IL-1β stimulation to identify the trans-factors associated with the production of PGF2α. Because of its reported interaction with AREs, NFR2 also known as Nuclear factor (erythroid-derived 2)-like 2 (NFE2L2) is a likely candidate to link increased PGF2α release...
under inflammatory conditions (Nishinaka and Yabe-Nishimura, 2005).

Clinical and pathological implications: Prostaglandins are important regulators of female reproductive functions. In this respect, gene inactivation studies in the mouse have shown that COX-2 and EP2 null mice share similar phenotypes where ovulation and implantation are impaired (Lim et al., 1997; Kennedy et al., 1999). The PGF2α/FP system is much less documented than that of PGE2, but the group of Jabbour in Edinburgh has documented its involvement in endometrial pathologies including menstrual disorders and endometrial cancer (Milne and Jabbour, 2003; Jabbour and Sales, 2004; Sales et al., 2004; Jabbour et al., 2005, 2006; Abera et al., 2010; Catalano et al., 2011) while we (Breuiller-Fouche et al., 2010; Phillips et al., 2011) and others (Olson and Ammann, 2007) associate it with premature delivery. With that in mind, it is likely that AKR1B1 contribute to these pathologies and conversely conditions where AKR1B1 is affected such as diabetes and metabolic syndrome may exhibit a specific prognosis pattern in affected women. In addition, we claim that the female reproductive system constitutes a powerful model to understand the contribution of PGs to physio-pathological conditions in other systems. This is supported by Figure 5 showing the relation between AKR1B1 and PG biosynthetic enzymes in response to IL-18 and their association with PGF2α production in smooth muscles, cardiomyocytes, and endothelial cells. This is all the more important as AKR1B1 known to be highly responsive to diverse physiologic and pathologic conditions and its multiple substrate processing ability combine to create complex combinations well illustrated in Figure 8. For instance, PGF2α has been associated with cardiac hypertrophy (Lai et al., 1996) and a new hamster AKR1B was identified and proposed to contribute to cardiomyopathy through PGF2α release (Sakamoto and Sugamoto, 2011). Moreover, knowing that high glucose induce AKR1B1 expression we see that as long as glucose levels remain high, there will be little increase in PGF2α release, but when glucose goes down, the high levels of AKR1B1 will generate increased production of PGF2α (Figure 9). In turn, high PGF2α will trigger pro-ischemic responses (Yuhki et al., 2010) that fortunately will be compensated by the feedback loop release of PGE2 exerting opposite action thus bringing back homeostasis (Mandal et al., 2005). Normal tension results from the action of EP2 opposing FP receptors in the vascular system, but adjacent tissues also express other EP receptors responsible for inflammation and pain. Therefore the relative silent condition resulting from PGE2 compensation of overproduction of PGF2α by AKR1B1 will likely generate pain or hyperalgesia increasing the probability to take NSAIDs. Not that far ago, the best pain killer on the market with minimal gastrointestinal toxicity was rofecoxib (Vioxx), the most powerful medication to date eradicating pain by reducing mPGE1 generated PGE2. If someone taking Vioxx was under the silent (increased AKR1B1) condition, then the compensatory PGE2/EP2 would not be available to counteract the ischemic response to PGF2α thus increasing the risks of heart failure. Interestingly, even after adjustment of risk factors, reexamination of Vioxx events (Baron et al., 2008) reveal that people suffering from type two diabetes were eight times more at risk to die from Vioxx than non-diabetics. By contrast people taking both Vioxx and Aspirin, thus blocking the AKR1B1 generated PGF2α were protected. Therefore, it is worth investigating if and how AKR1B1 potentially contributed to the failure of Vioxx.

In conclusion, the ability of AKR1B1 to release PGF2α brings a new angle to look at mechanisms responsible for pathophysiologic conditions in the woman reproductive and other systems especially in association with diabetes.

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