

Prostaglandin E2 triggers cytochrome P450 17 α hydroxylase overexpression via signal transducer and activator of transcription 3 phosphorylation and promotes invasion in endometrial cancer

JIEQI KE, ZHEN SHEN, MIN LI, CHENG PENG, PING XU, MEIMEI WANG,
YI ZHU, XUEFEN ZHANG and DABAO WU

Department of Obstetrics and Gynecology, Anhui Provincial Hospital, Hefei, Anhui 230001, P.R. China

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Abstract. Prostaglandin E2 (PGE2) is the most common prostaglandin in the human body, meaning that its malfunction impacts on the development of numerous diseases. Prostaglandin E synthase 2 (PTGES2) is involved in the synthesis of PGE2. In the present study, immunohistochemistry of PTGES2 was performed in 152 patients with endometrial cancer and in 66 patients with normal endometria. The results indicate a notable association among increased expression of PTGES2 and age ($P=0.0092$) and the depth of myometrial invasion ($P<0.0001$). Reverse transcription-quantitative polymerase chain reaction and western blot analysis demonstrated that cytochrome P450 17 α hydroxylase (CYP17), an enzyme for androgen synthesis, is overexpressed following PGE2 stimulation via signal transducer and activator of transcription 3 (STAT3) phosphorylation. ELISA also detected increased androgen (testosterone) secretion. Further invasion of endometrial cancer cells was induced at high androgen levels and when CYP17 was overexpressed. Furthermore, the present study observed that CYP17 is overexpressed via STAT3 phosphorylation in endometrial cancer cells, which grow at a high concentration of PGE2, resulting in increased androgen secretion. Concentrations of estrogen and progesterone were not elevated, while the concentration of androgens was. Overall, a high concentration of androgens caused increased invasion of endometrial cancer cells. A high concentration of androgens, which is initiated by a high expression of PTGES2 and a high concentration of PGE2, is an important promoter of myometrial invasion in endometrial cancer.

Introduction

Endometrial cancer is the most common reproductive cancer in women (1). In China, the incidence of endometrial cancer is 7.44 cases per 1,000,000 individuals (2), and in the USA, the disease causes 8,590 mortalities every year (3).

Prostaglandin E2 (PGE2) is the most common prostaglandin in humans and was first reported in 1936 (4). A previous study has revealed that PGE2 has a significant function in reproductive, neuronal, metabolic and immune systems (5). In two previous studies by Che *et al* (6) and Zhu *et al* (6,7), certain cytokines, such as interleukin (IL)-6 and oncostatin M, were revealed to promote proliferation and invasion of endometrial cancer cells. Certain studies have also reported that PGE2 serves a role in the development of ovarian, breast and colorectal cancer (8-10). However, to the best of our knowledge, no studies have investigated the mechanism of the role served by PGE2 in endometrial cancer.

The signal transducer and activator of transcription 3 (STAT3) transcription factor was identified as a downstream effector of inflammatory mediators that modulates gene expression and metabolism (11). STAT3 underwent phosphorylation and activated the expression of numerous STAT3-regulated genes (12). STAT3 is usually highly expressed in cancer cells, and high levels of STAT3 prompt a poor outcome in ovarian cancer, glioblastoma, breast cancer and prostate cancer (13-17).

The cytochrome P450 (CYP) superfamily of enzymes mediates the catalytic conversion of drugs to reactive products that bind to macromolecules, including proteins and DNA. CYP enzymes account for ~75% of total drug metabolism (18). CYP17 α hydroxylase (CYP17), an enzyme involved in androgen synthesis, has been implicated in the pathogenesis of numerous cancer types (19). A previous study has observed that inhibiting CYP17 is beneficial in prostate carcinoma (20), suggesting that CYP17 may be an important factor in hormone-associated cancer types.

In the present study, the PGE2 synthase-promoted expression of a P450 enzyme in endometrial cancer cells was analyzed, and the results identified CYP17 as the main enzyme involved in this process. These factors resulted in enhanced invasion in endometrial cancer cells.

Correspondence to: Professor Dabao Wu, Department of Obstetrics and Gynecology, Anhui Provincial Hospital, 17 Lu Jiang Road, Hefei, Anhui 230001, P.R. China
E-mail: wudabao5907@126.com

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Materials and methods

Reagents and antibodies. PGE2 (cat. no. 363-24-6) and androgen (cat. no. 1424-00-6) were purchased from Sigma-Aldrich (Merck KGaA, Darmstadt, Germany). JSI-124 (cat. no. 2222-07-3) was obtained from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). Prostaglandin E synthase 2 (PTGES2) antibody (cat. no. 10881-1-AP, dilution at 1:1,000) was purchased from Protein Tech Group, Inc. (Chicago, IL, USA). Antibodies against CYP17 (cat. no. ab125022, dilution at 1:2,000), CYP19 (cat. no. ab35604, dilution at 1:1,000), IL-1 (cat. no. ab8320, dilution at 1:2,000), pSTAT3 (phosphorylated Y705, cat. no. ab76315, dilution at 1:20,000 and phosphorylated S727, cat. no. ab32143, dilution at 1:2,000) were purchased from Abcam (Cambridge, UK). The ELISA kits for estrogen (cat. no. CEA461Ge, 96T), androgen (testosterone, cat. no. CEA458Ge, 96T) and progesterone (cat. no. CEA459Ge, 96T) were purchased from Cloud-Clone Corp. (Houston, TX, USA).

Patients and samples. Tissue samples for immunohistochemistry (IHC) were obtained from 152 patients (all females aged between 34 and 79 years, with a mean age of 53.6 years) with endometrial cancer who had undergone surgical resection at Anhui Provincial Hospital (Hefei, China) between 2010 and 2016, and from 66 patients with normal endometria during curettage of the uterus. Clinicopathological characteristics including age, FIGO stage and grade (21), myometrial invasion and nodal metastasis were included. The project was approved by the Human Investigation Ethics Committee of Anhui Provincial Hospital and informed consent was obtained from all patients prior to the study.

Cell lines and culture conditions. The human endometrial cancer Ishikawa cell line, the 293T cell line and primary endometrial (PE) cells were obtained from Dr Feizhou Jiang (Department of Obstetrics and Gynecology, The First Affiliated Hospital of Soochow University, Suzhou, China). Ishikawa, PE and 293T cells were grown in Dulbecco's modified Eagle's medium (DMEM)/F-12 (Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) supplemented with 10% fetal bovine serum (Gibco; Thermo Fisher Scientific, Inc.). Ishikawa, PE and 293T cells were incubated at 37°C in a humidified atmosphere containing 5% CO₂. All experiments were performed at the third passage following thawing.

Total RNA extraction, reverse transcription-quantitative polymerase chain reaction (RT-qPCR). Total RNA was isolated from Ishikawa cells using TRIzol reagent (cat no. 15596-026; Invitrogen; Thermo Fisher Scientific, Inc.) and cDNA was prepared using the reverse transcriptase kit (Takara Biotechnology Co., Ltd., Dalian, China). RT-qPCR was conducted using an ABI Prism 7500 sequence detection system (Applied Biosystems; Thermo Fisher Scientific, Inc.) and performed with the SYBR Green PCR Master mix (Toyobo Life Science, Osaka, Japan). According to supplier's protocol, the PCR conditions comprised 95°C for 30 sec, followed by 40 cycles of amplification (95°C for 3 sec and 60°C for 30 sec). The 2^{-ΔΔC_q} method was used to analyze the relative changes in gene expression (22). The results were expressed relative to

the internal reference gene GAPDH. Sequences of the primer pairs used are listed in Table I.

Western blot analysis. For western blot analysis, cells were lysed in cell lysis buffer (Beyotime Institute of Biotechnology, Nantong, China) for 30 min at 4°C. Total protein, determined using a bicinchoninic acid assay (cat. no. P0010S; Beyotime Institute of Biotechnology). A total of 15 μg protein per lane, was fractionated using SDS-PAGE (10% gel) and was transferred onto polyvinylidene fluoride membranes, blocking with 5% skim milk which dissolved in DMEM for 2 h. The membranes were then incubated with the appropriate aforementioned primary antibodies (IL-1, CYP17, CYP19, pSTAT3-S727 and pSTAT3-Y705) for 24 h in 4°C, followed by incubation with horseradish peroxidase-conjugated secondary antibody (cat. no. sc-280786, dilution at 1:10,000 for 2 h in 4°C, Santa Cruz Biotechnology, Inc.). Following three further washes in TBS, the proteins were detected and visualized using an electrochemiluminescence system (Pierce; Thermo Fisher Scientific, Inc.) and GAPDH (cat. no. ab184531; dilution at 1:10,000; Abcam) was used as an internal control.

IHC. The tissue sections were initially fixed in 10% formalin solution at 4°C for 2 days and paraffin embedded. The tissue sections were subsequently subjected to microtome sectioning (5 μm). The sections were placed on glass slides and immersed with 100% ethanol (5 min) and boiling water (30 min) three times. The endogenous peroxidase activity was blocked by immersing the sections in freshly prepared 10% H₂O₂ and 10% methanol in 1X PBS for 20 min. The sections underwent trypsin treatment (0.1% trypsin in 0.1% CaCl₂) for 10 min to cleave the protein crosslinks to assess the antigen and epitope. Nonspecific antigens were blocked using 4% bovine serum albumin (BSA; Sigma-Aldrich; Merck KGaA, Darmstadt, Germany) for 2 h at room temperature. The membranes were incubated with anti-PTGES2 (1:100; ProteinTech Group, Inc.) overnight at 4°C. Following incubation, the sections were thoroughly washed with 1X PBS and incubated with a goat anti-rabbit secondary antibody (dilution, 1:3,000; cat no. ab6721; Abcam) for 1 h at room temperature. Following washing to prevent non-specific binding, the sections were stained with diaminobenzidine (DAB; cat no. ab64238; Abcam). The percentage of positively stained cells was rated as follows: 0 points, 0%; 1 point, 1-25%; 2 points, 26-50%; 3 points, 51-75%; and 4 points, ≥75%. The staining intensity was rated in the following manner: 0 points, negative staining; 1 point, weak intensity; 2 points, moderate intensity; and 3 points, strong intensity. Subsequently, immunoreactivity scores for each case were obtained by multiplying the values of the two parameters described above. The average score for all 5 random fields at x200 magnification was used as the histological score (HS). Tumors were categorized into 2 groups based upon the HS: The low-expression group (HS, 0-6) and the high-expression group (HS, 7-12).

Transwell invasion assays. For Transwell invasion assays, the upper side of an 8-μm pore, 6.5-mm polycarbonate Transwell filter (Corning Incorporated, Corning, NY, USA) chamber was uniformly coated with Matrigel basement membrane matrix (BD Biosciences, Bedford, MA, USA) for 2 h at 37°C prior

Table I. Primer sequences for quantitative polymerase chain reaction analysis.

Gene	Forward (5'-3')	Reverse (5'-3')
PTGES2	CTTCCTTTTCTGGGCTTCG	GAAGACCAGGAAGTGCATCCA
GAPDH	GAAGGTGAAGGTCCGAGTC	GAAGATGGTGTATGGGATTTC
TNF- α	TGGCCTCCCTCTCATCAGTT	ATCGGCTGGCACCACCTAGTT
CYP19	TGGAAATGCTGAACCCGATAC	AATTCCCATGCAGTAGCCAGG
CYP17	AGAATTCTCTGGTCCGCC	TTCTCCAGTTTCTGGCCA
IL-1	GCCCTAAACAGATGAAGTGCTC	GAACCAGCATCTTCCTCAG
IL-10	GGCACCCAGTCTGAGAACAG	ACTCTGCTGAAGGCATCTCG

PTGES2, prostaglandin E synthase 2; TNF- α , tumor necrosis factor- α ; CYP, cytochrome P450; IL, interleukin; siCYP17, cytochrome P450 17 α hydroxylase small interfering RNA.

to the cells being added. A total of 2×10^4 Ishikawa cells were seeded in serum-free DMEM/F-12 into the upper chamber of a Transwell filter (in triplicate) and were incubated in 37°C for 48 h. Invasive cells, which had reached the lower chamber, which contained DMEM/F-12 supplemented with 10% FBS, were fixed in 4% paraformaldehyde, stained in 0.5% crystal violet (Beyotime, 20°C, 10 min) and counted using a confocal microscope (Olympus, Shibuya, Japan). A total of 5 fields was counted for each Transwell filter. Each field was counted and images were captured at x200 magnification.

Transfection. Control groups and siCYP17 groups were created. Ishikawa cells with DMEM medium only was used as the control group. To inhibit the expression of the target gene, high-performance liquid chromatography-purified small interfering CYP17 siRNAs (1 μ l in 50 μ l DMEM medium; 5'-3': GCUGGAGAAGAUCAUUUGU.) were prepared according to the sequence of the target gene. A scrambled siRNA with no homology to any known human mRNA was used as a negative control (siCo). siRNA oligonucleotide duplexes were synthesized by Shanghai GenePharma Co., Ltd. (Shanghai, China). The sequences of the siRNA oligonucleotides are provided in Table I. Cells were seeded onto 6-well plates at 70-80% confluence and grown overnight prior to transfection. Transfection of the cells with the siRNA or siCo was accomplished using the Lipofectamine 2000 transfection reagent (Invitrogen; Thermo Fisher Scientific, Inc.) according to the manufacturer's protocols in 48 h.

Plasmids. Flag-CYP17 was subcloned into a PiggyBac vector (Shanghai GenePharma Co., Ltd.) and transfected with a help vector. To obtain a stable and pure CYP17-overexpressing (CYP17 OE) cell population, selection with 300 μ g/ml hygromycin (Santa Cruz Biotechnology, Inc.) was performed in 37°C until the control cells died. For the immunofluorescence staining, CYP17-Flag was subcloned to the pLenti-GIII-CMV-IRES-puro-SV40-GFP vector (Shanghai GenePharma Co., Ltd.). To obtain a pure cell population, 2 μ g/ml puromycin (Santa Cruz Biotechnology, Inc.) selection was performed after 24 h of transfection at 37°C. The DMEM/F12 medium was then changed after 12 h when the blank control cells died. All the cells were visually GFP-positive under a fluorescence microscope. CYP17 plasmids were obtained from Shanghai

GenePharma Co., Ltd. The plasmids were transfected with Lipofectamine 3000 into Ishikawa cells according to the manufacturer's protocols.

ELISA. Concentrations of the estrogen, androgen and progesterone hormones were detected in culture medium of PE cells and Ishikawa cells using solid phase sandwich ELISA according to the manufacturer's protocols (Cloud-Clone Corp.). The hormone assay sensitivity was 0.1 pg/ml and the assay range was 1.03-20,000 pg/ml. For statistical analysis, culture medium was independently collected three times. The ELISA kits used were as follows: ELISA kit for Estradiol (estrogen, cat. no. CEA461Ge, 96T), ELISA kit for Testosterone (cat. no. CEA458Ge, 96T) and ELISA kit for Progesterone (cat. no. CEA459Ge, 96T), all purchased from Cloud-Clone Corp.

Statistical analysis. All quantitative data are presented as the mean \pm standard deviation. Data were analyzed using one-way analysis of variance (ANOVA) with a least significant difference test. All statistical analyses were performed using SPSS 17.0 (SPSS, Inc., Chicago, IL, USA). $P < 0.05$ was considered to indicate a statistically significant difference. All experiments were performed ≥ 3 times.

Results

PTGES2 expression is increased in endometrial cancer. PTGES2 is involved in the synthesis of PGE2. Recent studies have indicated that PGE2 may be a mitogen with a role in a number of cancer types (8-10). IHC was performed in normal endometria and endometrial cancer tissues. Compared with that in the normal endometria, the expression of PTGES2 was significantly upregulated in the endometrial cancer tissues (Fig. 1A and B; Table II). Statistical analysis revealed that increased expression of PTGES2 was significantly associated with the age of the patient ($P = 0.0092$) and the depth of myometrial invasion ($P < 0.0001$), but not with any other characteristics (FIGO stage, grade and nodal metastasis, Table III). As PTGES2 is required for the synthesis of PGE2, an ELISA was performed to determine the PGE2 concentration in Ishikawa cells, a typical endometrial cancer cell line. A significantly higher concentration of PGE2 was observed in these cells compared with that observed in the PE cells and

Table II. Expression of PTGES2 in normal endometria and endometrial cancer.

Groups	Patients, n	HS of PTGES2, n		P-value
		Low expression group (HS<6)	High expression group (HS≥6)	
Normal endometria	66	49	17	
Endometrial cancer	152	56	96	<0.001 ^a

^aP<0.05, determined using the χ^2 test. PTGES2, prostaglandin E synthase 2; HS, histological score.

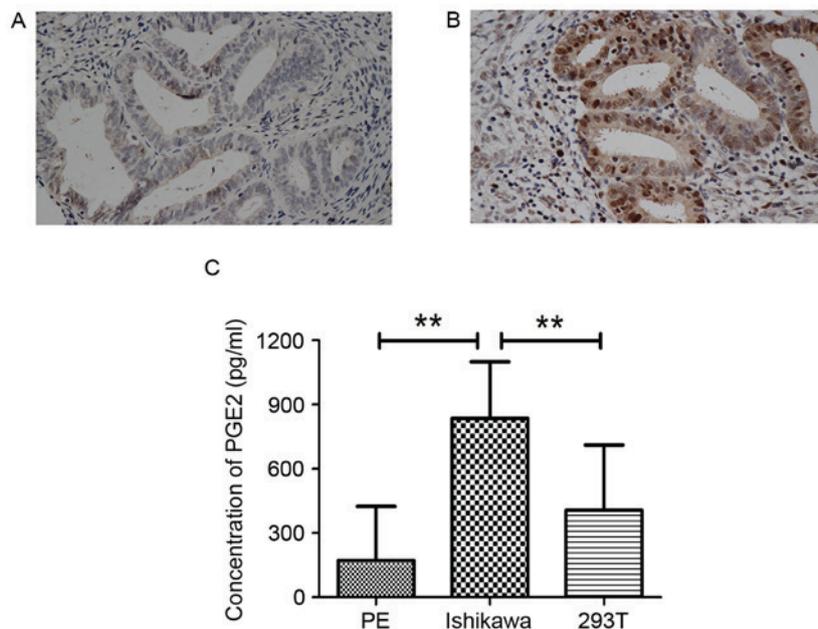


Figure 1. PTGES2 expression in different endometrial cells and PGE2 concentration in different cell types. Representative micrographs of PTGES2 staining in (A) endometrium and (B) endometrial cancer tissues (magnification, x400). (C) ELISA for PGE2 concentration in PE cells, Ishikawa cells and 293T cells. All cell types were cultured for 48 h. **P<0.01, analyzed by one-way analysis of variance. PTGES2, prostaglandin E synthase 2; PGE2, prostaglandin E2; PE, primary endometrial.

the 293T cells (Fig. 1C). This confirmed that endometrial cells grow in a high concentration of PGE2. Therefore, the Ishikawa cells were selected for further study.

PGE2 promotes CYP17 expression in endometrial cancer cells via STAT3. RT-qPCR was performed to test the expression of certain potential mRNA targets identified in previous studies (23-27). As the results indicated, CYP17 expression was significantly increased following PGE2 stimulation, whereas the differences among other cytokines were not significant (Fig. 2A). Subsequent western blot analysis was performed to investigate CYP17 expression changes following PGE2 stimulation (Fig. 2B). As with the RT-qPCR results, CYP17 expression increased mainly in these reported targets. In a previous study, prostaglandin E2 receptor 4 (EP4) was identified as the receptor of PGE2 (28). As EP4 is not a direct DNA promoter, it is unable to directly stimulate CYP17 expression. Subsequently, phosphorylation of the DNA binding protein, STAT3, was examined in Ishikawa cells. The results demonstrated that two STAT3 phosphorylated residues at S727 and Y705 exhibited increased CYP17 expression in

PGE2-stimulated Ishikawa cells. However, this increase was more notable at S727 than at Y705 and the increase of CYP17 was prevented by the presence of JSI-124, an inhibitor of phosphorylation (Fig. 2C). These results suggest that STAT3 may be a downstream protein of PGE2 and that PGE2 increases CYP17 expression.

High expression of CYP17 promotes androgen secretion and invasion in endometrial cancer cells. CYP17 is known to be the key enzyme involved in the synthesis of sex hormones. The concentrations of 3 hormones, estrogen, androgen (testosterone) and progesterone, in Ishikawa cells, were determined using ELISA. The results indicated that the concentration of androgen in the Ishikawa cells was more elevated than the concentrations of the other 2 hormones, compared with that in the PE cells or the control group (Fig. 3A). For further study, Ishikawa cells inhibited with CYP17 siRNA (siCYP17) and others overexpressing CYP17 (CYP17 OE) were prepared (Fig. 3B-D). CYP17 OE Ishikawa cells had been transfected with Flag-CYP17 plasmids, and exhibited an increased androgen concentration (Fig. 3E). Subsequently,

Table III. Associations between PTGES2 expression and clinicopathological characteristics in endometrial cancer.

Characteristics	Patients, n	HS of PTGES2, n		P-value
		Low expression group (HS<6)	High expression group (HS≥6)	
Total	152	56	96	
Age, years				
≥55	78	21	57	0.0092 ^a
<55	74	35	39	
FIGO stage				
Stage I-II	126	48	78	0.4804
Stage III-IV	26	8	18	
Grade				
G1	49	18	31	0.9856
G2-G3	103	38	65	
Myometrial invasion (depth)				
≤1/2 myometrium	48	35	13	<0.0001 ^a
>1/2 myometrium	104	21	83	
Nodal metastasis				
Positive	22	7	15	0.5990
Negative	130	49	81	

^aP<0.05, determined using the χ^2 test. HS, histological score; PTGES2, prostaglandin E synthase 2; FIGO, International Federation of Gynecology and Obstetrics.

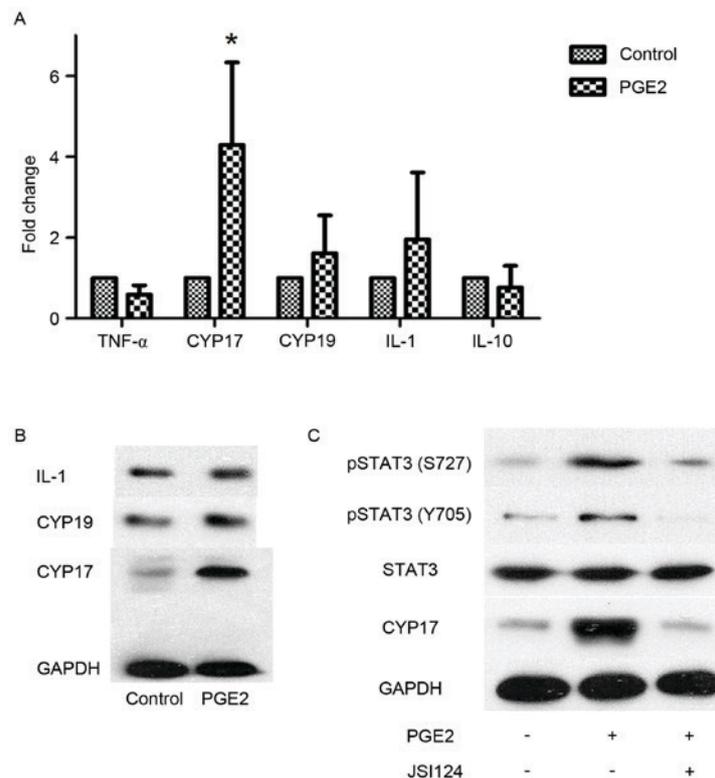


Figure 2. PGE2 promotes CYP17 expression in endometrial cancer cells via STAT3. (A) Reverse transcription-quantitative polymerase chain reaction analysis for TNF- α , CYP17, CYP19, IL-1 and IL-10 in Ishikawa cells following stimulation with PGE2 (1×10^{-9} mol/l). *P<0.05, analyzed by one-way analysis of variance. (B) Western blot analysis for IL-1, CYP19 and CYP17 in Ishikawa cells following stimulation with PGE2 (1×10^{-9} mol/l). (C) Western blot analysis for pSTAT3 S727, pSTAT3 Y705 and CYP17 in Ishikawa cells following stimulation with PGE2 (1×10^{-9} mol/l) with or without JSI-124 inhibition. PGE2, prostaglandin E2; CYP, cytochrome P450; STAT3, signal transducer and activator of transcription 3; TNF- α , tumor necrosis factor α ; IL, interleukin; PGE2, prostaglandin E2; pSTAT3, phosphorylated STAT3.

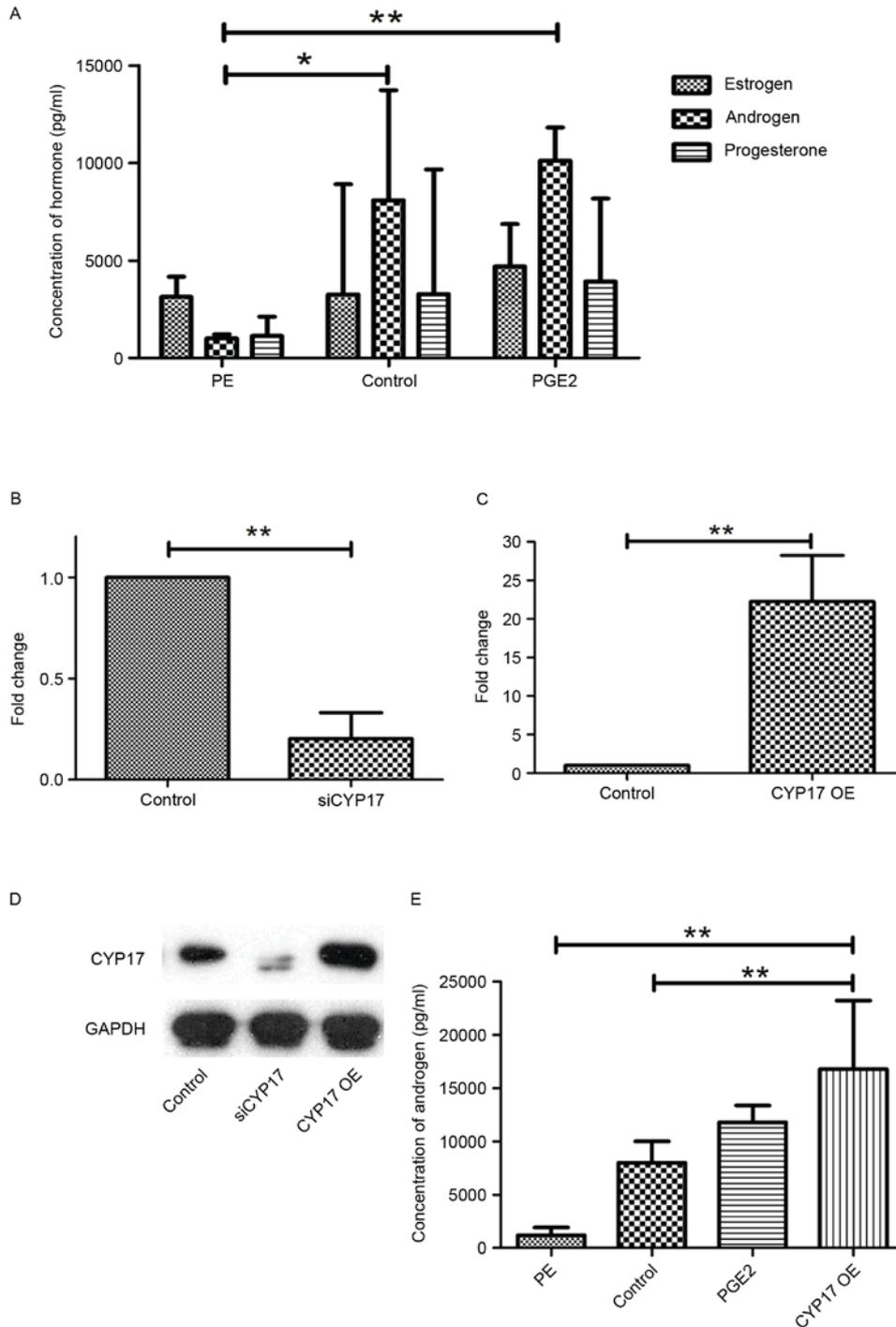


Figure 3. High expression of CYP17 promotes the secretion of androgens. (A) ELISA for estrogen, androgen and progesterone concentrations in PE cells, Ishikawa cells (control) and PGE2-stimulated Ishikawa cells. All cells were cultured for 48 h. Reverse transcription-quantitative polymerase chain reaction analysis for Ishikawa cells (B) following transfection of siCYP17 and (C) in the CYP17 OE group. (D) Western blot analysis of Ishikawa cells in control groups, siCYP17 groups and CYP17 OE groups. (E) ELISA was performed to determine the concentration of androgens in PE cells, Ishikawa cells (control), PGE2-stimulated Ishikawa cells and CYP17 OE Ishikawa cells. All cells were cultured for 48 h. *P<0.05, **P<0.01. CYP17, cytochrome P450 17 α hydroxylase; PE, primary endothelial; siCYP17, CYP17 small interfering RNA; CYP17 OE, CYP17 overexpression.

invasion was evaluated using a Transwell assay, the results of which indicated that siRNA Ishikawa cells had significantly lost their invasion ability when compared with the original Ishikawa cells following PGE2 stimulation (Fig. 4A and B). Ishikawa cells were classified as follows: The control group, the low androgen group (treated with 10⁻⁷ g/l androgen), the high androgen group (treated with 10⁻⁵ g/l androgen) and

the CYP17 OE group. The Transwell results demonstrated that, compared with the control groups, invasion in the high androgen group and the CYP17 overexpression group was significantly increased (Fig. 4C and D). This suggests that CYP17 overexpression and a high concentration of androgen promotes invasion of endometrial cancer cells, initiated by PGE2 stimulation.

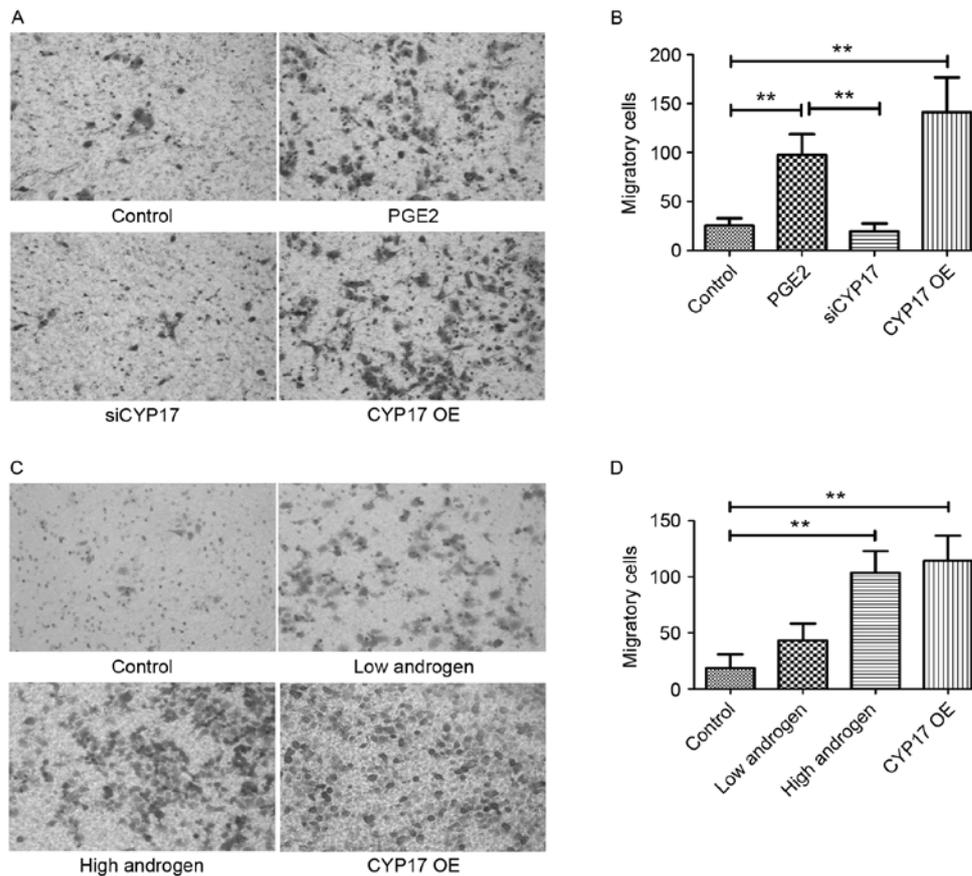


Figure 4. High expression of CYP17 promotes invasion in endometrial cancer cells. (A) Transwell migration assay images of Ishikawa cells in control, PGE2-stimulated, siCYP17 and CYP17 OE groups. Cells were stained with crystal violet. (B) The number of Ishikawa cells in control, PGE2 stimulated, siCYP17 and CYP17 OE groups (averaged across 5 random images). (C) Transwell migration assay images of Ishikawa cells in control, low androgen (treated with 10^{-7} g/l androgen for 48 h), high androgen (treated with 10^{-5} g/l androgen for 48 h) and CYP17 OE groups. Cells were stained with crystal violet. (D) The number of Ishikawa cells in control, low androgen, high androgen and CYP17 OE groups (averaged across five random images). ** $P < 0.01$, analyzed by one-way analysis of variance. Micrographs were taken at x200 magnification. CYP17, cytochrome P450 17 α hydroxylase; PGE2, prostaglandin E2; siCYP17, CYP17 small interfering RNA; CYP17 OE, CYP17 overexpression.

Discussion

The mediators and cellular effectors of inflammation are vital components of the local tumor microenvironment (29). Inflammation that promotes tumor development is an acknowledged enabling hallmark of cancer (30). PGE2 is an important inflammatory factor that acts as a tumor promoter in several cancer types (31,32). In the present study, PTGES2, the synthase of PGE2, was revealed to be highly expressed in endometrial cancer cells, resulting in high expression of PGE2 in the microenvironment of endometrial cancer cells. As demonstrated in previous studies (28,32), EP4 is the key PGE2 receptor in endometrial cancer, but the details of its function in cancer cells are complex and remain unclear (28). In the present study, PGE2 promoted CYP17 expression in endometrial cancer cells via STAT3.

The results of the present study indicated that CYP17 was more highly expressed following PGE2 stimulation. PGE2 is known to serve a role in endometrial cancer cells through its receptor, EP4 (28). With this in mind, in the present study, it was subsequently observed that STAT3 phosphorylation was increased with increased CYP17 expression. Therefore, STAT3 may promote CYP17 expression, via EP4, following

PGE2 stimulation in endometrial cancer cells. Subsequently, the concentration of androgens in endometrial cancer cells was determined and was revealed to be increased following CYP17 overexpression. As a result, invasion of endometrial cancer cells increased, accompanying cell behavior changes.

The cytochrome P450 family 17, subfamily A, member 1 gene provides instructions for making a member of the CYP enzyme family. Similar to other CYP enzymes, CYP17 is involved in the synthesis of steroid hormones (33). This group of hormones includes sex hormones, such as androgen and estrogen, which are required for normal sexual development and reproduction. The CYP17 enzyme performs two important reactions in this process, and existing evidence indicates that unopposed androgens contribute to the tumorigenesis and promotion of endometrial cancer (34,35). Patients with polycystic ovary syndrome, who usually have high androgen levels, have an increased risk of developing endometrial cancer (35,36). The results of the present study support this association and propose a potential mechanism that may begin with inflammation in the endometrium. Inflammation of the endometrium increased PGE2 levels in the microenvironment and resulted in increased CYP17 expression in endometrial cells and a higher concentration of

total androgen in the extracellular environment. This change further results in tumorigenesis and invasion of endometrial cancer cells.

With regards to the mechanism by which PGE2 acts upon cellular gene expression, the present study observed that STAT3 serves an important role in promoting CYP17 expression in endometrial cancer cells. Previous studies have demonstrated that the STAT3 signaling pathway is persistently activated in tumors (37,38). pSTAT3 forms homodimers or heterodimers, translocates into the nucleus and transactivates the expression levels of cyclin D1, vascular endothelial growth factor and matrix metalloproteinases-2/-9 genes. The protein products of these genes are involved in the regulation of tumor cell growth, apoptosis, angiogenesis and metastasis (39-42). It was clearly demonstrated that STAT3 is activated in endometrial cancer cells (43), and that STAT3 can stimulate PGE2 production (44). The results of the present study support a role for STAT3 in endometrial cancer, suggesting that a STAT3 inhibitor may be beneficial in the treatment of advanced-stage endometrial cancer when combined with PGE2/EP4 agonists.

In summary, the present study demonstrated a probable mechanism of invasion in endometrial cancer cells. The high expression of PGE2 in the microenvironment of endometrial cancer cells, possibly caused by inflammation, results in CYP17 overexpression in endometrial cancer cells via STAT3. Overexpression of CYP17 results in increased synthesis of androgens, which increases invasion of endometrial cancer cells. These findings also suggest that inflammation in the tumor microenvironment and high androgen levels serve important roles in tumorigenesis and invasion.

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Availability of data and materials

The datasets generated and analyzed in the present study are included in the published article.

Authors' contributions

JK and DW conceived and designed the study. ZS, ML, CP, PX and YZ performed the experiments. MW provided the mutants. JK wrote the paper. XZ reviewed and edited the manuscript, and performed part of experiments such as IHC. JK reviewed and edited the manuscript. All authors read and approved the manuscript.

Ethics approval and consent to participate

Ethical approval was given by the medical ethics committee of Anhui Provincial Hospital. All patients provided consent to participate in this research.

Patient consent for publication

Patients in the manuscript provided written informed consent for the publication of any associated data and accompanying images.

Competing interests

The authors declare that they have no competing interests.

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