Aberrant activation of signal transducer and activator of transcription-3 (STAT3) signaling in endometriosis

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STUDY QUESTION: Are STAT3 signaling molecules differentially expressed in endometriosis?

SUMMARY ANSWER: Levels of phospho-STAT3 and HIF1A, its downstream signaling molecule, are significantly higher in eutopic endometrium from women with endometriosis when compared with women without the disease.

WHAT IS KNOWN ALREADY: Endometriosis is an estrogen-dependent inflammatory condition. Interleukin 6 (IL-6) is an inflammatory survival cytokine known to induce prolonged activation of STAT3 via association with the IL-6 receptor.

STUDY DESIGN, SIZE, DURATION: Cross-sectional measurements of STAT3 and HIF1A protein levels in eutopic endometrium from women with endometriosis versus those without.

PARTICIPANTS/MATERIALS, SETTING, METHODS: Levels of phospho-STAT3 (pSTAT3) and HIF1A were examined in the endometrium of patients with and without endometriosis as well as in a non-human primate animal model using western blot and immunohistochemical analysis.

MAIN RESULTS AND THE ROLE OF CHANCE: Levels of pSTAT3 were significantly higher in the eutopic endometrium from women with endometriosis when compared with women without the disease in both the proliferative and secretory phases. HIF1A is known to be stabilized by STAT3 and IL-6. Our immunohistochemistry results show abundant HIF1A expression within the eutopic endometrial epithelial cells of women with endometriosis. Furthermore, pSTAT3 and HIF1A proteins are co-localized in endometriosis. This aberrant activation of pSTAT3 and HIF1A is confirmed by sequential analysis of eutopic endometrium using a baboon animal model of induced endometriosis. Lastly, we confirmed this IL-6 induction of both STAT3 phosphorylation and HIF1A mRNA expression in Ishikawa human endometrial adenocarcinoma cell line.

LIMITATIONS, REASONS FOR CAUTION: Ishikawa cancer cell line was used to study a benign disease. The peritoneal fluid contains various inflammatory cytokines in addition to IL-6 and so it is possible that other cytokines may affect the activity and expression of STAT3 signaling molecules.

WIDER IMPLICATIONS OF THE FINDINGS: Our results imply that aberrant activation of STAT3 signaling plays an important role in the pathogenesis of endometriosis. Our findings could progress in our understanding of the etiology and pathophysiology of endometriosis and potential therapeutic interventions by targeted pharmacological.

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Key words: endometriosis / STAT3 / HIF1A / IL6
Introduction

Endometriosis is a major cause of infertility and chronic pain, affecting 1 in 10 women of reproductive age and 35–50% of infertile women (Bulun, 2009; May et al., 2011). With an estimated annual societal cost of $22 billion, the economic impact of this disease is profound in the USA (Xiong et al., 2013). Because of a dearth of reliable biomarkers and considerable symptom overlap with other gynecological pathologies, this disease is difficult to diagnose. Time from onset to clinical diagnosis averages 8–11 years (Bulun, 2009). Surgical resection of ectopic lesions and hormonal suppression are currently the gold standards for treatment, but both approaches are associated with a high incidence of relapse and adverse side effects. Although several theories have been proposed for causality, the etiology of this disease is still unclear. Research examining the molecular mechanisms which permit the establishment of this disease at onset will enable earlier and more effective treatment interventions.

Endometriosis is an estrogen-dependent inflammatory condition associated with elevated tissue-, peripheral- and peritoneal-cytokines (Bulun, 2009). Interleukin-6 (IL-6) is a well-characterized pro-inflammatory cytokine previously implicated with this disease (Heinrich et al., 2003). Recent in vitro studies have shown that levels of IL-6 are greater in human endometrial stromal cells (hESCs) derived from the endometriosis biopsies of women with endometriosis when compared with women without the disease (Tsuo et al., 2000). IL-6 is known to signal via the Signal Transducer and Signal transducer and activator of transcription (STAT) family of transcription factors, specifically STAT3 (Wang et al., 2013).

The transcription factor, STAT3 is localized in the cytoplasm until activated by phosphorylation. A variety of cytokines which are leukemia inhibitory factor (LIF), IL-6, IL-11, and epidermal growth factor (EGF) can activate STAT3 (Zhong et al., 1994). After STAT3 activation, it translocates to the nucleus with the formation of homodimers or heterodimers and binds to promoter regions for target gene expression (Darnell, 1997). Using an oligodeoxynucleotide transferred in utero, implantation and decidualization are interrupted by transient and local suppression of STAT3 activity (Nakamura et al., 2006), and also phosphorylation of STAT3 is certainly required for embryonic development (Takeda et al., 1997). Phosphorylated STAT3 is connected to a number of tumor-promoting processes including maintenance of the stem cell, block differentiation, promoting growth and angiogenesis, and regulating the immune response and tumor microenvironment (Carro et al., 2010).

Hypoxia has a pathophysiological effect through the process of disease and regulation of gene expression (Semenza, 2000a,b). Hypoxia-inducible factors (HIFs) are a family of transcription factors implicated in cellular adaptation to low oxygen levels via transcriptional regulation. HIF1 is the master transcription factor and is composed of HIF1α and HIF1β subunits (Bulun et al., 2000). HIF1β subunit is constitutively expressed, whereas HIF1α levels vary in response to hypoxia. HIF1α is normally present only under hypoxic conditions, but can be stabilized by STAT3, nitric oxide (NO) and IL-6 via the epidermal growth factor receptor (EGFR), Ras, mechanistic target of rapamycin (mTOR) and Phosphatidylinositol-3-Kinase and Protein Kinase B (PI3K/AKT) pathways (O'Donnell et al., 2006).

Phosphorylated STAT3 induces HIF1α expression (Couto et al., 2012) while targeting STAT3 blocks HIF1α and VEGF expression (Xu et al., 2005). However, there are currently no studies investigating the correlation between STAT3 and HIF1α-specific signaling in the context of endometriosis. In the present study, we investigated levels of total and activated STAT3 and HIF1α in the endometrium from women with and without endometriosis. Our findings provide new insight into the etiology of endometriosis and provide a new molecular framework useful for the design of new therapeutic strategies.

Materials and Methods

Human endometrium samples

The study has been reviewed and approved by the Institutional Review Boards of Michigan State University, Spectrum Health Medical System (Grand Rapids, MI), Greenville Health System (Greenville, SC), and the University of North Carolina (Chapel Hill, NC). Written informed consent was obtained from all human subjects. Human endometrial samples were obtained through the Michigan State University’s Center for Women’s Health Research Female Reproductive Tract Biorepository, the Greenville Hospital System, and the University of North Carolina. To compare protein expression patterns of eutopic endometrium with and without endometriosis, endometrial biopsies were obtained at the time of surgery from 43 regularly cycling women between the age of 18 and 45. For control eutopic endometrium, five samples were collected from the proliferative phase, seven were from the early secretory phase, four were from the mid secretory phase, and four were late secretory. For endometriosis eutopic endometrium, 10 samples were collected from the proliferative phase, and 13 were from the secretory phase. The presence or absence of disease was confirmed during surgery. Women laparoscopically negative for this disease were placed into the control group, whereas women laparoscopically positive for this disease were placed in the endometriosis group. The endometriosis patients consisted of 6 stage I, 9 stage II, 5 stage III and 3 stage IV of endometriosis. Use of an IUD or hormonal therapies in the 3 months preceding surgery was exclusionary. Histologic dating of endometrial samples was performed by board certified pathologist and subsequently confirmed by an experienced Fertility specialist (B.A.L.).

Baboon endometrium samples

Use of this animal model was reviewed and approved by the Institutional Animal Care and Use Committees (IACUcs) of both the University of Illinois at Chicago and Michigan State University. Endometriosis was induced by intraperitoneal inoculation of menstrual endometrium on two consecutive menstrual cycles as previously described (Afshar et al., 2013). For baboon endometrium, eutopic endometrial tissues were collected from five early secretory phase baboons at pre-inoculation, and 3, 6, and 9 months of post-inoculation.

Cell culture and treatment

The uterine endometrial epithelial cell line, Ishikawa (endometrial adenocarcinoma) were grown in Dulbecco’s Modified Eagle’s Medium with F12 (Invitrogen, Grand Island, NY, USA), supplemented with 10% (v/v) fetal bovine serum (Invitrogen), 50 units/ml penicillin and 50 μg/ml streptomycin (Invitrogen) in an atmosphere of 5% CO2 and 95% air at 37°C. To determine the effects of IL-6, E2+Medroxyprogesterone Acetate (MPA), and WP1066 (Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA) on STAT3 and HIF1α activation, these reagents and inhibitors were directly added to subconfluent Ishikawa cells and then incubated for the indicated time. All experiments were performed in triplicate with independent protein lysates.

Western blot analysis

Endometrial tissue was lysed using a lysis buffer (10 mM Tris–HCl (pH 7.4), 150 mM NaCl, 2.5 mM EDTA, and 0.125% (v/v) Nonidet P-40 supplemented...
with both a protease inhibitor cocktail (Roche, Indianapolis, IN, USA) and a phosphatase inhibitor cocktail (Sigma Aldrich, St. Louis, MO, USA). Thirty µg of total protein was separated on 8% SDS–PAGE gels and transferred into a polyvinylidene difluoride membrane (Millipore Corp., Bedford, MA, USA). After blocking with 0.5% (w/v) casen for 2 h in phosphate-buffered saline (PBS) with 0.1% (v/v) Tween 20 (Sigma Aldrich), membranes were incubated with either anti-STAT3 (Cell Signaling, Danvers, MA, USA), anti-phospho-STAT3 (Cell Signaling), or anti-HIF1A (BD Bioscience, San Jose, CA, USA) antibodies. Total Actin (Santa Cruz Biotechnology, Inc.) levels were examined for loading controls. Following incubation with primary antibody, membranes were exposed to a horse-radish peroxidase–linked (HRP) – secondary antibody and positive immunostained observed using an enhanced chemiluminescence HRP substrate. All experiments were performed in triplicate with independent protein lysates. The band intensity was determined by relative densitometry using ImageJ (National Institute of Health, USA), and normalized against the bands obtained for actin or STAT3.

**Immunohistochemical and immunofluorescence analyses**

Uterine sections were blocked with 10% normal goat serum in PBS (pH 7.5) for immunohistochemistry (IHC) and 10% normal goat serum in PBST (0.01% Triton X-100 in PBS) for immunofluorescence (IF). Sections were exposed to appropriate primary antibody ([anti-STAT3, Cell Signaling], [anti-phospho-STAT3, Cell Signaling] or [anti-HIF1A, BD Bioscience]) in 10% normal goat serum in PBS (pH 7.5) overnight at 4°C. For IHC, sections were incubated with the appropriate secondary antibody (Vector Laboratories, Burlingame, CA, USA). Following exposure to the horseradish peroxidase-conjugated streptavidin substrate, positive immunoreactivity (brown precipitate) was detected using the Vectastain Elite DAB kit (Vector Laboratories). A semi-quantitative grading system (H-score) was used to compare the immunohistochemical staining intensities. The H-score was calculated using the following equation: H-score = Σ Pi (i+1), where i = intensity of staining with a value of 1, 2 or 3 (weak, moderate or strong, respectively) and Pi is the percentage of stained cells for each intensity, varying from 0 to 100% (Ishibashi et al., 2003). For immunofluorescence, the sections were incubated 2 h at RT with the following secondary antibodies: Alexa Fluor 488-conjugated anti-mouse IgG (Invitrogen) and Alexa Fluor 594-conjugated anti-rabbit IgG (Invitrogen). Nuclei of the cells were counterstained with 4′,6-diamidino-2-phenylindole (DAPI; Vector Laboratories).

**Statistical analysis**

Statistical analyses were performed using Student’s t-test, Mann–Whitney U-test, or repeated measures ANOVA or one way ANOVA followed by Tukey’s post hoc multiple range test using the Instat package from GraphPad (San Diego, CA, USA). The Spearman correlation coefficient was used to assess correlations between the levels of pSTAT3 and HIF1A in control and endometriosis. All data are presented as means ± SEM. P < 0.05 was considered statistically significant.

**Results**

**Aberrant activation of STAT3 in eutopic endometrial tissue from women with endometriosis**

To determine whether STAT3 signaling is dysregulated in endometriosis, we first examined the expression of total STAT3 and phosphorylated STAT3 (pSTAT3) in endometrium from patients with and without endometriosis using western blot (Fig. 1A and B). The expression levels of total STAT3 did not differ in the presence of endometriosis. However, levels of pSTAT3 were significantly higher in the endometrium derived from women with endometriosis (the mean of relative band intensity ± SEM: 2.28 ± 0.38) when compared with controls (1.00 ± 0.21) (Student’s t-test, P = 0.0158, Fig. 1A and B). To examine the cell-specific expression of STAT3 and pSTAT3, we next performed immunohistochemical analysis of endometrium from women with and without endometriosis. STAT3 expression was consistently detected in both the stromal and epithelial compartments of endometrium in both the control (the mean of H-score ± SEM: 252.25 ± 14.45) and endometriosis groups (274.35 ± 9.47) (Mann–Whitney U-test, P = 0.25, Fig. 1C and D). In control women pSTAT3 protein was not detected in endometrial cells except decidual cells of late secretory phase (Supplementary Fig. S1). Interestingly, the level of pSTAT3 protein was significantly increased in epithelial cells of endometriosis patients (255.65 ± 9.15) when compared with control patients (14.75 ± 5.39) (Mann–Whitney U-test, P < 0.0001, Fig. 1E and F). These results suggest that dysregulation of STAT3 signaling molecules may play an important role in the pathogenesis of endometriosis.

**HIF1A expression in endometrium from women with endometriosis**

HIF1A is a known target of pSTAT3 and a key mediator of angiogenesis, inflammation, proliferation, self-renewal, and extracellular invasion (Henze and Acker, 2010). The western blot analysis of HIF1A levels show that levels of HIF1A were markedly increased in the endometrium of patients with endometriosis (the mean of relative band intensity ± SEM: 24.68 ± 2.83) when compared with controls (1.00 ± 0.24) (Student’s t-test, P = 0.0002, Fig. 2A and B). Additionally, subsequent immunohistochemistry analysis of uterine paraffin wax sections confirm that HIF1A protein levels are significantly higher in the endometrial epithelium of patients with endometriosis (the mean of H-score ± SEM: 144.35 ± 15.99) when compared with the patients without endometriosis (16.75 ± 6.35) (Mann–Whitney U-test, P < 0.0001, Fig. 2C and D).

**Correlation between pSTAT3 and HIF1A in endometriosis**

Figure 3A shows the correlation between pSTAT3 and HIF1A proteins in both women without and with endometriosis. There was a significant positive correlation between pSTAT3 and HIF1A in the endometrial epithelial cells (Spearman correlation coefficient r = 0.7943, P < 0.0001). To determine whether pSTAT3 colocalized with HIF1A proteins with respect to endometriosis, we performed double immunofluorescence for pSTAT3 and HIF1A (Fig. 3B). Our immunostaining results show that pSTAT3 and HIF1A proteins were colocalized in 13% of epithelial cells from endometriosis patients. In contrast, 1% of epithelial cells were colocalized in patients without endometriosis. These results suggest a strong correlation exists between STAT3 activity and HIF1A expression in the endometrium which may play an important role in the development and progression of endometriosis.

**pSTAT3 and HIF1A expression during progression of endometriosis in a baboon model**

A baboon model has previously been developed to study the pathophysiology of endometriosis (Fazleabas et al., 2002; Braundmeier and
Intraperitoneal inoculation with autologous menstrual tissue under laparoscopic guidance results in the formation of endometriotic lesions with histological and morphological characteristics similar to those seen in women (Fazleabas et al., 2002). To determine if aberrant activation of STAT3 signaling was also evident in this non-human primate model, we examined pSTAT3 and HIF1A expression in the eutopic endometrium of baboons following experimental induction of the disease. pSTAT3 was strongly detected in both the stroma and epithelium of baboon endometrium following induction of the disease (Fig. 4), but levels of STAT3 protein were unchanged (Supplementary Fig. S2). pSTAT3 and HIF1A protein were weakly detected in the endometrium of pre-inoculation (control) animals (the mean of H-score ± SEM: 40.00 ± 11.40 and 12.00 ± 5.83, respectively). The levels of pSTAT3 proteins were significantly increased at 6 months post-

**Figure 1** Aberrant activation of phospho-signal transducer and activator of transcription-3 (pSTAT3) in eutopic endometrial tissue from women with endometriosis. (A) Western blot analysis of STAT3 and pSTAT3 proteins in human endometrium of control and endometriosis. Actin was used as sample-loading control. (B) Quantification of pSTAT3/STAT3 western blot data obtained by densitometric analysis. Representative photomicrograph of immunohistochemical staining of STAT3 (C) and pSTAT3 (E) proteins in human endometrium of control and endometriosis. The immunohistochemical histological score (H-score) of STAT3 (D) and pSTAT3 (F) proteins. The results represent the mean ± standard error of the mean (SEM). *P < 0.05; ***P < 0.001.
inoculation (124.00 ± 21.59, repeated measures ANOVA, \(P < 0.01\)) and 9 months post-inoculation (184 ± 26.94, \(P < 0.001\)) during endometriosis progression (Fig. 4A). Likewise, the levels for HIF1A proteins were significantly increased in the epithelial compartment at 6 (148.00 ± 28.71) and 9 (222 ± 25.98) months post-inoculation (repeated measures ANOVA, \(P < 0.01\) and \(P < 0.001\), respectively) (Fig. 4B). These results suggest that aberrant activation of STAT3 may induce HIF1A proteins during the pathogenesis of endometriosis.

**Figure 2** Aberrant expression of hypoxia-inducible factor I-alpha (HIF1A) in eutopic endometrial tissue from women with endometriosis. (A) Western blot analysis of HIF1A proteins in human endometrium of control and endometriosis. Actin was used as sample-loading control. (B) Quantification of HIF1A western blot data obtained by densitometric analysis. (C) Representative micrograph of immunohistochemical staining of HIF1A proteins in human endometrium of control and endometriosis. (D) The immunohistochemical histological score (H-score) of HIF1A proteins. The results represent the mean ± SEM. ***\(P < 0.001\).

**Figure 3** Correlation and colocalization analysis of phospho-STAT3 and HIF1A in eutopic endometrial tissue from women with endometriosis. (A) Correlation of between pSTAT3 and HIF1A H-score in control (\(n = 20\)) and endometriosis (\(n = 23\)) (\(P < 0.001, r = 0.7943\)). (B) Colocalization of pSTAT3 and HIF1A in the endometrium between women with and without endometriosis by immunofluorescence analysis.
IL-6 contributes to the elevated levels of pSTAT3 and HIF1A expression

Previous studies have shown that IL-6 is over-expressed within the endometrium of women with endometriosis and pSTAT3 is regulated by inflammatory cytokines such as IL-6 (Kroon et al., 2013; Karalok et al., 2014). To determine whether IL-6 can induce phosphorylation of STAT3 and HIF1A expression in endometrial epithelial cells, we treated Ishikawa cells with IL-6 for 12 h and subsequently used western blot analysis to examine the expression levels of STAT3, pSTAT3 and HIF1A. Our results showed that IL-6 treatment significantly increased pSTAT3 (4.23 ± 0.31) and HIF1A (3.45 ± 0.68) expression compared with the control (1.00 ± 0.07 and 1.00 ± 0.08, respectively) but STAT3 levels were unchanged (Fig. 5). These results indicate that this induction of pSTAT3 is not transcriptional regulation of STAT3. Estradiol (E2) + Medroxyprogesterone (MPA) are known moderators of uterine STAT3 function (Lee et al., 2013). Furthermore, E2 + MPA and IL-6 treatment significantly increased pSTAT3 levels (12.17 ± 0.50) when compared with E2 + MPA (1.26 ± 0.08, P < 0.001) and IL-6 alone (4.23 ± 0.31, P < 0.001). These results suggest that IL-6 induces phosphorylation of STAT3 and then HIF1A expression.

Activated STAT3 is required for HIF1A expression

We next examined whether HIF1A expression could be interrupted by inhibition of STAT3 phosphorylation. To examine whether HIF1A expression is downstream of STAT3 activation in uterine epithelial cells, we treated Ishikawa cells with IL-6 in the presence or absence of WP1066, a STAT3 inhibitor, and examined HIF1A, pSTAT3 and STAT3 protein levels using western blot analysis. Our results show a remarkable time-dependent induction of pSTAT3 by IL-6 which could be prevented by pretreatment with WP1066. In addition, WP1066 treatment remarkably reduced HIF1A expression after 6 h of IL-6 treatment (Fig. 6). These results suggest that the HIF1A induction seen in the endometrial epithelial cells is downstream of pSTAT3 induction.

Discussion

Aberrant activation of STAT3 has been identified as both abnormal and oncogenic (He and Karin, 2011; Fagard et al., 2013). STAT3 signaling participates in oncogenesis by stimulating cell proliferation, promoting angiogenesis, mediating immune evasion and conferring resistance to...
apoptosis. Our results herein show that levels of pSTAT3 are significantly higher in the eutopic endometrium from women with endometriosis when compared with controls (Fig. 1). This suggests that tight regulation of STAT3 activity is important for normal uterine function. The relationship between STAT3 and endometriosis in humans is not well known. Given its relevance to the inflammatory process, surprisingly little is known about endometrial STAT3 and its regulation in health and disease. Examining of molecular mechanisms of uterine STAT3 signaling may suggest effective therapies for endometriosis and other uterine disorders.

This study is the first to build on the hypothesis that aberrant activation of STAT3 and HIF1A signaling within the eutopic endometrium is relevant to endometriosis due to dysregulation of cell proliferation, inflammation and steroid hormone signaling. To better understand the correlation of these two transcription factors in endometriosis, we examined correlation and colocalization of pSTAT3 and HIF1A (Fig. 3). Furthermore, this study shows that inhibition of STAT3 can prevent HIF1A expression in epithelial cells (Fig. 6). HIF1A has previously been shown to up-regulate many of the aberrant proteins and factors associated with endometriosis, including increased cellular proliferation (Semenza, 2000a,b), aberrant ESR2 expression (Juhasz-Boss et al., 2011; Wu et al., 2012), and angiogenic factors VEGF (Maybin et al., 2011) and Cyr61 (Gashaw et al., 2008). Leptin is also a known target of HIF1A and can induce endometrial cell proliferation (Wu et al., 2007; Oh et al., 2013). Silencing HIF1A (siRNA) suppresses hypoxia-induced VEGF (Spinella et al., 2007). Our results suggest that elevated pSTAT3 induces and stabilizes HIF1A within the eutopic endometrium of women with endometriosis, subsequently leading to aberrantly increased angiogenesis, MMPs, and proliferation within the eutopic endometrial compartment (Gilabert-Estelles et al., 2007; Aznaurova et al., 2014; Lu et al., 2014).

We describe herein a correlation between pSTAT3 and HIF1A expression levels, which is disease-dependent (Fig. 3). Previous studies using the non-human primate model suggest that the presence of ectopic endometrium (endometriotic lesions) alters the eutopic endometrial gene expression profile (Li et al., 2013; Meola et al., 2013). As in cancer, hypoxia is likely an important effector of the microenvironment of the ectopic endometriotic implants. Indeed, HIF1A has been shown to regulate expressions of ESR1 and ESR2 expression (Wu et al., 2012).
These results suggest that dysregulation of STAT3 and HIF1A signaling may play an important role in the pathogenesis of endometriosis. Western blot analysis showed an increase in both pSTAT3 and HIF1A expression in Ishikawa cells following IL-6 treatment (Fig. 5). Additionally, we observed more colocalization of HIF1A and pSTAT3 proteins in endometrial epithelial cells from women with endometriosis (Fig. 3B). These results suggest that cytokines and disease-specific regulation of pSTAT3 may represent a sentinel difference within the eutopic endometrium from women with and without this disease.

Previous studies have described use of a non-human primate model for the study of endometriosis (D’Hooghe et al., 1994; Fazleabas, 2006). Intraportal inoculation with autologous menstrual tissue results in the formation of endometriotic lesions with histological and morphological characteristics similar to those seen in women (Fazleabas et al., 2002). This baboon model also shows similar markers of progestosterone resistance as previously described in humans (Braundmeier and Fazleabas, 2009). Sequential analysis of eutopic endometrium from these baboons throughout the progression of this disease reveal increased pSTAT3 and HIF1A expression (Fig. 4). Angiogenesis plays an important role in endometriosis. The angiogenic potential of both the endometrium and the peritoneal environment influences lesion establishment, and endometriotic lesions require an adequate blood supply to survive in their eutopic sites (McLaren, 2000; Taylor et al., 2002). Therefore, these results suggest that aberrant activation of STAT3 may induce HIF1A proteins during endometriosis progression. We confirmed HIF1A induction of pSTAT3 in Ishikawa endometrial epithelial cancer cell lines (Fig. 6).

Inflammatory cytokines are responsible for the IL-6 elevation found at higher concentrations within the peritoneal fluid (Buyalos et al., 1992) and serum of women with endometriosis (Pellicer et al., 2002). In animal models of endometriosis, IL-6 is also increased in the serum of rats with surgically induced endometriosis (Boutten et al., 1992). Our results show that phosphorylation of STAT3 can be remarkably increased by IL-6 (Fig. 5) and that inhibition of STAT3 phosphorylation can reduce HIF1A expression (Fig. 6).

JAKs and STATs are critical components of many cytokine receptor systems, regulating growth, survival, differentiation, and pathogen resistance. JAK1 and JAK2 account for STAT3 phosphorylation upon docking with IL-11/gp130 receptor complex (Kamimura et al., 2003). WP1066 inhibits STAT3 activity by inhibiting its upstream transcription factor JAK (Ferrajoli et al., 2007). WP1066 shows a decrease of HIF1A levels in Ishikawa cells, suggesting that STAT3 can regulate angiogenesis through HIF1A. Additionally, S3I-201 inhibits STAT3 activity by inhibiting STAT3 dimerization (Fletcher et al., 2009), whereas cryptotanshinone inhibits STAT3 activity by inhibiting its phosphorylation (Lu et al., 2013). Furthermore, Ishikawa cells are human endometrial epithelial cancer cell lines. Ishikawa cell line, which is one of the few uterine cell lines that expresses functional ESR and PGR (Croxtall et al., 1990).

This cancer cell line is modestly responsive to estrogen, but has lost its inhibitory response to progesterone, despite expression of receptors for both hormones. We have observed activation of STAT3 in Ishikawa cells treated with IL-6 (Fig. 5). Therefore, additional studies are necessary to fully understand the action of STAT3 in other endometrial epithelial cell lines with other STAT3 inhibitors.

In summary, we report for the first time that phosphorylated STAT3 and HIF1A are highly expressed and activated in the endometrium of patients with endometriosis. Our studies provide a better understanding for the aberrant gene expression previously described by many and attributed to progesterone resistance (Young and Lessey, 2010). Progress in our understanding of the etiology and pathophysiology of endometriosis and potential therapeutic interventions by targeted pharmacological agents has been hampered due, in part, to the lack of defined molecular mechanisms. These findings will provide new etiological insight into the development of this disease as well as a molecular framework for the design of new therapeutic strategies.

**Supplementary data**

Supplementary data are available at http://humrep.oxfordjournals.org/.

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**Authors’ roles**

B.G.K., J.-Y.Y., T.H.K. and J.F.L. performed experiments, analyzed the data and wrote the manuscript. J.-H.S. provided helpful discussion and critical analysis of data. S.D.F., A.T.F., and S.L.Y. contributed to sample collection and critical revision of the manuscript. B.A.L and J.-W.J. contributed to the design of the study, interpretation of the data and wrote the manuscript.

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**Conflict of interest**

None declared.

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