GATA-3 Stimulates Proliferation of Endometriotic Cells

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ABSTRACT

Objectives: This study aimed to investigate whether there was a correlation between the expression of GATA binding protein 3 (GATA-3) and the proliferation of endometrium cells from women with endometriosis.

Material and methods: Endometrial tissue was collected from 20 women with endometriosis (experimental group) and 10 women without endometriosis (control group). The primary endometrial cells were then cultured and identified. Each group of cells were further divided into four groups. The control group was divided into control-no treatment (C-N), control-negative control (C-NC, transfected with a scrambled siRNA negative control), control-lentiviral transfection control (C-LC) and control-GATA-3 lentiviral transfection (C-G3); the experimental group was divided into experiment-no treatment (E-N), experiment-negative control (E-NC), experiment-lentiviral transfection control (E-LC) and experiment-siGATA-3 lentiviral transfection (E-siG3). RT-PCR and western blot were used respectively to detect the endometrium cells’ GATA-3 mRNA and protein expression. Further, Cell Counting Kit-8 was used to measure the endometrium cells’ proliferation.

Results: The relative expression of GATA-3 mRNA and protein in the E-N group was significantly higher than in the D-N group and E-siG3 group (P < 0.05). The relative cell proliferation rate in the C-G3 group at 48 h, 72 h and 96 h was significantly higher than in the C-N, C-NC and C-LC groups (P < 0.05), while in the experimental group, the proliferation rate of the E-siG3 group was significantly lower than in the E-N, E-NC and E-LC groups (P < 0.05).

Conclusions: This study suggests the role of GATA-3 in promoting proliferation of endometrial cells; therefore, GATA-3 may play an essential role in the development of endometriosis.

Key words: endometriosis; GATA-3; endometriotic cell; proliferation

INTRODUCTION

Endometriosis, characterized by endometrial-like tissue and stroma outside the endometrial cavity, is known as a commonly encountered disease affecting 5–15% of women in their reproductive age [1]. Although 20–25% of women with endometriosis have no symptoms, patients may present with chronic pelvic pain, dysmenorrhea and infertility [2]. It has been nearly 90 years since this disease was initially described by Sampson [3], and several theories have been proposed to understand and explain the development of endometriosis [4]; however, the current understanding of the exact pathophysiology of endometriosis remains unclear.

Endometrial epithelial cells have been found by many recent studies working as a big source of cytokine production, which plays a very important role in the immune, menstrual and reproductive functions of endometrium [5, 6]. For instance, one study suggested that the imbalance of T helper type 1 and 2 (Th1/Th2) cytokines are of great importance in the development of endometriosis [7]. Interestingly, in lymphocytes, GATA binding protein 3 (GATA-3) has been proved to work as one of the most important cytokine expression regulators [8]. GATA-3, as a factor of the specific transcription of Th2 cell has its binding sites all cover the promoter regions of Th2 cytokines [9]. Importantly, it has been found that during the menstrual cycle, GATA-3 is expressed cyclically in the human endometrial epithelium [8]. Additionally, Peng et al. [10] suggested that the expression of GATA-3 in endometrial epithelial cells of women with endometriosis was higher than that of healthy women. Specially, it has been proposed that number of proliferating endothelial cells increase in the endometrium of endometriosis patients [11, 12], which implies that the proliferation of endometrial cells is important in the patho-
genesis of endometriosis. However, to our knowledge, the role of GATA-3 in the proliferation of endometrial cells has not been investigated.

Objectives

In this study, endometrial tissue was collected from 20 women with endometriosis and 10 women without endometriosis, and the expression of GATA-3 in the endometrium cells of endometriosis patients and normal controls was compared. Further, the proliferation of endometrium cells from endometriosis patients and normal controls was investigated as well. This study may help to provide a deeper understanding on the mechanisms of GATA-3 that is involved in the proliferation of endometrial cells.

MATERIALS AND METHODS

Sample collection

A total of 20 patients (aged 37–49) with late stage endometriosis (experimental group) and 10 patients (aged 37–49) with carcinoma in situ of the cervix without endometriosis (control group) were included. At least six months prior to surgery, no patients received any gonadotropin-releasing hormone (GnRH) nor other hormone or anti-inflammatory medication. The endometrial tissues of all patients were obtained by excision or curettage at Shengjing Hospital affiliated to China Medical University.

The informed consent was signed by all patients or their first-degree relative before operation. Additionally, this study was approved by the Ethics Committee of Shengjing Hospital affiliated to China Medical University and was performed in accordance with the ethical standards. All patients or their first-degree relatives signed an informed consent form prior to operation. Besides, the study was approved by the Ethics Committee of Shengjing Hospital affiliated to China Medical University, and it was carried out based on the ethical standards.

Primary endometrial cell culture and identification

The endometrial tissue samples from all groups were cut into pieces and digested with 0.1% type I collagen enzyme for 1 h at 37°C. The cell suspension was passed through an 80 mesh sieve, then centrifuged at 850 rpm for 5 min. After removing the supernatant, Dulbecco Modified Eagle Medium (DMEM)/F-12 was added to resuspend cells. Then cell suspension was centrifuged again at 800 rpm for 10 min to remove neutrophils and erythrocytes. After that, cells were gently resuspended in DMEM/F-12 supplemented with 10% fetal bovine serum (FBS), and cell suspensions were cultured in flasks at 37°C with 5% CO₂. The endometrial cells were identified by immunocytochemical staining of keratin and vimentin.

Groups

The primary endometrial cells of the experimental and control groups were divided into four groups: the control group into control-no treatment (C-N), control-negative control (C-NC, transfected with negative control scrambled siRNA), control-lentiviral transfection control (C-LC) and control-GATA-3 lentiviral transfection (C-G3); the experimental group into experiment-no treatment (E-N), experiment-negative control (E-NC), experiment-lentiviral transfection control (E-LC) and experiment-siGATA-3 lentiviral transfection (E-siG3).

Lentiviral transfection

The GATA-3 RNA interference (RNAi) lentiviral vector, GATA-3 lentiviral vector, complementary DNA (cDNA) sequential synthesis, and scrambled siRNA negative control were constructed by Shanghai GeneChem Co., Ltd (Shanghai, China). Lentiviral transfection was conducted using a lentiviral vector particle kit purchased from Shanghai GeneChem Co., Ltd (Shanghai, China). The target sequence of siGATA-3 was 5’-AACAUUGCAGGGUCAGCAGA-3’ and of the scrambled control was 5’-TTCTCGGAACTGTACGTT-3’. In detail, cells were incubated with the recombinant lentivirus at the optimal multiplicity of infection (MOI) in serum-free DMEM/F-12 containing 5 mg/mL polybrene at 37°C and 5% CO₂. The media was changed after 15 h of transfection. The transfection efficiency was examined by fluorescence microscopy and flow cytometry after 48 h, and then was confirmed by real-time quantitative polymerase chain reaction (RT-PCR) and western blot.

RT-PCR

Total mRNA was isolated from cells using Trizol Reagent (Takara, Otsu, Shiga, Japan). First-strand cDNA was synthesized using the PrimeScript First Strand cDNA Synthesis Kit. The SYBR PrimeScript RT-PCR Kit (Takara, Otsu, Shiga, Japan) was used for quantitative PCR with glyceraldehyde-3-phosphatedehydrogenase (GAPDH) as an internal control gene. In this assay the specific primers used were: GATA-3 sense, 5’-CGAGATGGCCAGGGGACACTA-3’; GATA-3 reverse, 5’-TGTTCTGACGTAGTGCCACAGG-3’; GAPDH sense, 5’-GGACGTCTCAAGGCGAACAC-3’; GAPDH reverse, 5’-ATGTTGTGAAAGCAGCGCATG-3’. All reactions were performed in the LightCycler 2.0 System (50°C for 2 min; 95°C for 10 min; 45 cycles of 95°C for 15 s and 60°C for 1 min). The GATA-3 relative expression value was calculated using the 2^ΔΔCT method [12].

Western blot

Total protein was collected from the cells of the eight groups using a protein extraction kit (Keygen Biotechnology, Nanjing, China). About 50 μg of protein was separated by
GATA-3 mRNA and protein expression levels were measured by real-time fluorescence quantitative polymerase chain reaction and western blot respectively. The mRNA and protein expression of GATA-3 were significantly higher in the C-N group compared to the E-N group. Additionally, there was no statistical difference found between the C-N and C-LC groups or between the E-NC and E-LC groups.

### Results

#### Relative expression of GATA-3 mRNA

As shown in Table 1 and Figure 1A, the relative expression of GATA-3 mRNA in the E-N group was 2.30 ± 0.03, which was significantly higher than in the D-N (1.00 ± 0.10) and E-siG3 group (0.62 ± 0.20) (P < 0.05). In control groups, the relative expression of GATA-3 mRNA in the C-G3 group (4.64 ± 0.14) was statistically higher than that in the C-N group (P < 0.05). No statistically difference was found between the C-NC and C-LC groups or between the E-NC and E-LC groups.

#### Relative expression of GATA-3 protein

The relative expression of GATA-3 protein was shown in Table 1 and Figure 1B. When comparing the E-N group (2.43 ± 0.12) with the C-N group (1.00 ± 0.19), a significant difference was found between the two groups (P < 0.05). When comparing the C-N group with the C-G3 group, a significantly higher relative expression of GATA-3 protein was found in the C-G3 group (4.10 ± 0.51, P < 0.05). Additionally, there was no statistical difference found between the C-NC and C-LC groups or between the E-NC and E-LC groups.

### Discussion

Endometriosis is an inflammatory disease involving a possible shift towards the Th2-mediated immune response [13, 14]. As a Th2 cell-specific transcription factor, GATA-3 has been found to play important roles in cell proliferation and differentiation of the mammary glands [15], nephric duct [16] and nervous system [17]. In this study, GATA-3 was up-regulated in the primary endometrial cells of endometriosis patients compared to controls. In addition, the
proliferation rate of endometrial cells in the endometriosis group was significantly higher than in the control group. Importantly, after the endometrial cells of endometriosis patients were transfected with GATA-3 siRNA, the proliferation rate significantly decreased. However, in the control group, when the endometrial cells were transfected with GATA-3, their proliferation rate increased significantly.

As mentioned above, cytokines play important roles in the immune and menstrual functions of the endometrium [18].

GATA-3 regulates the expression of cytokines in lymphocytes [8]. Some studies have demonstrated the role of GATA-3 in the endometrium throughout the menstrual cycle and in women with endometriosis [10, 19, 20]. Consistent with the findings above, this study revealed increased expressions of GATA-3 mRNA and protein in the primary endometrial cells of endometriosis patients compared to controls. Therefore, GATA-3 may be used as a biomarker in the diagnosis and treatment of patients with endometriosis.
found that blocking GATA-3 can reduce the proliferation of endometrial cells. Additionally, we also demonstrated that IL-4 stimulated the proliferation of endometrial cells in patients with endometriosis. Importantly, Zhuo et al. have suggested that GATA-3 can promote the proliferation of endometriotic stromal cells. Taken together, we speculated that the role of GATA-3 in the proliferation of endometrial cells in patients with endometriosis may be associated with cytokine IL-8.

Furthermore, IL-4, another Th2 cytokine, has been linked to increased mRNA and protein expression in peripheral blood and peritoneal fluid of patients with endometriosis. The action of IL-4 in Th2 development induces the early expression of GATA-3. In turn, the expression of Th2-specific cytokines including IL-4 can also be induced by GATA-3 with the contributions of other transcription factors. Importantly, Zhuo et al. demonstrated that IL-4 stimulated the proliferation of endometriotic stromal cells. Taken together, we speculated that IL-4 may be another important cytokine implicated in the GATA-3-induced proliferation of endometrial cells in endometriosis patients.

**CONCLUSIONS**

Our results suggest the role of GATA-3 in promoting proliferation of endometrial cells. Additionally, we also found that blocking GATA-3 can reduce the proliferation of endometrial cells. GATA-3 may be used as a biomarker in the clinical diagnosis and treatment of endometriosis. Further studies are required to elucidate and confirm the exact mechanism.

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**Author disclosure and contributions**

All authors declare no competing interests.

P.C. acquired funding, designed the study, collected samples, conducted the experiment and statistical analysis and created tables and figures, and wrote the manuscript. F.R. acquired funding and collected samples. DB.W. designed the study and helped in writing the manuscript. All authors have seen and approved the final version of this manuscript.

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