MOLECULAR AND CELLULAR MECHANISMS OF DISEASE

# miR-141-3p affects apoptosis and migration of endometrial stromal cells by targeting KLF-12



Yiwei Zhang<sup>1</sup> • Juan Yan<sup>2</sup> • Xiaowei Pan<sup>2</sup>

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#### Abstract

Endometriosis is an estrogen-dependent disease that is characterized by pelvic pain and infertility. MicroRNAs have been shown to implicate in the progression of endometriosis. In our study, we used real-time PCR to evaluate the expression of miR-141-3p in endometrial samples. In addition, western blot analysis was used to assess the expression of Krüppel-like factor 12 (KLF-12). The proliferation and migration of ectopic endometrial stromal cells (ESCs) were determined by MTT assay and Transwell assay, respectively. Cell apoptosis was evaluated using a Cell Death Detection ELISA Plus kit. The results showed that miR-141-3p and KLF-12 were significantly different in paired ectopic and eutopic endometrial samples. miR-141-3p overexpression significantly restrained the proliferation and migration and promoted the apoptosis of ectopic ESCs, whereas a decreased level of miR-141-3p was associated with opposite results. Furthermore, dual-luciferase reporter assay confirmed that KLF-12 was a novel target of miR-141-3p, while it also decreased the effects of miR-141-3p on the proliferation, apoptosis, and migration of ectopic ESCs. Our data suggested that enhanced expression of miR-141-3p suppressed the proliferation and migration of ectopic ESCs and promoted their apoptosis via targeting KLF-12. Our results may provide a novel potential therapeutic target for the treatment of endometriosis.

Keywords miR-141-3p · Proliferation · Apoptosis · Ectopic endometrial stromal cells · Krüppel-like factor 12

# Introduction

Endometriosis is a chronic and estrogen-dependent gynecological disorder that is characterized by the presence and growth of functional endometrial-like tissue outside the uterus

#### Highlights

5. KLF-12 is directly regulated by miR-141-3p.

6. KLF-12 reverses the miR-141-3p-induced suppression of cell proliferation and migration.

7. KLF-12 inhibits the miR-141-3p-mediated induction of cell apoptosis.

⊠ Yiwei Zhang yiwei\_angz@163.com

<sup>2</sup> Department of Gynecology, Shijiazhuang First Hospital, Shijiazhuang 050011, Hebei, People's Republic of China [8]. Endometriosis can severely decrease the quality of life because of pelvic masses, pelvic pain, dysmenorrhea, infertility, and cancerous lesions [2, 27]. Although the disorder is known to be a benign disease, the migratory and invasive properties of ectopic endometrial stromal cells (ESCs) are similar to those of malignant cancer cells [1]. However, the mechanisms underlying the migration of ectopic ESCs are not well known.

MicroRNAs, well known as small noncoding RNAs with 20-22 nucleotides, regulate the expression of target genes and control a variety of cellular functions, including apoptosis, proliferation, migration, and invasion [13, 21, 34]. Numerous studies suggested that microRNAs are involved in the development of human diseases, including cardiovascular disorders, gynecological diseases, cancer, and inflammatory diseases [20, 28, 33]. Furthermore, abnormal levels of microRNAs have been associated with the progression of several female reproductive tract diseases, including preeclampsia [15], endometrioid endometrial adenocarcinoma [35], uterine leiomyomata [4], ovarian adenocarcinoma [3], and endometriosis [24]. Multiple studies have indicated that miR-141-3p may be decreased in endometriosis [10, 26]. However, the role of miR-141-3p in endometriosis is still largely unknown.

<sup>1.</sup> miR-141-3p is obviously decreased in endometriosis.

<sup>2.</sup> KLF-12 is significantly upregulated in endometriosis.

<sup>3.</sup> Restoration of miR-141-3p expression in ectopic ESCs inhibits cellular proliferation and migration.

<sup>4.</sup> Restoration of miR-141-3p expression in ectopic ESCs induces the apoptosis in ectopic ESCs.

<sup>&</sup>lt;sup>1</sup> Department of Obstetrics and Gynecology, The Third Hospital of Hebei Medical University, 139 Ziqiang Road, Shijiazhuang 050051, Hebei, People's Republic of China

In this work, we explored the roles and potential mechanisms of miR-141-3p in the development of endometriosis. We found that the level of miR-141-3p was significantly decreased in ectopic endometrium, while Krüppel-like factor 12 (KLF-12) was upregulated. Functional experiments indicated that miR-141-3p overexpression was associated with a clear decline in the proliferation and invasion of ectopic ESCs, and it promoted apoptosis of ectopic ESCs. Moreover, KLF-12 is a direct target of miR-141-3p, obviously attenuated the inhibitory effect of miR-141-3p on the proliferation and migration of ectopic ESCs. These findings may provide insight into developing a new therapeutic biomarker for endometriosis treatment.

# Materials and methods

### **Tissue acquisition**

Twenty patients with endometriosis at our hospital were selected for collection of paired ectopic (EC) endometrial samples and eutopic (EU) endometrial samples. Control samples were collected from the endometrium of patients with hysteromyoma. All participants had regular periods and had no hormone therapy within 3 months before surgery. Patients with polycystic ovary syndrome or inflammatory were excluded from our study. Each patient signed an informed consent form, and the hospital ethics committee approved this study. Their clinical profiles are shown in Table 1.

# Isolation and culture of normal, eutopic, and ectopic ESCs

We isolated primary human endometrial stromal cells according to a previously described method with slight modification [19]. Briefly, the tissues were minced into 1 mm pieces after washing PBS for three times and then digested with 0.1% type

Table 1	Clinical	characteristics	of	study	subjects
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Study subjects	Controls	Endometriosis	
Age (years)	$29.52 \pm 8.06$	33.13±6.13	
Gravidity			
0	19 (76.0%)	4 (20.0%)	
≥1	6 (24.0%)	16 (80.0%)	
Parity			
0	13 (52.0%)	17 (85.0%)	
≥1	12 (48.0%)	3 (15.0%)	
Cycle phase			
Proliferative	16 (64.0%)	15 (75.0%)	
Secretory	9 (36.0%)	5 (25.0%)	

Age is presented as mean  $\pm\,SD;$  the other variables are presented as the number (%)

II collagenase for 1 h. Cells were centrifuged with 500*g* for 5 min and then resuspended in Ham's F12/DMEM (1:1). Cell suspension was passed through 100- $\mu$ m and 40- $\mu$ m nylon mesh to separate glandular epithelial cells. After 12 h of adherence, the culture medium was replaced to remove nonadherent glandular epithelial cells and blood cells. After purification, stromal cells were grown in Ham's F12/DMEM (1:1) including heat-inactivated 10% fetal bovine serum (FBS; Gibco, Grand Island, NY, USA), 1% penicillin/streptomycin, and 0.25 µg/mL amphotericin B. Cells were incubated in a humidified atmosphere consisting of 5% CO<sub>2</sub> and 95% air at 37 °C. In order to avoid contamination by epithelial cells, the isolated stromal cells were immune-stained with vimentinand cytokeratin-specific antibodies.

#### **Construction of adenovirus**

The AdMax (Microbix) system was used to generate adenoviruses harboring KLF-12 (NCBI Reference Sequence: NM\_007249.4) according to a previously described method. After being packaged, HEK293A cells amplified the viruses, which were then purified and dialyzed with CsCl banding and 10 mmol/L Tris-buffered saline, respectively. The Adeno-X Rapid Titer kit (BD, Franklin Lakes, NJ, USA) was used to measure the titering in HEK293A cells, following the manufacturer's protocol.

# **Cell transfection**

The ectopic ESCs were respectively transfected with miR-NC (negative control), miR-141-3p mimic, miR-141-3p inhibitor, mutant KLF-12 3'-UTR, wild-type KLF-12 3'-UTR, and Ad-Con or Ad-KLF-12 using Lipofectamine 2000 reagent (Thermo, Waltham, MA, USA) following the manufacturer's illustration. Afterward, cells were incubated at 37 °C for 24 h.

# Quantitative real-time polymerase chain reaction assay

We isolated total RNA, using an RNA Isolation Kit (Takara Biotechnology, Dalian, China) in accordance with the manufacturer's instructions. cDNA was synthesized by using PrimeScript RT reagent kit (Takara, Tokyo, Japan). Following the manufacturer's protocols, we measured the expression of miR-141-3p with an ABI 7500 Real-time PCR system (Bio-Rad Laboratories Inc., Hercules, CA, USA), using a TaqMan miRNA assay (Qiagen, Shanghai, China). We chose U6 as the internal control for measuring the expression of miRNA. In addition, we used the ABI 7500 Real-time PCR system to measure the KLF-12 expression through SYBR Green Master Mix (Thermo), with glyceraldehyde 3phosphate dehydrogenase (GAPDH) as the internal control for mRNA expression.

#### Western blot

We respectively used a total protein extraction kit (Kaiji Biological Inc., Nanjing, China) and the BCA kit (Beyotime, Shanghai, China) to extract the protein and determine the protein concentration. SDS-PAGE was performed to separate the proteins, which were then electrotransferred to polyvinylidene difluoride (PVDF) membranes and incubated with the appropriate primary and secondary antibodies. The optical density (OD) value of the bands was measured by an enhanced ECL kit and a film, while the Image-Pro Plus software (Pierce Biotechnology Inc., Rockford, IL, USA) was used to quantify the protein levels.  $\beta$ -Actin was used for normalization.

# **Dual-luciferase reporter assay**

First, we respectively synthesized the wild-type KLF-12 3'-UTR (KLF12 3'-UTR Wt) and mutant KLF-12 3'-UTR (KLF-12 3'-UTR Mut) of KLF-12 and then cloned them into the pGL3 luciferase reporter vector (Promega, Madison, WI, USA). Then, 24-well plates were used to seed cells until reaching 70% confluence. We subsequently cotransfected cells with KLF12 3'-UTR Wt or KLF-12 3'-UTR Mut and pRL-TK plasmid (Promega), along with 30 nM miR-141-3p mimic, or negative control was transfected into cells using the Lipofectamine 2000. The dual-luciferase reporter assay was performed to analyze Renilla luciferase activity after transfection for 48 h.

### MTT assay

Cell viability was evaluated using the MTT assay (Sigma-Aldrich Co., St. Louis, MO, USA). After transfection for 24 h, ectopic ESCs were grown into 96-well plates. Subsequently, 20  $\mu$ L of MTT solution was mixed into the media and incubated for 4 h, and we then added 200  $\mu$ L of dimethyl sulfoxide to solubilize the formazan crystals for 15 min. The OD value was determined at 490 nm.

### Transwell assay

Adjusting the density of ectopic ESCs to  $5 \times 10^3$  cells per well, we then seeded the upper Transwell chamber and added 400 µL of serum-free cell culture medium. One milliliter of medium containing 5% FBS was added to the lower chamber, and the cells were incubated for 6 h. Using a microscope (Olympus, Tokyo, Japan) with a × 10 objective, we counted the number of cells that had migrated through the pores in five independent visual fields. Migrated cells were stained with Giemsa dye for observation.

#### Cell apoptosis analysis

A Cell Death Detection PLUS ELISA (Roche Diagnostics, Indianapolis, IN, USA) was performed to detect DNA fragmentation as a molecular marker of apoptosis. Cells were grown in 96-well plates and transfected with miR-NC, miR-141-3p mimic, or miR-141-3p inhibitor. Cytoplasmic fractions were collected and transferred to streptavidin-coated 96-well plates, followed by incubated with anti-histone antibody and anti-DNA antibody for 2 h. The absorbance of each well was determined at 405 nm using a microplate reader and normalized to the concentration of protein.

### **Statistical analysis**

The SPSS software (ver. 13.0; SPSS, Chicago, IL) was used to analyze all data, which are expressed as means  $\pm$  SD. One-way analysis of variance followed by the Student–Newman–Keuls test was used to determine significant differences of multiple groups. Statistical significance was defined as P < 0.05.

# Results

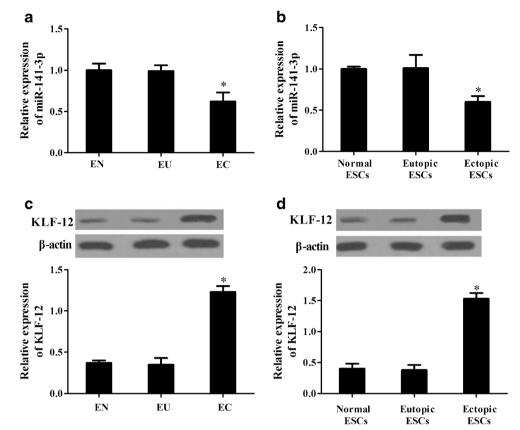
# The level of miR-141-3p is clearly decreased in endometriosis

The expression of miR-141-3p in the patients with and without endometriosis was measured by RT-qPCR. We found that miR-141-3p was prominently decreased in ectopic endometrium compared with normal endometrium and eutopic endometrium (P < 0.05, Fig. 1a). In addition, miR-141-3p in ectopic ESCs was significantly lower than that in normal ESCs and eutopic ESCs (P < 0.05, Fig. 1b). Furthermore, we conducted a western blot analysis to determine the expression of KLF-12 in the endometrial samples from patients with and without endometriosis. As shown in Fig. 1c, KLF-12 was clearly increased in ectopic endometrium. Similarly, we observed that KLF-12 was markedly upregulated in ectopic ESCs (P < 0.05, Fig. 1d). These data indicated that the expression of miR-141-3p and KLF-12 was shown as inverse relationship.

# Restoration expression of miR-141-3p inhibits the proliferation and migration of ectopic ESCs

To study the effects of miR-141-3p in endometriosis, we transfected the miR-141-3p mimic or miR-141-3p inhibitor into ectopic ESCs and then evaluated the proliferation and migration of ESCs. As evident in Fig. 2a, compared with the miR-NC group, miR-141-3p was significantly elevated in the ESCs transfected with miR-141-3p mimic. The expression of

Fig. 1 The expression of miR-141-3p and KLF-12 in endometrial tissues and ESCs is different. a, b Differential levels of the miR-141-3p in endometrial tissues (a) and ESCs (b) were evaluated by qRT-PCR. c, d Differential expressions of the KLF-12 in endometrial tissues (c) and ESCs (d) were detected by western blot. EN normal endometrium, EU eutopic endometrium, EC ectopic endometrium. \*P < 0.05compared with the normal endometrium or ESC group



miR-141-3p was prominently reduced after transfection with miR-141-3p inhibitor. As shown in Fig. 2b, compared with the miR-NC group, miR-141-3p overexpression showed a lower growth rate; however, upregulation of miR-141-3p significantly promoted ectopic ESC proliferation. Moreover, the results of Transwell assays indicated that miR-141-3p overexpression substantially reduced the migration of ectopic ESCs, whereas suppression of miR-141-3p had an opposite effect (P < 0.05, Fig. 2c). These results revealed that miR-141-3p impeded the proliferation and migration of ectopic ESCs.

# Enforced expression of miR-141-3p induces apoptosis in ectopic ESCs

The incidence of apoptosis was detected using a Cell Death Detection PLUS ELISA.

Enhanced expression of miR-141-3p significantly promoted apoptosis in the ectopic ESCs, while knockdown of miR-141-3p showed the opposite effect (P < 0.05, Fig. 3a). Besides, Bcl-2 and Bax expression was determined by western blot. As shown in Fig. 3b, miR-141-3p overexpression

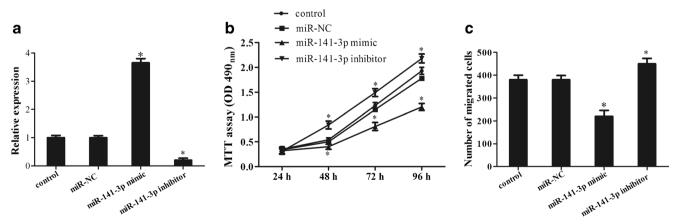


Fig. 2 miR-141-3p overexpression inhibited the proliferation and migration of ectopic ESCs. a Ectopic ESCs transfected with the miR-141-3p mimic or inhibitor to overexpress or suppress miR-141-3p expression, respectively. b MTT assay was performed to determine the

cell viability of ectopic ESCs after transfection with miR-NC or miR-141-3p mimic or miR-141-3p inhibitor. c Cell migration was measured using Transwell assay. \*P < 0.05 compared with the miR-NC group

downregulated the expression of Bcl-2 and raised the expression of Bax, whereas inhibiting the expression of miR-141-3p remarkably elevated Bcl-2 expression and decreased Bax expression. These findings demonstrated that miR-141-3p contributes to the apoptosis of ectopic ESCs.

# KLF-12 is directly regulated by miR-141-3p

We cotransfected cells with wild-type KLF-12 3'-UTR or mutant KLF-12 3'-UTR along with miR-141-3p mimic or miR-NC and then performed dual-luciferase reporter assay to explore whether KLF-12 is a direct target of miR-141-3p. The predictive binding sites between miR-141-p and KLF-12 are shown in Fig. 4a. We found that the luciferase activity of pGL3-KLF-12-WT was decreased compared with the pGL3-KLF-12-Mut (P < 0.05, Fig. 4b). In addition, the protein level of KLF-12 was detected. The results indicated that miR-141-3p overexpression significantly decreased the protein level of KLF-12, while the miR-141-3p inhibitor-transfected cells had a significantly increased level (P < 0.05, Fig. 4c). These data showed that KLF-12 is a direct target of miR-141-3p.

# KLF-12 reverses the effects of miR-141-3p on the proliferation and migration of ectopic ESCs

To understand the potential mechanism of miR-141-3p in the regulation of proliferation and migration during endometriosis, the proliferation and migration of ESCs were detected after cotransfection with Ad-KLF-12 and miR-141-3p mimic. MTT assay indicated that miR-141-3p mimic inhibited the proliferation of ectopic ESCs, and this effect was reversed by KLF-12 overexpression (P < 0.05, Fig. 5a). Similarly, the inhibitory role of miR-141-3p in migration was blocked by gain function of KLF-12 (P < 0.05, Fig. 5b).

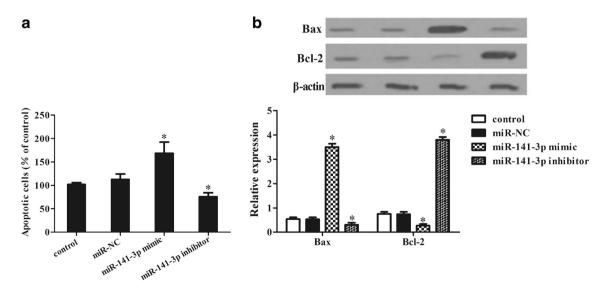
### KLF-12 reversed the roles of miR-141-3p in apoptosis

Furthermore, we evaluated the apoptosis of ectopic ESCs after cotransfection with Ad-KLF-12 and miR-141-3p mimic. As shown in Fig. 6a, overexpression of KLF-12 inhibited the increase in ectopic ESC apoptosis induced by miR-141-3p (P < 0.05). Consistent with these results, overexpression of KLF-12 downregulated Bax expression and raised the level of Bcl-2 in miR-141-3p mimic-transfected ectopic ESCs (P < 0.05, Fig. 6b). These findings support that miR-141-3p facilitated the apoptosis of ectopic ESCs via targeting KLF-12.

# Discussion

Aberrant expression of miRs is implicated in the endometriosis [10]. It has been reported that miR-141-3p is decreased in certain kinds of cancer [12, 36, 38]. Recent report revealed that miR-141 was notably decreased in the ectopic endometrium [29]. In this study, we demonstrated that the levels of miR-141-3p were significantly reduced in ectopic endometrium, suggesting that miR-141-3p may be involved in the development of endometriosis.

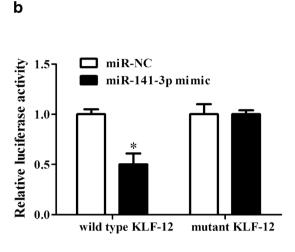
Previous research showed that the migratory ability of endometriotic cells may play an important role in the development of endometriosis [22]. miR-141-3p plays an integral role in the progression and development of tumors. Recent



**Fig. 3** miR-141-3p promoted ectopic ESCs apoptosis. **a** Cell apoptosis was examined in ectopic ESCs transfected with miR-141-3p mimic or miR-141-3p inhibitor. **b** The expression of Bax and Bcl-2 was examined

in ectopic ESCs transfected with miR-141-3p mimic or miR-141-3p inhibitor by western blot. \*P < 0.05 compared with the miR-NC group





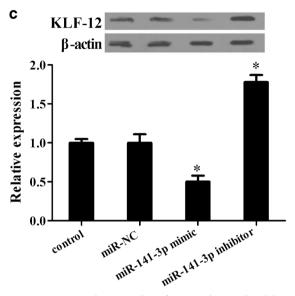


Fig. 4 miR-141-3p directly targeted KLF-12 in ectopic ESCs. a The putative binding sites for miR-141-3p in KLF-12. b Relative luciferase activity in ectopic ESCs cotransfected with miR141-3p or miR-NC, and wild-type or mutant 3'-UTR KLF-12 was measured by dual-luciferase

reporter assay. c The expression of KLF-12 in ectopic ESCs transfected with miR-141-3p mimic, or miR-141-3p inhibitor was detected by western blot.  ${}^{*}P < 0.05$  compared with the miR-NC group

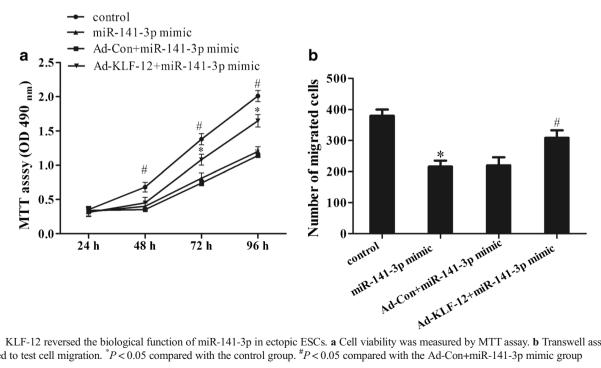


Fig. 5 KLF-12 reversed the biological function of miR-141-3p in ectopic ESCs. a Cell viability was measured by MTT assay. b Transwell assay was utilized to test cell migration. P < 0.05 compared with the control group. P < 0.05 compared with the Ad-Con+miR-141-3p mimic group

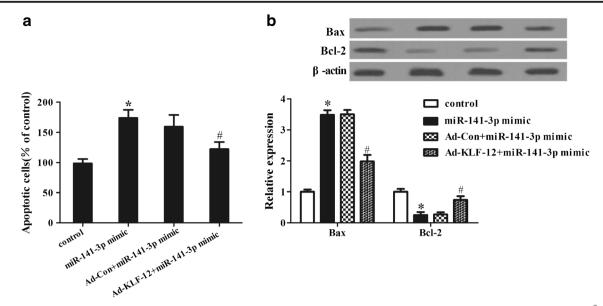


Fig. 6 KLF-12 attenuates miR-141-3p-induced apoptosis. a Cell apoptosis was examined in ectopic ESCs transfected with miR-141-3p mimic or miR-141-3p mimic and Ad-KLF-12. b The expression of Bax and Bcl-2 was examined in ectopic ESCs transfected with miR-141-3p

mimic or miR-141-3p mimic and Ad-KLF-12 by western blot. \*P < 0.05 compared with the control group. \*P < 0.05 compared with the Ad-Con+miR-141-3p mimic group

report showed that overexpression of miR-141-3p promoted cell migration and invasion in cervical cancer [17]. However, Hou et al. reported that miR-141-3p inhibited the proliferation, migration, and invasion of hepatocellular carcinoma cell lines [11]. miR-141-3p served as a tumor suppressor through inhibiting the proliferation and invasion of NSCLC cells [18]. Similarly, miR-141-3p also inhibited the growth and metastasis of papillary thyroid cancer cells [7]. Consistently, our data demonstrated that enforced expression of miR-141 inhibited cell migration and triggered cell apoptosis in ectopic ESCs. However, the opposite trends were observed upon inhibition of miR-141-3p.

miRNAs are known to exert their biological functions through binding to the 3'-UTR of target genes. These targets have been suggested to constitute molecular pathways potentially involved in the development of endometriosis. Members of the Krüppel-like factor (KLF) family of transcription factors have critical roles in regulating cell differentiation, physiologic function, and phenotypic modulation [5, 14, 30, 32]. Genetic studies have demonstrated that KLF-12 plays an important role in embryonic development, tumor genesis, and progress [23, 25, 31]. For example, KLF-12 promoted colorectal cancer growth by regulating early growth response protein 1 [16]. Mounting studies have confirmed that KLF-12 acts as an oncogene in multiple cancers, including gastric cancer [6], colorectal cancer [16], and basal-like breast carcinoma [9]. In addition, a recent study demonstrated that KLF12 acts as a negative regulator during the decidualization of ESCs

[12, 30, 37] . In the present study, the results indicated that the protein levels of KLF-12 differed in ectopic versus eutopic endometrial tissue. In addition, we demonstrated that KLF-12 was a direct target of miR-141-3p. Importantly, the results of rescue experiments and functional assays strongly demonstrated that restoration of KLF-12 abolished the inhibitory effects of miR-141-3p on proliferation and migration in ectopic ESCs. Moreover, enforced expression of KLF-12 reversed the increased apoptosis in ectopic ESCs mediated by miR-141-3p overexpression.

In conclusion, we found that miR-141-3p may suppress cell proliferation and migration, promoted apoptosis via targeting KLF-12 in ectopic ESCs. Our study suggests that miR-141-3p may be a potential target in developing treatment against endometriosis.

Authors' contributions Yiwei Zhang and Juan Yan conceived and designed the study; Yiwei Zhang, Juan Yan, and Xiaowei Pan conducted experiments and collected the data; Yiwei Zhang and Juan Yan analyzed and interpreted the experiments results; Yiwei Zhang wrote the first draft of the manuscript; Yiwei Zhang, Juan Yan, and Xiaowei Pan revised the paper. All authors contributed to and approved the final manuscript.

#### Compliance with ethical standards

The hospital ethics committee approved this study.

**Conflict of interest** The authors declare that they have no conflict of interest.

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