Mifepristone inhibits proliferation, migration and invasion of HUUA cells and promotes its apoptosis by regulation of FAK and PI3K/AKT signaling pathway

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Purpose: The aim was to investigate mifepristone effects on endometrial carcinoma and the related mechanism.

Methods: HUUA cells were treated with DMEM containing different concentrations of mifepristone. HUUA cells treated with 100 μmol/L mifepristone were named the Mifepristone group. HUUA cells co-transfected with pcDNA3.1-PI3K and pcDNA3.1-AKT overexpression vectors were treated with 100 μmol/L mifepristone and named the Mifepristone + PI3K/AKT group. mRNA expression was detected by quantitative reverse transcription PCR. Protein expression was performed by Western blot. Cell proliferation was conducted by MTT assay. Wound-healing assay was conducted. Transwell was used to detect cells migration and invasion. Apoptosis detection was performed by flow cytometry.

Results: Mifepristone inhibited HUUA cells proliferation in a dose-dependent manner. Compared with HUUA cells treated with 0 μmol/L mifepristone, HUUA cells treated by 50–100 μmol/L mifepristone had a lower wound-healing rate, a greater number of migrating and invasive cells (P < 0.01), as well as a higher percentage of apoptotic cells and Caspase-3 expression (P < 0.01). When HUUA cells were treated with 50–100 μmol/L of mifepristone, FAK, p-FAK, p-PI3K and p-AKT relative expression was all significantly lower than HUUA cells treated with 0 μmol/L of mifepristone (P < 0.01). Compared with the Mifepristone group, HUUA cells of the Mifepristone + PI3K/AKT group had a lower cell growth inhibition rate and percentage of apoptotic cells (P < 0.01).

Conclusion: Mifepristone inhibited HUUA cells proliferation in a dose-dependent manner. Compared with HUUA cells treated with 0 μmol/L mifepristone, HUUA cells treated by 50–100 μmol/L mifepristone had a lower wound-healing rate, a greater number of migrating and invasive cells (P < 0.01), as well as a higher percentage of apoptotic cells and Caspase-3 expression (P < 0.01). When HUUA cells were treated with 50–100 μmol/L of mifepristone, FAK, p-FAK, p-PI3K and p-AKT relative expression was all significantly lower than HUUA cells treated with 0 μmol/L of mifepristone (P < 0.01). Compared with the Mifepristone group, HUUA cells of the Mifepristone + PI3K/AKT group had a lower cell growth inhibition rate and percentage of apoptotic cells (P < 0.01).

Keywords: Mifepristone, HUUA cells, proliferation, FAK, PI3K/AKT signaling pathway

Introduction
Endometrial carcinoma is one of the most common gynecological malignancies that often occurs in perimenopausal women and postmenopausal women.¹ The mortality caused by endometrial carcinoma is very high all over the world.² The incidence of endometrial carcinoma is reported to be closely related to life habits and regions.³,⁴ And unfortunately, the incidence of endometrial carcinoma has also increased year by year.⁵,⁶ Understanding of the occurrence and development mechanism of endometrial carcinoma can fundamentally improve the prevention, diagnosis and treatment of endometrial carcinoma.
Mifepristone, a common progesterone receptor antagonism agent, is widely used as an anti-pregnancy drug and also can be used during gynecological surgery operations.\textsuperscript{7,8} In recent studies, researchers have found that mifepristone also has inhibitory effects on a variety of tumor cells, such as breast cancer and prostate cancer, especially for inhibiting gynecological-related tumor cells.\textsuperscript{9-11} It could effectively inhibit the progesterone receptor contained in uterine fibroids and reduce the size of the uterine fibroids.\textsuperscript{12} Research studies have also found that mifepristone at doses of 5 mg and 10 mg daily could effectively maintain uterine fibroids treatment for 3–12 months.\textsuperscript{13,14} One explanation was that mifepristone could reduce progesterone levels as well as progesterone levels around the tumor. Therefore, it could finally achieve the effect of inhibiting the growth of uterine fibroids.\textsuperscript{15,16} Other explanations were that mifepristone could block ceramide glycosylation and promote cell apoptosis, reduce exocytosis of MDR-associated proteins and P-glycoprotein, enhance DNA repair capacity and regulate tumor suppressor genes expression.\textsuperscript{11}

There were also articles which reported the effect of mifepristone on endometrial carcinoma, while the exact mechanism has not yet been determined. In this research, we explored the effect of mifepristone on endometrial carcinoma cells proliferation, migration, invasion and apoptosis. The relevant mechanism has also been further studied. We hope this study could provide guidance for the clinical treatment of endometrial carcinoma by mifepristone.

**Material and methods**

**Cell culture and treatment**

HHUA cells were purchased from China Type Culture Collection Center. They were maintained in DMEM containing 10% FBS in a 5% CO\textsubscript{2}, 37°C incubator. These HHUA cells at logarithmic growth phase were collected and suspended in DMEM medium (10% FBS) containing 0 μmol/L, 25 μmol/L, 50 μmol/L, 75 μmol/L, and 100 μmol/L mifepristone, respectively. Then they were inoculated in 24-well plates at a density of 1×10\textsuperscript{5} cells per well. All plates were placed in the 5% CO\textsubscript{2}, 37°C incubator for continued incubation.

**Cells transfection and grouping**

pcDNA3.1-PI3K and pcDNA3.1-AKT overexpression vectors were constructed to co-transfect normal HHUA cells. These cells were treated with 100 μmol/L mifepristone and were set as the Mifepristone + PI3K/AKT group. In addition, for HHUA cells only treated with 100 μmol/L mifepristone, they were known as the Mifepristone group. Cells of these two groups were inoculated in 24-well plates at a density of 1×10\textsuperscript{5} cells per well. They were incubated in the incubator at 37°C, 5% CO\textsubscript{2}.

**Quantitative reverse transcription PCR**

HHUA cells treated with different conditions were collected after being cultured for 48 hours. Total RNA was obtained by using a Trizol reagent kit (Thermo Fisher Scientific, Waltham, MA, USA). PCR amplification reaction was performed in a 20 μL reaction system, including 1 μL of cDNA, 1 μL of forward primer and 1 μL of reverse primer in the system. The sequences of primers used in this research were as follows: FAK, forward, GAGCGTCTAATCCGACAG CAACAG, reverse, GCCCGTCACATTCTCGTACACCT; Caspase-3, forward, GTGGAATTGATCCGTGATG, reverse, GGAATCTGTTTCTTGGCATG; PI3K, forward, GGACCCGATGCGGTTAGA, reverse, GATGATG TGCAGGGAGGC; AKT, forward, ATGGCAACCTTCTATT GGCCTAC, reverse, GGGCGGACTCGTCACTAC; GAPDH, forward, GTCGATGGCTAGTCGTAGCATCGAT, reverse, TGCTAGCTGGCATGCCCGATGCATC. The cycling conditions were as follows: For FAK: 38 cycles of 95°C for 10 seconds, 58°C for 20 seconds and 72°C for 34 seconds. For Caspase-3: 50 cycles of 95°C for 10 seconds, 58°C for 20 seconds and 72°C for 34 seconds. For PI3K: 40 cycles of 95°C for 10 seconds, 58°C for 20 seconds and 72°C for 34 seconds. For AKT: 40 cycles of 95°C for 10 seconds, 58°C for 20 seconds and 72°C for 34 seconds. For GAPDH: 42 cycles of 95°C for 10 seconds, 58°C for 20 seconds and 72°C for 34 seconds. The relative gene expression was calculated by the comparative Ct formula $-\Delta\Delta$Ct.

**Western blot**

After 48 hours of incubation, HHUA cells were collected and lysed in RIPA buffer (Beyotime, Beijing, China) to extract total protein. A Pierce BCA kit (Thermo Fisher Scientific, Waltham, MA, USA) was used to determine protein concentration. Then equal cell lysate (30 μg) from each sample was separated by 10% SDS-PAGE gel. Protein was then transferred onto PVDF membrane. The membrane was blocked with 5% fat-free milk in TBS-T for 2 hours at room temperature. Primary antibody (mouse anti-human FAK and p-FAK antibody, mouse anti-rabbit Caspase-3, PI3K, AKT, p-PI3K and p-AKT antibody, 1:1,000, Santa Cruz Biotechnology, Santa Cruz, CA, USA) was added for 12 hours incubation at 4°C. Secondary antibody (Beijing Zhongshan Jingiao Biotechnology Co., Ltd., China, 1:5,000) was also added for an extra 1 hour of incubation. After the membrane...
was washed three times with TBST, the enhanced chemiluminescence kit was used to identify the reactive bands.

**MTT assay**

In this study, proliferative capacity of HHUA cells was determined using MTT assay. Cells were cultured for 24 hours, 48 hours, 72 hours, and 96 hours, respectively, in 24-well plates. A total of 20 μL MTT solution (5 mg/mL) was added into each well for 4 hours incubation at 37°C. The residual liquid in each well was discarded. DMSO at a dose of 150 μL was added into each well. Ten min shaking was performed to promote the complete dissolution of purple formazan crystals. At last, these 24-well plates were subjected to absorbance measurements by ELISA at 495 nm wavelength (OD495 value). The cell growth inhibition rate was calculated according to the following formula: cell growth inhibition rate=-(OD495 value of treatment group/OD495 value of control group)×100%. In this research, HHUA cells treated with 0 μmol/L mifepristone were used as a control group.

**Wound-healing assay**

HHUA cells were prepared as cell suspensions at a density of 2x10^4 cells/mL. Six-well plates with 1 mL cell suspensions per well were incubated in the incubator for 12 hours at 37°C, 5% CO₂. A scratch was made by using a 100 μL sterile pipette tip. Then the residual liquid in each well was removed and 1 mL DMEM (10% FBS) was added. Six-well plates were placed under an inverted microscope and photographed. All 6-well plates were returned to the incubator for 48 hours incubation at 37°C, 5% CO₂. Afterward, they were placed under the inverted microscope again for observation and photographing. The wound-healing rate was calculated according to the following formula: (initial scratch width−scratch width at 48 hours)×100%/initial scratch width.

**Transwell assay**

Cells were harvested and prepared as serum-free cell suspensions after 48 hours incubation in 24-well plates. They were seeded onto the upper chamber of Transwell at a density of 2x10^4 cells per well. DMEM containing 10% FBS was then added into the lower chamber of Transwell. After 24 hours incubation in the 5% CO₂, 37°C incubator, the penetrated cells were fixed with formaldehyde for 5 minutes. Crystal violet was used to dye for 10 minutes. Cells were placed under an inverted microscope. Five fields were randomly selected to count the number of cells that passed through the membrane. The number of cells that passed through the membrane was defined as migrating cell numbers.

Furthermore, invasion ability was detected using the same operating procedure. However, the membrane on the upper chamber was precoated with Matrigel. Also, the number of cells that passed through the membrane was considered as the number of invading cells.

**Apoptosis detection by flow cytometry**

Cells were collected after 48 hours incubation and fixed with pre-cooled 70% ethanol for 2 hours at 4°C. Then 0.2 mL of PI solution (50 mg/mL) was added for 30 minutes incubation at 4°C in darkness. Flow cytometry was used to detect apoptosis and apoptosis percentage was also analyzed.

**Statistical analysis**

All data were expressed as mean±SD. Student t-test was used for the comparison between two groups. Comparison among three or more groups was detected by one-way ANOVA test. SPSS 17.0 and GraphPad Prism 5.0 were used for statistical analysis of data. P<0.05 was considered statistically significant. In this research, all experiments were repeated three times.

**Results**

**Mifepristone inhibited HHUA cells proliferation in a dose-dependent manner**

This study examined the effects of different concentrations of mifepristone on HHUA cells proliferation by MTT assay. The results demonstrated that mifepristone inhibited HHUA cells proliferation in a dose-dependent manner. The cell growth inhibition rate was gradually increased with the increase of mifepristone concentration. When the concentration of mifepristone was at 100 μmol/L, cell growth inhibition rate was the highest (Figure 1).

**Figure 1** Mifepristone inhibited HHUA cells proliferation in a dose-dependent manner.
Mifepristone inhibited HHUA cells migration and invasion in a dose-dependent manner

Cell migration ability was detected by wound-healing assay and Transwell assay, respectively. According to the results, we noted that, for HHUA cells treated with 50–100 μmol/L mifepristone, their wound-healing rate and number of migrating cells was both markedly decreased when compared with HHUA cells treated with 0 μmol/L mifepristone (P<0.01) (Figure 2A and B). We also noticed that, compared with HUA cells treated with 0 μmol/L mifepristone, those treated with 50–100 μmol/L mifepristone had dramatically lower invasive cell numbers (P<0.01) (Figure 2C).

Mifepristone promoted HHUA cells apoptosis in a dose-dependent manner

Mifepristone promoted apoptosis of HHUA cells in a dose-dependent manner. Compared with the percentage of apoptotic HHUA cells treated with 0 μmol/L mifepristone, significantly increased apoptotic cells percentage was found when they were treated with 25–100 μmol/L mifepristone (P<0.01) (Figure 3A). Meanwhile, we also found that mifepristone could promote Caspase-3 expression in a dose-dependent manner. Caspase-3 mRNA and protein were both significantly increased after they were treated by 50–100 μmol/L of mifepristone (P<0.01) (Figure 3B and C).

Mifepristone inhibited FAK and p-FAK expression in a dose-dependent manner

We further explored the effect of mifepristone on the FAK pathway in HHUA cells. The results revealed that mifepristone could inhibit FAK mRNA and protein relative expression in a dose-dependent manner. When HHUA cells were treated with 50–100 μmol/L of mifepristone, FAK mRNA and protein relative expression was significantly lower than

![Figure 2](image-url)

**Figure 2** Mifepristone inhibited HHUA cells migration and invasion in a dose-dependent manner.

**Notes:** (A) Detection of wound-healing rate by wound-healing assay. (B) HHUA cells migration assay by Transwell. (C) HHUA cells invasion detection by Transwell. **P<0.01** when compared HHUA cells treated with 0 μmol/L mifepristone.
those treated with 0 μmol/L of mifepristone (P<0.01). Similar trend was also found in p-FAK expression (Figure 4A and B). These results indicated that mifepristone could inhibit the activity of the FAK pathway in HHUA cells in a dose-dependent manner.

**Figure 3** Mifepristone promoted apoptosis of HHUA cells in a dose-dependent manner.

**Notes:** (A) Detection of HHUA cells apoptosis by flow cytometry; (B) Detection of Caspase-3 mRNA relative expression in HHUA cells by qRT-PCR; (C) Detection of Caspase-3 protein relative expression in HHUA cells by Western blot. **P<0.01 when compared HHUA cells treated with 0 μmol/L mifepristone.

**Abbreviation:** qRT-PCR, quantitative reverse transcription PCR.

Mifepristone inhibited p-PI3K and p-AKT expression in a dose-dependent manner

PI3K/AKT signaling pathway has shown to be involved in tumorigenesis and development. Our findings indicated that different concentrations of mifepristone did not obviously affect PI3K and AKT expression. However, HHUA cells treated with 50–100 μmol/L mifepristone were with significantly lower p-PI3K and p-AKT expression than those treated with 0 μmol/L mifepristone (P<0.01) (Figure 5A–D). The results indicated that, in HHUA cells, mifepristone could inhibit p-PI3K and p-AKT expression in a dose-dependent manner.

Mifepristone inhibited HHUA cells proliferation and promoted its apoptosis by suppressing PI3K/AKT signaling pathway activity

We further investigated whether mifepristone affected HHUA cells proliferation and apoptosis by affecting PI3K/AKT signaling pathway activity. Compared with Mifepristone group, the relative expression of PI3K, p-PI3K, AKT and p-AKT was all significantly upregulated in HHUA cells of Mifepristone + PI3K/AKT group (P<0.01) (Figure 6A). Meanwhile, the cell growth inhibition rate and percentage of apoptotic cells in HHUA cells of Mifepristone + PI3K/AKT
group was both markedly lower than those of Mifepristone group \( (P<0.01) \) (Figure 6B and C).

**Discussion**

The present study investigated the effects of mifepristone on HHUA cells. The results indicated that mifepristone could inhibit HHUA cells proliferation, migration and invasion and promote its apoptosis in a dose-dependent manner. The mechanism of mifepristone on affecting HHUA cells was through inhibiting the activity of FAK pathway and PI3K/AKT signaling pathway.

Previous researches have proven that mifepristone could exert an anti-tumor effect, which could selectively inhibit the formation and development of multiple tumors.\(^{17-19}\) Liu et al\(^{20}\) proved that mifepristone could be used as a therapeutic drug for treating triple negative breast cancer. By in vivo and in vitro studies, they found that mifepristone significantly inhibited triple negative breast cancer cells proliferation and promote its apoptosis. The mechanism was through down-regulation of KLF5. It is well known that KLF5 is a key transcription factor, which is able to maintain triple negative breast cancer cancer stem cells.\(^{21,22}\) Jurado et al\(^{23}\) discovered

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**Figure 4** Mifepristone inhibited FAK and p-FAK expression in a dose-dependent manner.

**Notes:** (A) Detection of FAK mRNA expression by qRT-PCR. (B) Detection of FAK and p-FAK protein expression by Western blot. **\( P<0.01 \) when compared HHUA cells treated with 0 \( \mu \)mol/L mifepristone.

**Abbreviation:** qRT-PCR, quantitative reverse transcription PCR.

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**Figure 5** Mifepristone inhibited FAK and p-FAK expression in a dose-dependent manner.

**Notes:** (A) Detection of FAK mRNA expression by qRT-PCR. (B) Detection of FAK and p-FAK protein expression by Western blot. **\( P<0.01 \) when compared HHUA cells treated with 0 \( \mu \)mol/L mifepristone.

**Abbreviation:** qRT-PCR, quantitative reverse transcription PCR.
that mifepristone could greatly improve the therapeutic effect of cisplatin on cervical cancer. They recommend that anti-hormonal drugs combined with anti-cancer drugs could be used in the treatment of cervical cancer as well as other cancers. However, in-depth studies of the relevant mechanisms have not been conducted. Lu et al\textsuperscript{24} used different concentrations of mifepristone to investigate its effect on human endometrial carcinoma cells migration. They found that mifepristone could inhibit H19 transcriptional levels by promoting the methylation of the H19 promoter, which eventually resulted in the up-regulation of E-cadherin expression and led to an inhibitory effect on the migration of human endometrial carcinoma cells.

Further studies indicated that mifepristone suppressed the activity of the FAK pathway in HHUA cells. It could inhibit the expression of FAK as well as p-FAK in a dose-dependent manner. FAK, a regulator of focal adhesion dynamics, is involved in signal transduction events between cells and their extracellular matrix.\textsuperscript{25,26} FAK plays a significant role in adhesion and cooperative signaling of tumor cells’ growth factors. It increases tumor cells’ motility, invasiveness and proliferation, and viability.\textsuperscript{27} FAK has been found to be up-regulated in a variety of tumors, including breast cancer, pancreatic neuroendocrine tumors and gastric cancer.\textsuperscript{28–30} Inhibition of FAK phosphorylation could significantly inhibit tumor cell growth and metastasis.\textsuperscript{31} There were also studies demonstrated that high expression of FAK was found in both endometrial hyperplasia and endometrial carcinoma. It was participated in epithelial-mesenchymal transition and migration during the development and progression of endometrial carcinoma.\textsuperscript{32–37} Alowayed et al\textsuperscript{38} suggested that endometrial carcinoma cell proliferation and migration ability were both impaired after FAK activity was inhibited. We also validated that mifepristone inhibits the development of endometrial carcinoma by inhibiting the activity of FAK. In addition, our study also found that mifepristone could suppress the activity of PI3K/AKT signal pathway in a dose-dependent manner. It impaired HHUA cell proliferation by suppressing the phosphorylation of PI3K and AKT. Accumulated studies showed that dysregulated phosphorylation of PI3K/AKT

Figure 6 Mifepristone inhibited HHUA cells proliferation and promoted its apoptosis by suppressing PI3K/AKT signaling pathway activity.

Notes: (A) Detection of PI3K, p-PI3K, AKT and p-AKT protein expression by Western blot. (B) Detection of cell growth inhibition rate by MTT assay. (C) Detection of apoptosis by flow cytometry. **P<0.01 when compared with the Mifepristone group.
played an important role in the development of cancers.\textsuperscript{39–41} Lian et al\textsuperscript{42} thought that activated PI3K/AKT signaling pathway has a positive effect on the development of laryngeal squamous cell carcinoma. Liu et al\textsuperscript{43} illustrated that PI3K/AKT pathway could be used as a potential target of colorectal cancer. Previous research has proven that, FAK was one of the key genes regulating the PI3K/AKT signaling pathway.\textsuperscript{44} Activated FAK could phosphorylate PI3K, which further led to the phosphorylation of AKT.\textsuperscript{45} AKT participated in many tumor cellular processes and tumor development after it was activated.\textsuperscript{46–48} Therefore, we speculated that, mifepristone might suppress the activity of the PI3K/AKT signaling pathway by impairing the activation of FAK. This mechanism will be one of the focuses of our future research.

Conclusions
This research demonstrated that mifepristone inhibited HUUA cells proliferation, migration, invasion and promoted its apoptosis by inhibition of the FAK and PI3K/AKT signaling pathways. It provided a new theoretical basis for the treatment of endometrial carcinoma with mifepristone, and also provided a new potential target for the treatment of endometrial carcinoma.

Author contributions
Lin Sang: Substantial contributions to conception and design, data acquisition; Dawei Lu: Substantial contributions to conception and design, data analysis and interpretation; Jun Zhang: Drafting the article or critically revising it for important intellectual content; Shihua Du: Drafting the article or critically revising it for important intellectual content; Xingbo Zhao: Final approval of the version to be published. All authors contributed toward data analysis, drafting and critically revising the paper and agree to be accountable for all aspects of the work.

Disclosure
The authors report no conflicts of interest in this work.

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