

# Long noncoding RNA PCAT-1 promoted ovarian cancer cell proliferation and invasion by suppressing KLF6

H.-P. LIU<sup>1</sup>, D. LV<sup>2</sup>, J.-Y. WANG<sup>1</sup>, Y. ZHANG<sup>1</sup>, J.-F. CHANG<sup>1</sup>, Z.-T. LIU<sup>1</sup>, N. TANG<sup>1</sup>

<sup>1</sup>Reproductive Medicine Center, The 960<sup>th</sup> Hospital of the PLA Joint Logistics Support Force, Chengde, China

<sup>2</sup>Department of Pain Management, Tianjin First Center Hospital, Tianjin, China

**Abstract. – OBJECTIVE:** Recently, the role of long noncoding RNAs (lncRNAs) in tumor progression has caught attention in numerous researchers. In our investigation, lncRNA PCAT-1 was studied to identify how it functioned in the progression of ovarian cancer.

**PATIENTS AND METHODS:** LncRNA PCAT-1 expression was detected by quantitative Real-time polymerase chain reaction (qRT-PCR) in both ovarian cancer cells and tissue samples. Furthermore, to identify the function of PCAT-1 in ovarian cancer, cell proliferation, transwell assay and Matrigel assay were conducted. In addition, by performing qRT-PCR and Western blot assay, the underlying mechanism was explored.

**RESULTS:** PCAT-1 expression was remarkably higher in ovarian cancer samples when compared with that in corresponding normal tissues. Moreover, cell proliferation, migration and invasion were inhibited after PCAT-1 was knocked down in ovarian cancer cells. In addition, the mRNA and protein expression of KLF6 (Krüppel-like factor 6) was upregulated after PCAT-1 was knocked down. Furthermore, the KLF6 expression level was negatively correlated to the PCAT-1 expression level in ovarian cancer tissue samples.

**CONCLUSIONS:** We found that PCAT-1 promotes the progression of ovarian cancer through enhancing cell metastasis and proliferation via suppressing KLF6, which might be a novel therapeutic strategy in ovarian cancer.

**Key Words:** Long noncoding RNA, PCAT-1, Ovarian cancer,

## Introduction

Ovarian carcinoma is the second most general gynecologic malignancy, which has the highest mortality accounting for 5-6% cancer-related deaths among the female. Almost 22,500 new

patients were diagnosed with ovarian cancer and 14,100 ovarian cancer patients died in America in 2017<sup>1</sup>. The most patients with ovarian cancer are usually diagnosed at advanced stage. Though surgery and chemotherapy have been widely utilized as standard therapeutic strategies for ovarian cancer, 80% of these patients used to develop resistance to therapies and metastasis, leading eventually to the patients' death<sup>2,3</sup>. This severe situation underscores the urgency of early detection and new treatment for the patients with ovarian cancer.

Long non-coding RNAs (lncRNAs) are a cluster of non-coding transcripts longer than 200 nucleotides. However, evidence has proved that lncRNAs play an important role in a variety of biological behaviors, including the tumorigenesis. For example, by regulating the expression of miR-335, lncRNA MSTO2P facilitates cell proliferation and colony formation in gastric cancer indirectly<sup>4</sup>. LncRNA HULC is overexpressed in gastric cancer that can predict prognosis of the patients and serve as a potential serum marker for diagnosis<sup>5</sup>. LncRNA APOC1P1-3 represses cell apoptosis in breast cancer through inhibiting the process of alpha-tubulin acetylation<sup>6</sup>. Moreover, by regulation of AKT-MDM2-p53 signaling axis, lncRNA LOC572558 depresses cell proliferation and tumor growth in bladder cancer<sup>7</sup>. However, how lncRNA PCAT-1 functions in the proliferation of ovarian cancer and the underlying mechanism remain unexplored. In our study, we found out that PCAT-1 was remarkably upregulated in ovarian cancer tissues. Moreover, PCAT-1 enhances cell proliferation and metastasis in ovarian cancer cell *in vitro*. Moreover, our further experiments explored the potential underlying mechanism how PCAT-1 functioned in ovarian cancer development.

## Patients and Methods

### Tissue Specimens

Paired ovarian carcinoma were sequentially-enrolled from 52 ovarian carcinoma patients undergoing surgery between July 2016 and December 2017 in the 960th Hospital of the PLA Joint Logistics Support Force. The Ethics Committee of the 960th Hospital of the PLA Joint Logistics Support Force approved this study protocol, and all participants involved in this study given a written informed consent.

### Cell Culture

Human ovarian cancer cell lines A2780, TO-V112D, OVCAR-3 and SKOV3 and normal ovarian cell ISOE80 were maintained in the culture medium consisted of 10% fetal bovine serum (FBS; Gibco, Waltham, MA, USA), penicillin as well as Dulbecco's Modified Eagle's Medium (DMEM; Gibco, Waltham, MA, USA) at 37°C in a humidified atmosphere with 5% CO<sub>2</sub>.

### Lentivirus Expressing Short-Hairpin RNA and Cell Transfection

Lentivirus expressing short-hairpin RNA (shRNA) directed against PCAT-1 was produced by GenePharma (Shanghai, China). The complementary DNA encoding PCAT-1 was amplified and inserted into pcDNA3.1 (GenePharma, Shanghai, China), which was then transfection of ovarian cancer cells with Lipofectamine 2000 (Invitrogen, Carlsbad, CA, USA). The detection of PCAT-1 expression level in these cells was conducted using quantitative real-time polymerase chain reaction (qRT-PCR).

### RNA Extraction and RT-PCR

Total RNA from tissues and cells were separated by using TRIzol reagent (Invitrogen, Carlsbad, CA, USA). The total RNA was reverse-transcribed to complementary deoxyribonucleic acids (cDNAs) through reverse transcription Kit (TaKaRa Technology, Dalian, China). Thermocycling conditions was as follows: 30 s at 95°C, 5 s for 40 cycles at 55°C, 35 s at 60°C. 2<sup>-ΔΔCt</sup> method was utilized for calculating relative expression. The primer sequences are as follows: PCAT-1 forward 5'-TGAGAAGAGAAATCTATTG-3', PCAT-1 reverse 5'-GGTTTGTC-TC-CCGCTTTA-3'; β-actin, forward 5'-GATGGAAATCGTCAGAGGCT-3' and reverse 5'-TGCACTTAGTTGGAAATGC-3'.

### Western Blot Analysis

Total proteins were collected from cells via radioimmunoprecipitation assay (RIPA) buffer and then quantified by using a protein assay reagent (bicinchoninic acid method; Beyotime, Shanghai, China). The target proteins were separated by sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE). The blots were incubated with antibodies after replacing the polyvinylidene difluoride (PVDF) membrane. Next, rabbit anti-β-actin (Cell Signaling Technology, CST, Danvers, MA, USA) and rabbit anti-KLF6 (Cell Signaling Technology, CST, Danvers, MA, USA) were used for incubation of these membranes. Pierce and Warriner chemiluminescence substrate was utilized for visualizing Western Blotting substrate Immunoreactive bands (Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA).

### Transwell Assay

24 h after transfection, 2×10<sup>5</sup> cells in 100 μl serum-free DMEM were transformed to top chamber of 96-well culture inserts (Corning, Corning, NY, USA) coated with or without 50 μg Matrigel (BD, Bedford, MA, USA). 20% serum DMEM was added to the lower chamber of the culture inserts. 24 h later, these inserts were treated by methanol for 30 min and stained by hematoxylin for 20 min. An inverted microscope (×20) was utilized for counting migrated and invaded cells in three random fields.

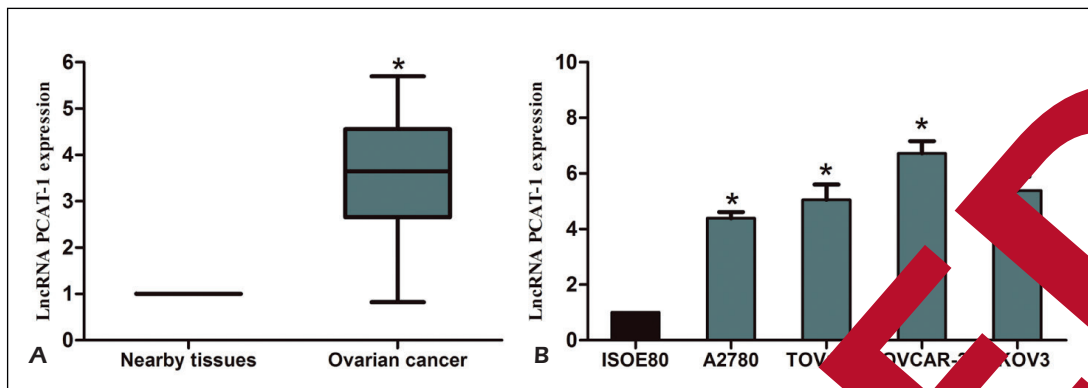
### Statistical Analysis

All statistical analyses were carried out using Statistical Product and Service Solutions (SPSS) 18.0 (SPSS Inc., Chicago, IL, USA). The differences between two groups were compared by Student *t*-test. The statistically significance was defined as *p*<0.05.

## Results

### PCAT-1 Expression Level in Ovarian Cancer Tissues and Cells

First, qRT-PCR was conducted for detecting PCAT-1 expression in 52 patients' tissues and four ovarian cancer cell lines. As a result, PCAT-1 was significantly upregulated in tumor tissue samples (Figure 1A). PCAT-1 level of ovarian cancer cells was higher than that of normal ovarian cell ISOE80 (Figure 1B).



**Figure 1.** Expression levels of PCAT-1 were increased in ovarian cancer tissues and cell lines. **A**, PCAT-1 expression was significantly increased in the ovarian cancer tissues compared with adjacent tissues. **B**, Expression levels of PCAT-1 relative to  $\beta$ -actin were determined in the human ovarian cancer cell lines and normal epithelial cell ISOE80 by qRT-PCR. Data are presented as the mean  $\pm$  standard error of the mean. \* $p < 0.05$ .

### Knockdown of PCAT-1 Inhibited Cell Proliferation in OVCAR-3 Ovarian Cancer Cell

In our study, we chose OVCAR-3 cell lines for the knockdown of PCAT-1. Then qRT-PCR was utilized for detecting the PCAT-1 expression (Figure 2A). Moreover, we conducted CCK-8 assay and found that after PCAT-1 was knocked down, the cell growth ability of OVCAR-3 cells was significantly repressed (Figure 2B).

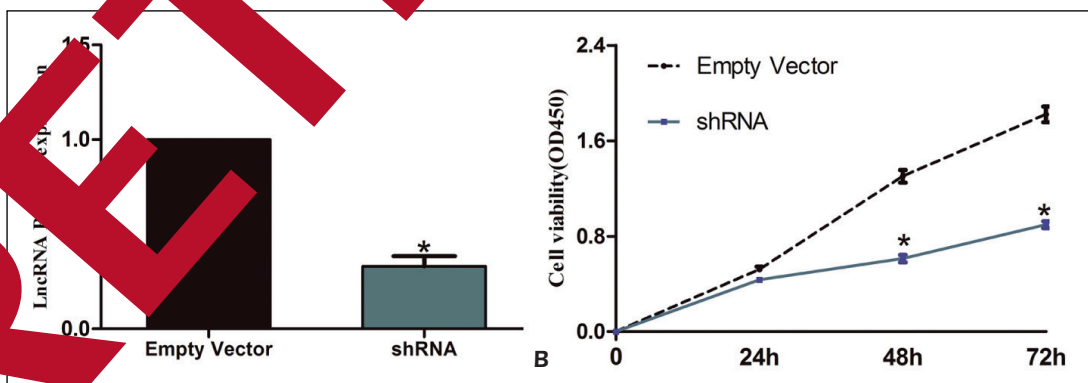
### Knockdown of PCAT-1 Inhibited Cell Migration and Invasion in OVCAR-3 Ovarian Cancer Cells

The results of transwell assay showed that after PCAT-1 was knocked down, the migrated

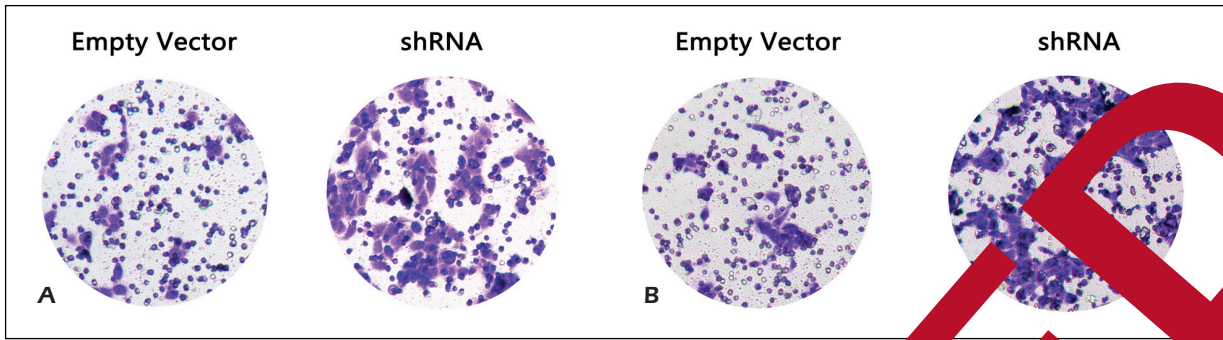
ability of ovarian cancer cells was significantly repressed (Figure 3A). The outcome of Matrigel assay also revealed that after PCAT-1 was knocked down in ovarian cancer cells, the number of invaded cells was remarkably decreased (Figure 3B).

### The Interaction Between KLF6 and PCAT-1 in OVCAR-3 Ovarian Cancer Cells

Then qRT-PCR results showed that expression level of KLF6 in ovarian cancer cells was higher in PCAT-1 shRNA (shRNA) group when compared with the KLF6 level in empty vector group (Figure 4A). Western blot assay found out that after PCAT-1 was knocked down, KLF6 could be upregulated at protein level (Figure 4B).



**Figure 2.** Knockdown of PCAT-1 inhibited OVCAR-3 ovarian cancer cell proliferation. **A**, PCAT-1 expression in OVCAR-3 ovarian cancer cells transduced with PCAT-1 shRNA (shRNA) and the empty vector was detected by qRT-PCR.  $\beta$ -actin was used as an internal control. **B**, CCK-8 assay showed that knockdown of PCAT-1 significantly inhibited cell growth in OVCAR-3 ovarian cancer cells. The results represent the average of three independent experiments (mean  $\pm$  standard error of the mean). \* $p < 0.05$ .



**Figure 3.** Knockdown of PCAT-1 inhibited OVCAR-3 ovarian cancer cell migration and invasion. **A**, Transwell assay showed that knockdown of PCAT-1 significantly decreased cell migration in OVCAR-3 ovarian cancer cells. **B**, Matrigel assay showed that number of invaded cells was significantly decreased via knockdown of PCAT-1 in OVCAR-3 ovarian cancer cells. The results represent the average of three independent experiments (mean  $\pm$  standard error of the mean).

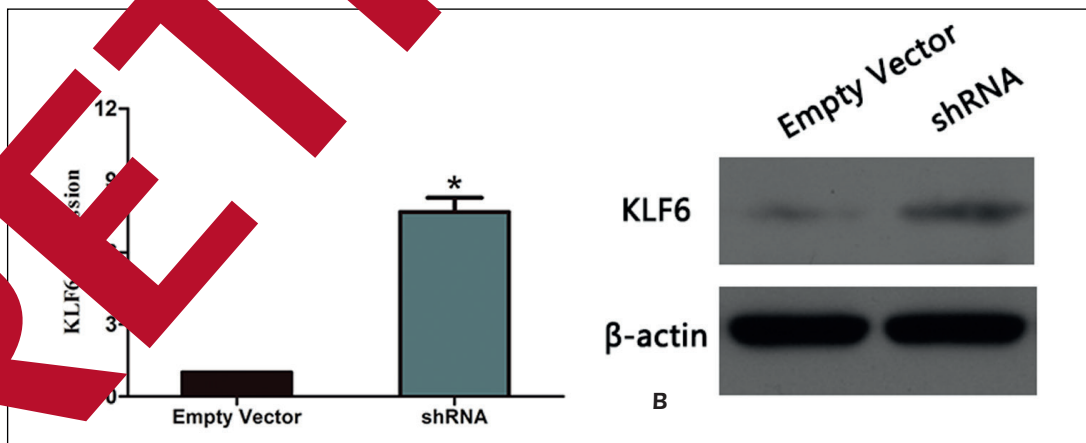
**The Interaction Between KLF6 and PCAT-1 in OVCAR-3 Ovarian Cancer Tissues**

We further found that KLF6 expression of ovarian cancer tissues was significantly lower compared with that of adjacent tissues (Figure 5A). Correlation analysis revealed that the negative association was seen between KLF6 expression level and PCAT-1 expression in ovarian cancer tissues (Figure 5B).

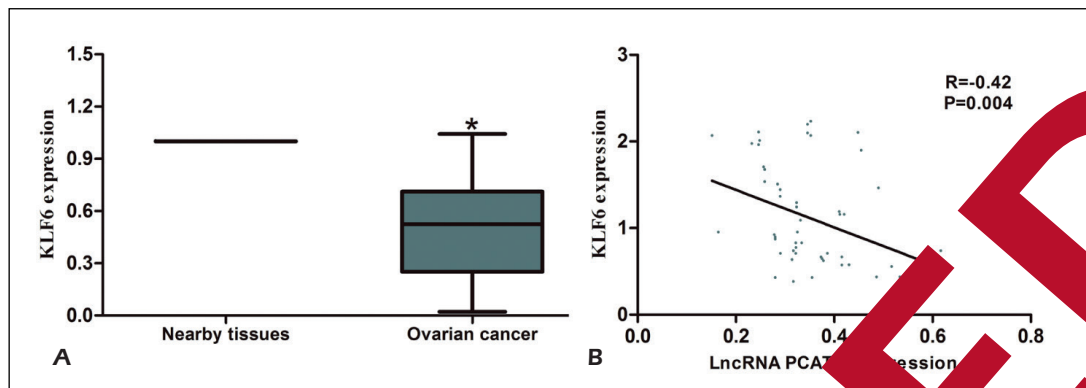
be potential indicators and therapeutic target for ovarian cancer. For instance, lncRNA AF-AS1 promotes cell proliferation and attenuates cell apoptosis in ovarian cancer<sup>8</sup>. LncRNA LINC01511 inhibits the progression of ovarian carcinoma stem cells and functions as a novel target for treating ovarian cancer<sup>9</sup>. Upregulated by oestrogen, lncRNA ElnRNA1 functions as an oncogene in the proliferation of epithelial ovarian cancer cell<sup>10</sup>. LncRNA ZFAS1 promotes cell proliferation, migration and chemoresistance in epithelial ovarian cancer by interacting with miR-150-5<sup>11</sup>. In addition, downregulation of lncRNA SPRY4-IT1 enhances cell metastasis in ovarian cancer by partly regulating epithelial-mesenchymal transition<sup>12</sup>.

**Discussion**

Evidence proved that lncRNAs participate in the development of ovarian cancer and could



**Figure 4.** Interaction between PCAT-1 and KLF6 in OVCAR-3 ovarian cancer cells. **A**, qRT-PCR results showed that KLF6 expression was higher in PCAT-1 shRNA (shRNA) compared with the empty vector. **B**, Western blot assay revealed that KLF6 protein expression was increased in PCAT-1 shRNA (shRNA) compared with empty vector. The results represent the average of three independent experiments. Data are presented as the mean  $\pm$  standard error of the mean. \* $p$ <0.05.



**Figure 5.** Interaction between PCAT-1 and KLF6 in ovarian cancer tissues. **A**, KLF6 was significantly downregulated in ovarian cancer tissues compared with adjacent tissues. **B**, The linear correlation between the expression levels of KLF6 and PCAT-1 in ovarian cancer tissues. The results represent the average of three independent experiments. Data are presented as the mean  $\pm$  standard error of the mean. \* $p < 0.05$ .

LncRNA PCAT-1 (Prostate cancer-associated ncRNA transcripts 1) was initially discovered in prostate cancer<sup>13</sup>. PCAT-1 was highly expressed in bladder cancer and might be a novel therapeutic target<sup>14</sup>. Overexpressed PCAT-1 promoted cell proliferation and cell metastasis in non-small cell lung cancer<sup>15</sup>. Bi *et al*<sup>16</sup> reported upregulating CDKN1A, PCAT-1 facilitates cell metastasis in gastric cancer. In addition, Bi *et al*<sup>17</sup> proposed that the PCAT-1 played an important role in tumorigenesis of hepatocellular carcinoma *via* TP53-miR-215-P53-KLF axis. In our research, we found that PCAT-1 was upregulated in both ovarian cancer samples and cells. Besides, after PCAT-1 knockdown, ovarian cancer cell proliferation and metastasis were found inhibited. Above results indicated that PCAT-1 promoted tumorigenesis of ovarian cancer and might act as an oncogene.

KLF6 (Krüppel-like factor 6) is widely known as a tumor suppressor in many carcinomas, which regulates diverse biological processes. For example, KLF6 constrains the progression of hepatocellular carcinoma dissemination by regulating a VAV3 signaling axis<sup>18</sup>. KLF6-E2F1 axis inactivated aggressive clear cell renal cell carcinoma and KLF6 functions as an anti-oncogene *via* transcriptional depression of E2F1<sup>19</sup>. KLF6 inhibited glioblastoma, which is related to tumor prognosis for the patients through targeting E2F1<sup>20</sup>. Moreover, as a target of miR-630, KLF6 constrains cell proliferation and cell invasion in epithelial ovarian cancer<sup>21</sup>. In the present study, KLF6 expression could be upregulated after knockdown of PCAT-1. Moreover, KLF6 expression in ovarian

cancer tissues was negatively related to PCAT-1 expression. All the results above suggested that PCAT-1 might promote tumorigenesis of ovarian cancer *via* suppressing KLF6.

## Conclusions

We identified that PCAT-1 was remarkably higher-expressed in ovarian cancer tissues and cells. Besides, PCAT-1 could enhance ovarian cancer cell migration and invasion through targeting KLF6. These findings suggest that PCAT-1 may contribute to therapy for ovarian cancer as a candidate target.

## Conflict of Interests

The authors declared no conflict of interest.

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