

LncRNA FEZF1-AS1 promoted chemoresistance, autophagy and epithelial-mesenchymal transition (EMT) through regulation of miR-25-3p/ITGB8 axis in prostate cancer

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Abstract. – **OBJECTIVE:** Accumulating evidence determined that lncRNA plays important roles in the development and occurrence of cancers. Prostate cancer is the second most common type of cancer and one of the top five cancers for the cause of male death in the world. Therefore, this study was to explore the regulatory mechanism of lncRNA in chemoresistance of PC.

MATERIALS AND METHODS: qRT-PCR was used to detect the mRNA expression of FEZF1-AS1, miR-25-3p and ITGB8. Western blot was applied to measure the protein expression of ITGB8, E-cadherin, N-cadherin, Vimentin, LC3II, ATG5 and Beclin-1. In addition, CCK-8 assay was used to assess proliferation of transfected cells. Luciferase reporter assay and RIP assay were used to determine the relationship among FEZF1-AS1, miR-25-3p and ITGB8.

RESULTS: In this study, the expression of FEZF1-AS1 and ITGB8 was up-regulated, whereas the expression of miR-25-3p was down-regulated in PC tumor tissues and PC/PTX cells. Luciferase reporter assay and RIP assay determined that miR-25-3p was a target of FEZF1-AS1 and ITGB8 was a target of miR-25-3p. Interestingly, knockdown of FEZF1-AS1 could inhibit cell viability and EMT and promoted cell autophagy in PC/PTX cells, but inhibition of miR-25-3p or promotion of ITGB8 could reverse the effect of FEZF1-AS1 in PC/PTX cells.

CONCLUSIONS: In this study, we found that lncRNA FEZF1-AS1 promoted chemoresistance, autophagy and epithelial-mesenchymal transition (EMT) through regulation of miR-25-3p/ITGB8 axis in PC, providing a new regulatory mechanism in PC and a novel therapeutic target.

Keywords:

FEZF1-AS1, Chemoresistance, Autophagy, EMT, Prostate cancer.

Introduction

Prostate cancer (PC) is an epithelial malignancy that occurs in the prostate, which is the second most common type of cancer and one of the top five cancers for the cause of male death in the world. Chemoresistance was the main problem in the treatment of PC. While alleviating the development of PC, drugs also increase the chemoresistance of cancer cells to drugs and reduce the effect of drug treatment on cancers². Paclitaxel (PTX) is a natural anti-cancer drug that has a significant alleviated effect on PC, but chemoresistance is also occurred³⁻⁶. However, chemoresistance studies for PC cells are still very limited.

LncRNA is a type of non-coding RNA of more than 200 bp in length⁷. LncRNAs play an important regulatory role in the treatment and diagnosis of cancer, and its regulation mechanism has also improved the understanding of cancer occurrence and development⁸⁻¹⁰. It is well known that lncRNA affects the expression of mRNA by binding to its target miRNA, thereby affecting a series of physiological metabolism, immunity, and cell cycle processes¹¹⁻¹⁴, including PC¹⁵. Such as lncRNA PCAT-1 could contribute to cell proliferation *via* regulating cMyc in PC¹⁶. Growing evidence determined that FEZF1-AS1 played multiple roles in cancers, including colorectal carcinoma, gastric cancer, osteosarcoma and PC¹⁷⁻²⁰. In this study, we found that FEZF1-AS1 is highly expressed in PC, but its role in drug resistance is not fully understood.

MiRNAs are also important regulators of the development, progression, chemoresistance of cancer cells^{21,22}. In PC, miR-29b has been shown to

be involved in cell proliferation, metastasis and EMT^{23,24}. With the development of sequencing technology, more and more miRNAs involved in cancer cell progression have been discovered, and miR-25-3p is one of them^{25,26}. miR-25-3p is a tumor suppressor that is low in cancers and inhibits the development and progression of cancer cells^{27,28}. However, research on the mechanism of drug resistance in PC cells needs further exploration.

Integrin beta-8 (ITGB8), a member of the integrin beta chain family, encodes a single-pass type I membrane protein containing a VWFA domain and four cysteine-rich repeats²⁹. ITGB8 is shown to be involved in cancer cell invasion, metastasis and chemoresistance^{30,31}, but its function in PC has not been fully explored.

In this paper, we focus on the functional study of lncRNA resistance in PC cells, further exploring its molecular regulation mechanism in drug resistance, improving the understanding of cancer cell resistance mechanism.

Materials and Methods

Patients, Tissues and Serum Samples

A total of 47 PC tissues and matched adjacent normal tissues, and PC serum samples were obtained from the West China Hospital, Sichuan University with the informed consent of patients. This study was approved by the Institutional Review Board of the West China Hospital, Sichuan University. All patients were diagnosed as PC and had not undergone chemotherapy or surgery before specimen collection. Normal serum samples were collected from 42 patients who were not diagnosed with PC. All tissues and serum were stored at -80°C.

Cell Culture and Transfection

PC cell lines (DU145 and LNCaP) were purchased from the American Type Culture Collection (ATCC; Manassas, VA, USA). DU145/PTX and LNCaP/PTX cells were obtained from DU145 and LNCaP treatment with paclitaxel. All cell lines were seeded in six-well plates and cultured in Dulbecco's modified Eagle's medium (DMEM; Life Technologies, Gaithersburg, MD, USA) contained with 10% fetal bovine serum (FBS; Atlanta Biologicals, Lawrenceville, GA, USA) at 37°C in a humidified atmosphere with 5% CO₂.

All plasmids and oligos, Si-FE2F1-AS1, si-ITGB8, anti-miR-25-3p, pcDNA-ITGB8 (ITGB8),

miR-25-3p mimic (miR-25-3p) and their negative control, were purchased from Ribobio (Guangzhou, China). A negative control siRNA (si-NC, R, Life Technologies) was also utilized. Reaching up to 70% confluence, all plasmids and oligos were transfected into DU145/PTX and LNCaP/PTX using Lipofectamine 2000 (Invitrogen-Fisher Scientific, Waltham, MA, USA) according to the manufacturer's instructions.

qRT-PCR

Total RNA was extracted from all tissues and cells by using TRIzol reagent (Invitrogen, Carlsbad, CA, USA) following the manufacturer's instructions. For miR-25-3p, RNA was transcribed to cDNA using the TaqMan microRNA Reverse Transcription Kit (Applied Biosystems, Foster City, CA, USA). For FE2F1 or ITGB8, RNA was transcribed to cDNA by using M-MLV Reverse Transcriptase (Invitrogen). U6 and Glyceraldehyde phosphate dehydrogenase (GAPDH) were used as reference genes for miR-25-3p and FE2F1 or ITGB8, respectively. SYBR Green Real-time PCR (Invitrogen) was used for quantitative RT-PCR, according to the manufacturer's protocol. Each sample had three replications. Relative expression levels were calculated and analyzed using the 2^{-ΔΔCt} method. The primers are as follows: U6-F 5'-CTCGCTTCGGCAGCAC-3'; U6-R 5'-AACGCTTCACGAATTTGCGT-3'; miR-25-3p-F 5'-CATTGCACTTGTCTCGGTCTGA-3'; miR-25-3p-R 5'-GCTGTCAACGATACGCTACGTAACG-3'; ITGB8-F 5'-CCCGTGACTTTCGTCTTGGA-3'; ITGB8-R 5'-CCTTTCGGGTGGATGCTAA-3'; FE2F1-F 5'-TTAGGAGCCTTGTTCTGTGT-3'; FE2F1-R 5'-GCGCATGTACTTAAGAAAGA-3'; GAPDH-F 5'-GGAGCGAGATCCCTCCAAAAT-3'; GAPDH-R 5'-GGCTGTTGTCATACTTCTCATGG-3'.

Western Blot

Cell samples were washed with Tris-buffered saline (TBS) and lysed using a lysis buffer (Beyotime, Shanghai, China) on ice for protein extraction. Then, a BCA protein assay kit (Beyotime) was used to detect the protein concentration. Subsequently, 50 µg of proteins were added onto 10% sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) to separation and then transferred to polyvinylidene difluoride (PVDF) membranes. The membranes were incubated with primary antibodies, including anti-ITGB8 anti-E-cadherin, anti-N-cadherin, anti-Vimentin, anti-LC3I, anti-LC3II, anti-ATG5,

anti-Beclin-1 and anti-GAPDH (1:2000, Cell Signaling Technology, Danvers, MA, USA). After that, the membranes were incubated with horseradish peroxidase (HRP)-conjugated secondary antibody solution (goat anti-mouse IgG, 1:2000 dilution, Cell Signaling Technology). The ECL Western blotting substrate (Promega, Madison, WI, USA) was applied to detect and visualize the blot signals.

Cell Viability

CCK-8 Kit (Dojindo Molecular Technologies, Kumamoto, Japan) was used to measure the cell viability according to the manufacturer's protocol. Briefly, all cells were seeded into 96-well plates at a density of 5000 cells/cm². Next, each well was added into 10 μ L CCK-8 solutions and incubated at 37°C for 0 h, 24 h, 48 h, and 72 h. The absorbance was detected at 450 nm using a microplate reader (Bio-Rad, Hercules, CA, USA) at 0 h, 24 h, 48 h, 72 h. DU145, LNCaP, DU145/PTX and LNCaP/PTX cells were treated at concentrations of 0, 2, 4, 8, 16 paclitaxel for 24 h, and the half-maximal inhibitory concentration IC₅₀ was calculated as the concentration of paclitaxel that increased cell viability by 50% using sigmoidal dose-responses curves (IC₅₀ Calculator software; AAT Bioquest, Sunnyvale, CA, USA).

Luciferase Reporter Assay

Wild-type and mutated FEZF1-AS1 3'-UTRs (FEZF1-AS1-WT and FEZF1-AS1-MUT) or ITGB8 3'-UTRs (ITGB8-3'UTR-WT and ITGB8-3'UTR-MUT) were inserted into the luciferase reporter vector (Promega, Madison, WI, USA). The wild-type or mutated FEZF1-AS1 3'-UTR-ITGB8 3'-UTRs containing binding sites with miR-25-3p, the mutated or wild-type FEZF1-AS1 lacked binding sites with miR-25-3p. DU145/PTX and LNCaP/PTX cells were seeded into 24-well plates. The FEZF1-AS1-WT and FEZF1-AS1-MUT, ITGB8-3'UTR-WT and ITGB8-3'UTR-MUT were transfected with miR-125-3p or miR-NC into DU145/PTX and LNCaP/PTX cells using Lipofectamine 2000 (Thermo-Fisher Scientific, Waltham, MA, USA).

RNA Immunoprecipitation (RIP)

DU145/PTX and LNCaP/PTX cells were used as the EZ-Magna RIP™ RNA-Binding Protein Immunoprecipitation Kit (Millipore, Billerica, MA, USA) to perform RIP experiments according to the manufacturer's protocol.

Anti-Ago2 or anti-IgG antibodies were used to incubate with cells according to the manufacturer's protocol. qRT-PCR was applied to detect the expression of FEZF1-AS1.

Tumor Xenograft Model

Female 6-week-old BALB/c nude mice were purchased from Vital River Inc. (Beijing, China) and injected subcutaneously in the flank regions with 2 $\times$ 10⁴ DU145/PTX or LNCaP/PTX cells stably transfected with sh-NC or sh-FEZF1-AS1. Tumor volumes were measured after injection for 7 d, 14 d, 21 d and 28 d. After injection for 28 d, tumor weights were calculated. Volume = (width $\times$ length $\times$ (width + length)) $\times$ 0.5. All animal experiments were approved by the Ethics Committee of West China Hospital, Sichuan University.

Statistical Analysis

All experiments were repeated three times. Detailed Student's *t*-test was used to analyze all comparisons in various groups. All data represent the mean $\pm$ standard deviation (SD) from three independent experiments. All statistical analyses were displayed and analyzed using GraphPad Prism 5.0 (GraphPad Software, San Diego, CA, USA). *P* < 0.05 was considered as statistically significant.

Results

FEZF1-AS1 Was Upregulated In PC Tumor Tissues And Associated With Paclitaxel Sensitivities In PC

As shown in Figure 1A, FEZF1-AS1 was up-regulated in PC tumor tissues compared with normal tissues. Moreover, FEZF1-AS1 is associated with PTX resistance sensitivity of PC (Figure 1B). In addition, with the increase of paclitaxel concentration, the cell viability of DU145/PTX and LNCaP/PTX cells was significantly higher than that of DU145 and LNCaP cells (Figure 1C and 1D). Additionally, the IC₅₀ of DU145/PTX and LNCaP/PTX cells were sharply higher than that of DU145 and LNCaP cells (Figure 1C and 1D). Interestingly, the expression level of FEZF1-AS1 in DU145/PTX and LNCaP/PTX cells was also significantly higher than that of DU145 and LNCaP (Figure 1E and 1F). These results indicated that abnormal expression of FEZF1-AS1 in PC is associated with paclitaxel resistance of PC cells.

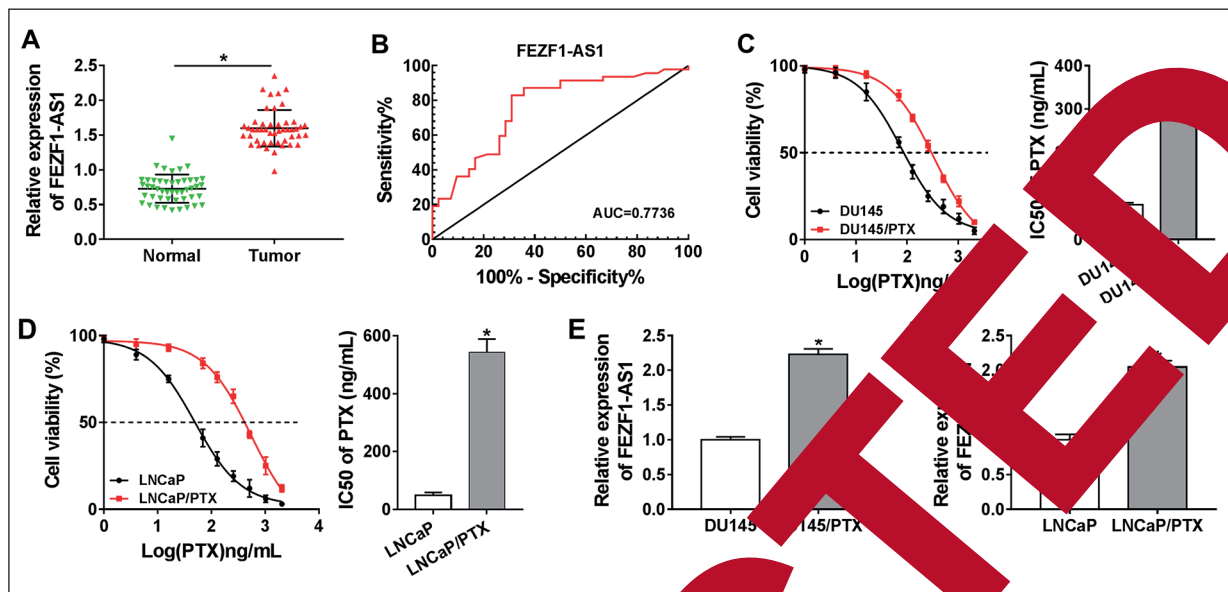


Figure 1. FEZF1-AS1 was up-regulated in PC tumor tissues and associated with paclitaxel sensitivities in PC. (A) The expression of FEZF1-AS1 in normal tissues and PC tumor tissues was detected with qPCR. (B) ROC curve analysis of the diagnostic value of FEZF1-AS1 for drug resistance in PC. (C and D) The IC₅₀ values of DU145/PTX, DU145, LNCaP/PTX and LNCaP cells treated with paclitaxel of different concentrations (0, 2, 4, 6 ng/mL) were detected. (E and F) The expression of FEZF1-AS1 in DU145/PTX, DU145, LNCaP/PTX and LNCaP cells was detected using RT-PCR. **p* < 0.05.

Knockdown of FEZF1-AS1 Could Inhibit Cell Viability and EMT and Promoted Cell Autophagy in PC/PTX Cells

To further clarify the function of FEZF1-AS1 in PC resistance, DU145/PTX and LNCaP/PTX cells transfected with si-NC, si-FEZF1-AS1 and si-FEZF1-AS1 transfection efficiency was detected. The expression of FEZF1-AS1 in DU145/PTX and LNCaP/PTX cells (Figure 2A). In addition, we also found that decreased expression of FEZF1-AS1 significantly reduced the IC₅₀ of DU145/PTX and LNCaP/PTX cells (Figure 2C and 2D). Moreover, cell proliferation of si-FEZF1-AS1 group was significantly reduced compared to si-NC group in DU145/PTX and LNCaP/PTX cells (Figure 2D and 2E). LC3II, LC3II, ATG5 and Beclin-1 are important regulators of autophagy in cells which reflect the autophagy state of cells. As shown in Figure 2H, the protein expression of LC3II/LC3I, ATG5 and Beclin-1 were significantly inhibited by si-FEZF1-AS1 transfection in DU145/PTX and LNCaP/PTX cells. In addition, FEZF1-AS1 also affected the EMT process of DU145/PTX and LNCaP/PTX cells. E-cadherin, N-cadherin and Vimentin have proven to be important biomarkers for EMT. In this paper, a decreased FEZF1-AS1 expression significantly

inhibited the protein expression of N-cadherin and Vimentin and promoted the expression of E-cadherin protein in DU145/PTX and LNCaP/PTX cells (Figure 2I to 2K). Therefore, all data determined that a decreased expression of FEZF1-AS1 inhibited proliferation and EMT as well as promoted autophagy in DU145/PTX and LNCaP/PTX cells, revealing that the low expression of FEZF1-AS1 contributed to reducing the paclitaxel resistance of PC cells and inhibited the growth and metastasis of PC/PTX cells.

MiR-25-3p Was a Target MiRNA of FEZF1-AS1

Target genes for FEZF1-AS1 candidates were predicted by starBase 3.0. The results showed that miR-25-3p was the target miRNA of FEZF1-AS1 (Figure 3A). To further validate the relationship between FEZF1-AS1 and miR-25-3p in DU145/PTX and LNCaP/PTX cells, analysis of the luciferase reporter assay showed that luciferase activity was significantly reduced when miR-25-3p was combined with FEZF1-AS1-WT DU145/PTX and LNCaP/PTX cells (Figure 3B and 3C). At the same time, miR-25-3p was shown to be underexpressed in PC tissues and negatively correlated with expression of FEZF1-AS1 (Figure 3D and 3E). In addition, the expression of miR-25-3p

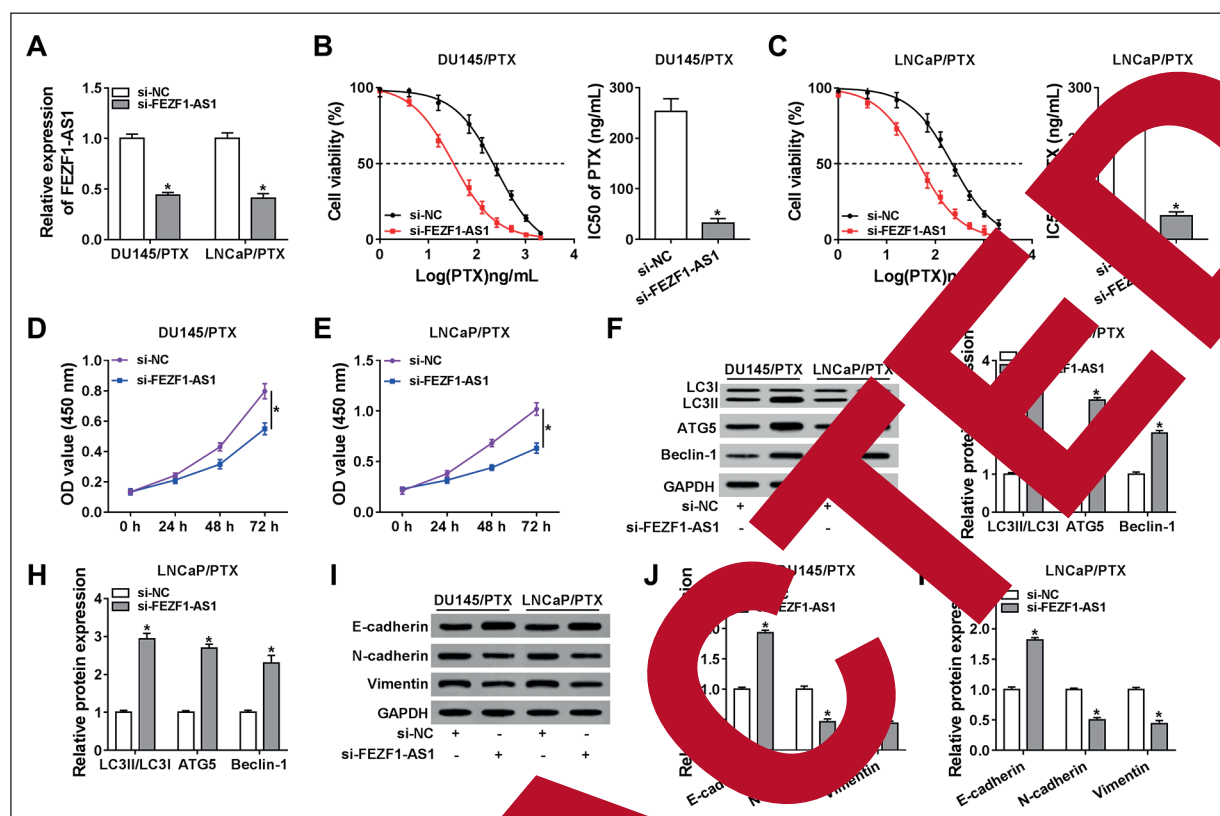


Figure 2. Knockdown of FEZF1-AS1 could inhibit cell viability and promoted cell autophagy in PC/PTX cells. (A) The expression of FEZF1-AS1 was detected in si-NC and si-FEZF1-AS1 groups in DU145/PTX and LNCaP/PTX cells with qRT-PCR. (B and C) IC₅₀ values of DU145/PTX and LNCaP/PTX cells transfected with si-NC or si-FEZF1-AS1 treated with paclitaxel of different concentrations (0–16 ng/mL) were detected. (D and E) Cell proliferations were assessed in si-NC and si-FEZF1-AS1 groups of DU145/PTX and LNCaP/PTX cells with CCK-8 assay. (F to H) The protein expressions of LC3II/LC3I, ATG5 and Beclin-1 were measured in si-NC and si-FEZF1-AS1 groups of DU145/PTX and LNCaP/PTX cells with Western blot. (I to K) The protein expressions of E-cadherin, N-cadherin and Vimentin were measured in si-NC and si-FEZF1-AS1 groups of DU145/PTX and LNCaP/PTX cells with Western blot. **p* < 0.05.

was significantly higher in DU145/PTX and LNCaP/PTX cells than in DU145 and LNCaP cells (Figure 3F). The expression of miR-25-3p in si-FEZF1-AS1 group was significantly higher than that of si-NC group in DU145/PTX and LNCaP/PTX cells (Figure 3G and 3H). Not only that, the mass array showed FEZF1-AS1 was enriched for miR-25-3p in DU145/PTX and LNCaP/PTX cells (Figure 3I and 3J). These results reflected that miR-25-3p was a target miRNA of FEZF1-AS1 and played a role in the paclitaxel resistance mechanism of PC.

FEZF1-AS Regulated Cell Viability, Proliferation and Autophagy by Inhibition of miR-25-3p in PC/PTX Cells

To further clarify the specific regulatory mechanisms of FEZF1-AS1, overexpressing or

downregulating miR-25-3p of DU145/PTX and LNCaP/PTX cells was obtained by transfection of the vector of miR-25-3p or anti-miR-25-3p (Figure 4A and 4B). As shown in Figure 4C to 4F, overexpression of miR-25-3p remarkably reduced IC₅₀ and cell proliferation whereas downexpression of miR-25-3p could significantly reverse the effect of inhibition of FEZF1-AS1 on DU145/PTX and LNCaP/PTX cells. A similar effect was observed in the protein expression of autophagy-related factors. miR-25-3p transfection contributed to the protein expression of LC3II/LC3I, ATG5 and Beclin-1, promoting autophagy in DU145/PTX and LNCaP/PTX cells. In addition, the effects of inhibition of FEZF1-AS1 were significantly inhibited by anti-miR-25-3p transfection in DU145/PTX and LNCaP/PTX cells (Figure 4C to 4I). Besides,

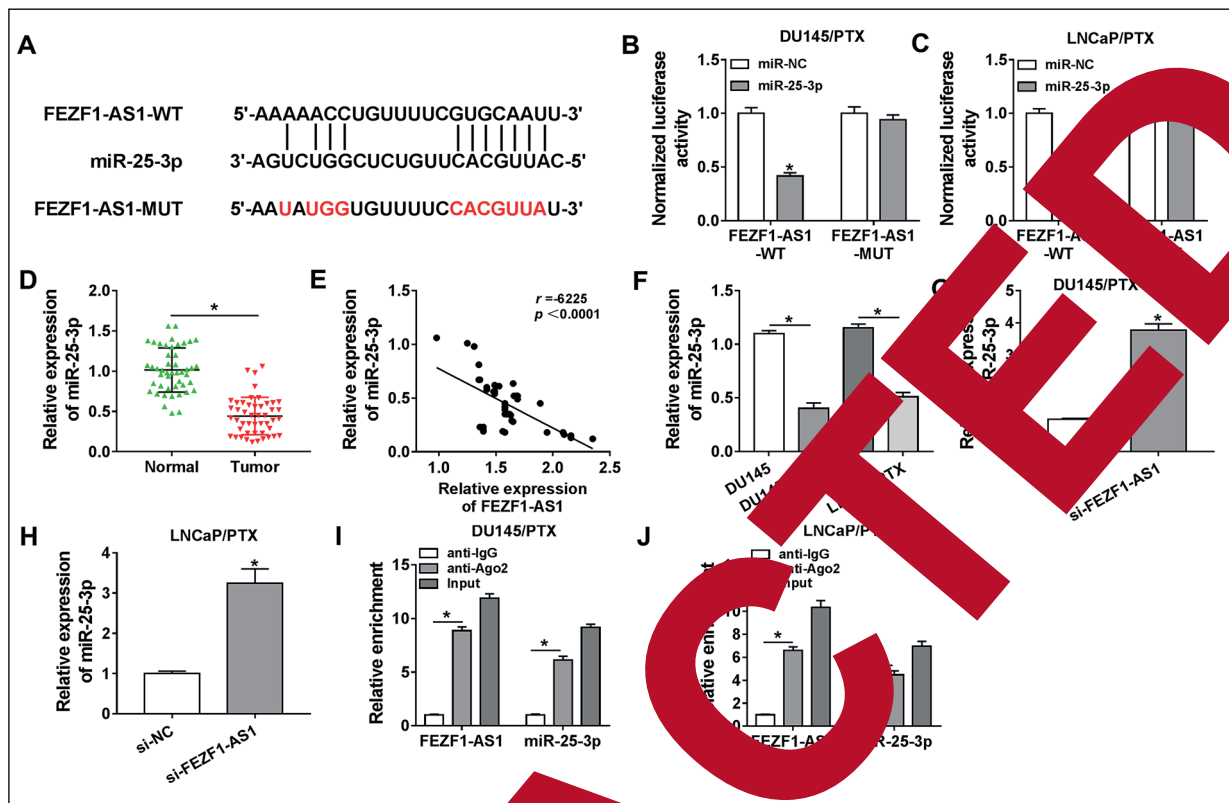


Figure 3. miR-25-3p was a target miRNA of FEZF1-AS1. (A) Binding sites for miR-25-3p on the FEZF1-AS1. (B and C) The luciferase activities were measured in DU145/PTX and LNCaP/PTX cells co-transfected with FEZF1-AS1-WT or FEZF1-AS1-MUT and miR-NC or miR-25-3p. (D) The expression of miR-25-3p in normal tissues and PC tumor tissues was detected with qRT-PCR. (E) The correlation between FEZF1-AS1 and miR-25-3p was measured in tumor tissues through Spearman correlation analysis. (F) The expression of miR-25-3p in DU145/PTX, DU145, LNCaP/PTX and LNCaP cells was detected using qRT-PCR. (G and H) The expression of miR-25-3p was detected in si-NC and si-FEZF1-AS1 groups of DU145/PTX and LNCaP/PTX cells with qRT-PCR. (I and J) The enrichment of FEZF1-AS1 was detected in DU145/PTX and LNCaP/PTX cells using RIP assay. * $p < 0.05$.

miR-25-3p overexpression increased E-cadherin protein expression, inhibited N-cadherin and Vimentin protein expression in DU145/PTX and LNCaP/PTX cells. The inhibition of miR-25-3p significantly reversed the effects of DU145/PTX and LNCaP/PTX cells transfected with si-FEZF1-AS1 (Figure 4J to 4L). Taken together, FEZF1-AS1 promoted cell proliferation, and autophagy, and MT in DU145/PTX and LNCaP/PTX cells by targeting miR-25-3p, suggesting that the miR-25-3p/FEZF1-AS1 axis may be a novel regulatory mechanism of FEZF1-AS1/miR-25-3p play an important regulatory role to predict the prognosis of PC.

FEZF1-AS Regulated ITGB8 Expression through Targeting miR-25-3p

We predicted the potential target gene of miR-25-3p using starBase 3.0. ITGB8 was a

downstream target gene for miR-25-3p and highly expressed at protein expression in tumor tissues and DU145/PTX and LNCaP/PTX cells (Figure 5A, 5D and 5E). Luciferase reporter assay showed that, when miR-25-3p was combined with ITGB8-3UTR-WT, instead of ITGB8-3UTR-MUT, luciferase activities sharply reduced (Figure 5B and 5C). More than that, overexpression miR-25-3p significantly inhibited the protein expression of ITGB8 in DU145/PTX and LNCaP/PTX cells (Figure 5F). In addition, knocking out FEZF1-AS1 significantly suppressed the protein expression of ITGB8, which was reversed by downregulation of miR-25-3p (Figure 5G and 5H). These data indicate that ITGB8 is the target gene for miR-25-3p and FEZF1-AS1 regulated the protein expression of ITGB8 by targeting miR-25-3p in DU145/PTX and LNCaP/PTX cells.

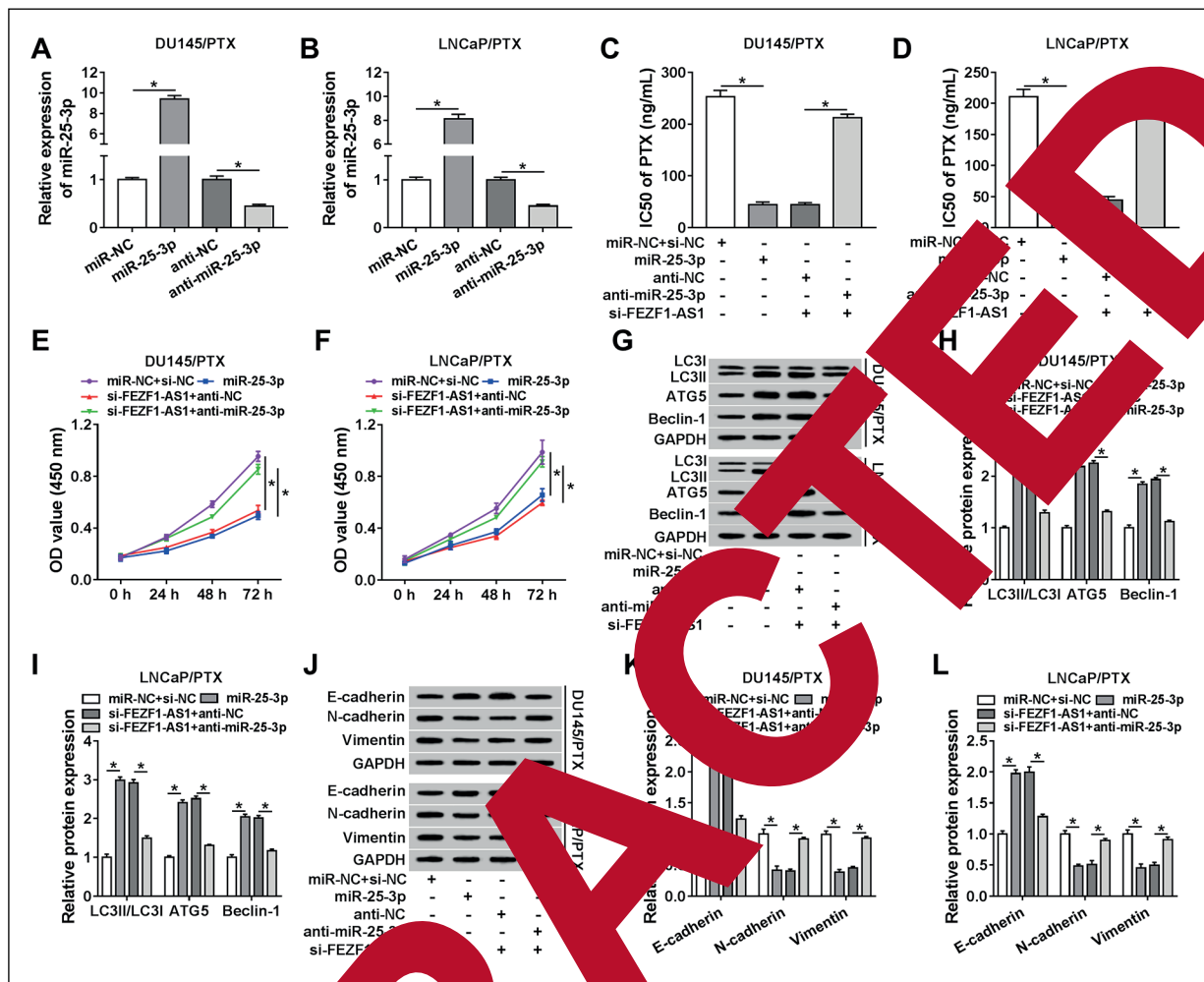


Figure 4. FEZF1-AS regulates cell viability and autophagy by inhibition of miR-25-3p in PC/PTX cells. (A and B) The expression of miR-25-3p was detected in miR-NC, miR-25-3p, anti-NC and anti-miR-25-3p groups of DU145/PTX and LNCaP/PTX cells with qRT-PCR. (C and D) Cell viability of DU145/PTX and LNCaP/PTX cells was measured in miR-NC + si-NC, miR-25-3p, si-FEZF1-AS1 + anti-NC, si-FEZF1-AS1 + anti-miR-25-3p groups. (E and F) Cell proliferations of DU145/PTX and LNCaP/PTX cells were measured in miR-NC + si-NC, miR-25-3p, si-FEZF1-AS1 + anti-NC, si-FEZF1-AS1 + anti-miR-25-3p groups with CCK-8 assay. (G to I) The protein expression of LC3II/LC3I, ATG5 and Beclin-1 was measured in miR-NC + si-NC, miR-25-3p, si-FEZF1-AS1 + anti-NC, si-FEZF1-AS1 + anti-miR-25-3p groups of DU145/PTX and LNCaP/PTX cells with Western blot. (J to L) Relative protein expression of E-cadherin, N-cadherin and Vimentin was measured in miR-NC + si-NC, miR-25-3p, si-FEZF1-AS1 + anti-NC, si-FEZF1-AS1 + anti-miR-25-3p groups of DU145/PTX and LNCaP/PTX cells with Western blot. * $P < 0.05$.

The Effect of Low FEZF1-AS Expression on Cell Viability, PT and Autophagy Reverses the Promotion of ITGB8 in DU145/PTX Cells

To further clarify the role of ITGB8 in the regulation mechanism of FEZF1-AS, we obtained high- and low-expression ITGB8 of DU145/PTX and LNCaP/PTX cells to explore the function of ITGB8 (Figure 6A and 6B). The CCK-8 assay showed that the inhibition of ITGB8 significantly reduced IC50 and proliferative capacity, promot-

ed protein expression of LC3II/LC3I, ATG5 and Beclin-1 in DU145/PTX and LNCaP/PTX cells, whereas increased expression of ITGB8 attenuated these effects of DU145/PTX and LNCaP/PTX cells transfected with si-FEZF1-AS1 (Figure 6C to 6I). Moreover, si-ITGB8 transfection enhanced E-cadherin protein expression and reduced N-cadherin and Vimentin protein expression. The effects of si-FEZF1-AS1 on E-cadherin, N-cadherin and Vimentin protein expression were reversed through promotion of ITGB8 in

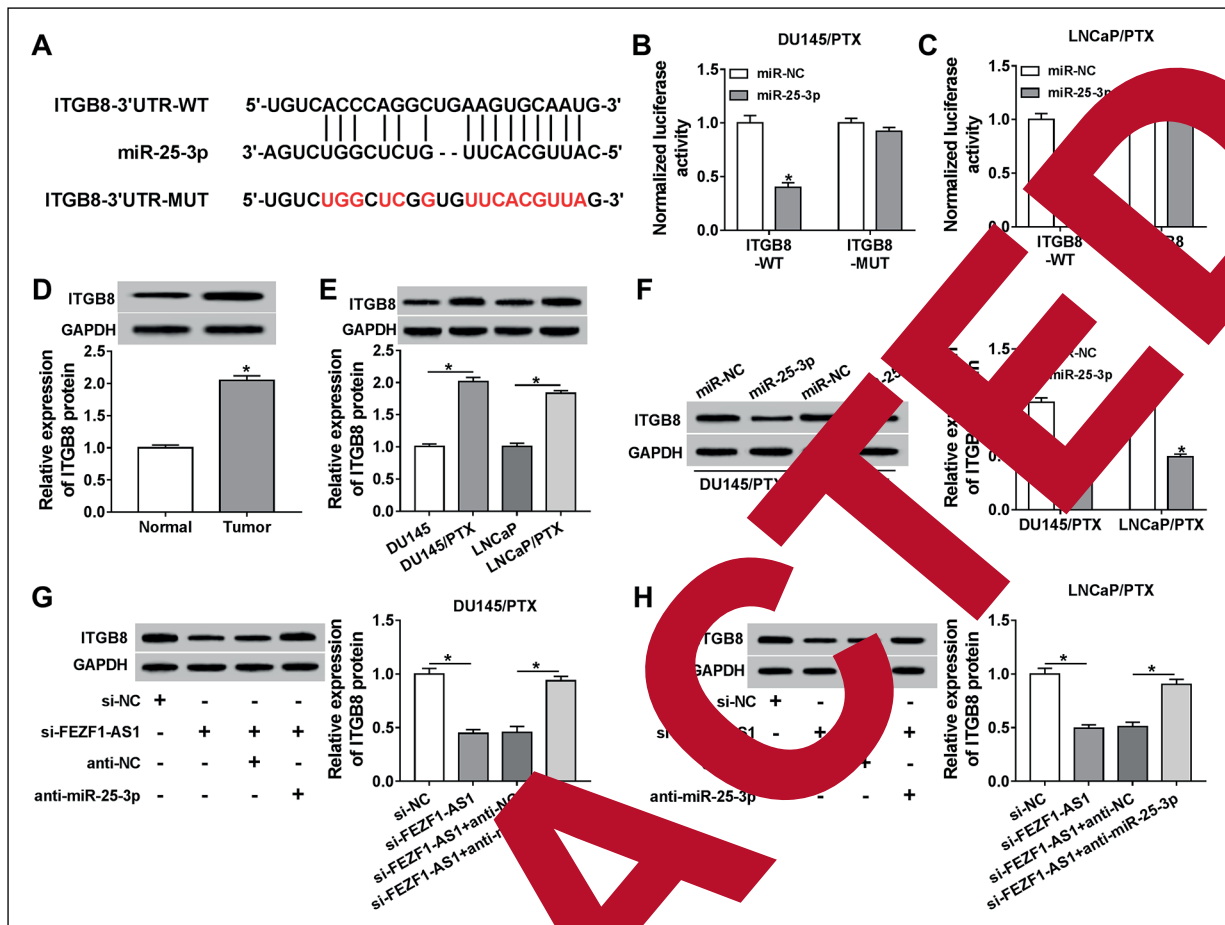


Figure 5. FEZF1-AS regulated ITGB8 expression through miR-25-3p. (A) Predicted binding sites for miR-25-3p on the ITGB8 transcript. (B and C) The luciferase activities were measured in DU145/PTX and LNCaP/PTX cells co-transfected with ITGB8-WT or ITGB8-MUT and miR-NC or miR-25-3p. (D) The protein expression of ITGB8 in normal tissues and PC tumor tissues was detected with Western blot. (E) The protein expression of ITGB8 in DU145/PTX, DU145, LNCaP/PTX and LNCaP cells was detected using Western blot. (F) The protein expression of ITGB8 was detected in si-NC, si-FEZF1-AS1, si-FEZF1-AS1 + anti-NC, si-FEZF1-AS1 + anti-miR-25-3p groups of DU145/PTX and LNCaP/PTX cells with Western blot. * $p < 0.05$.

DU145/PTX and LNCaP/PTX cells (Figure 6J to 6L). These results indicated that the effects of low FEZF1 expression on cell viability, EMT and autophagy were reversed by promotion of ITGB8 in DU145/PTX cells.

Knockdown of FEZF1-AS1 Suppressed Cell Growth and Decreased PTX Sensitivities of PC Cells in Vivo

We have verified the regulatory mechanism of FEZF1-AS1 in mice. The results showed that reducing the expression of FEZF1-AS1 in DU145/PTX and LNCaP/PTX cells significantly inhibited tumor volume and reduced tumor weight (Figure 7A and 7B). In addition, the expression of FEZF1-AS1 and the mRNA and protein ex-

pression of ITGB8 in sh-FEZF1-AS1 group were higher than that in sh-NC group (Figure 6C and 6D). The expression of miR-25-3p in sh-FEZF1-AS1 group was higher than that in sh-NC group (Figure 6C). The results showed that low expression of FEZF1-AS1 could reduce PTX sensitivities and inhibit tumor growth via regulating miR-25-3p/ITGB8 axis.

Discussion

With the continuous development of molecular biology and bioinformatics technology, more and more molecular mechanisms of cancer have been discovered. Accumulating

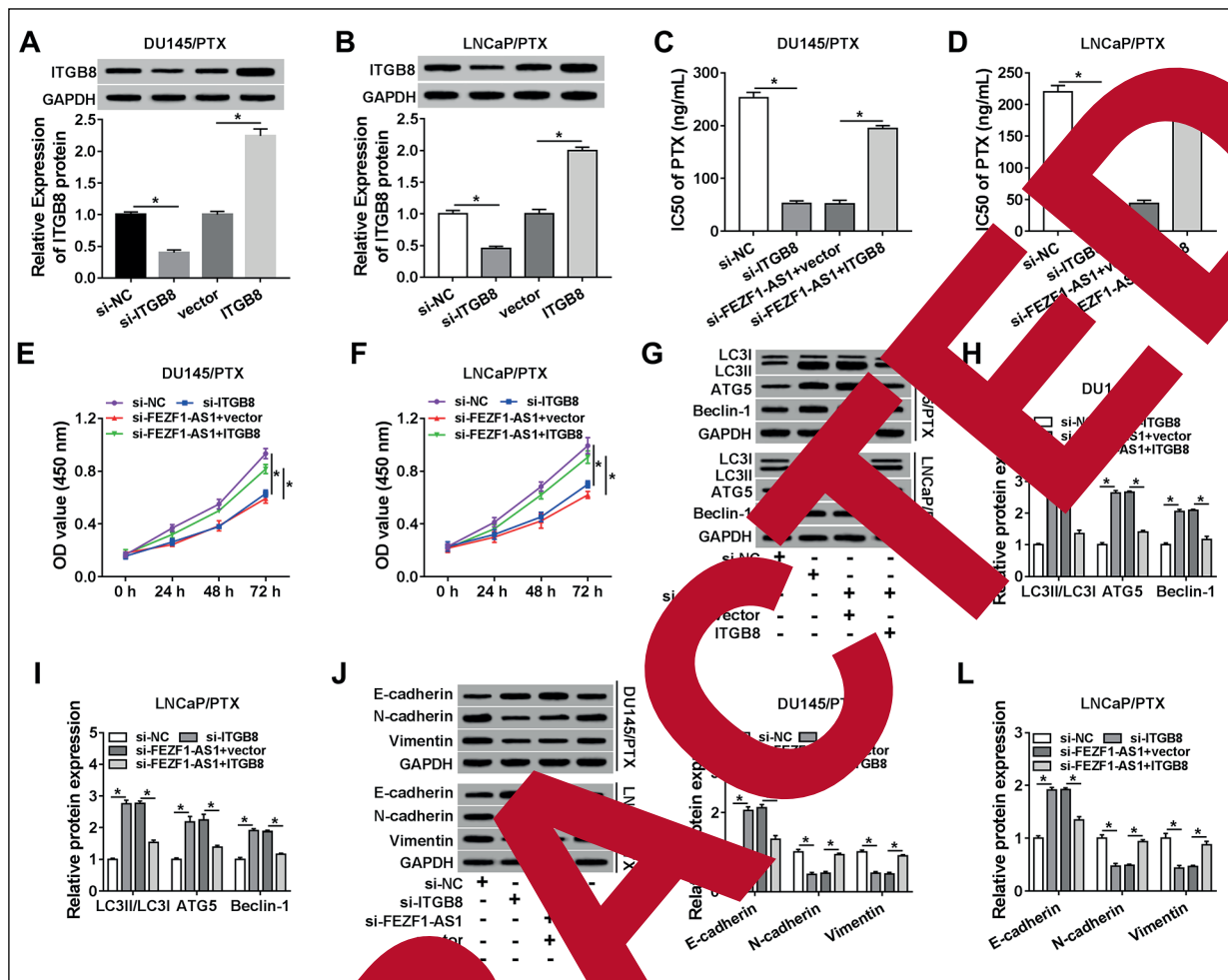


Figure 6. The effects of low FEZF1-AS1 expression on cell viability, EMT and autophagy were reversed by promotion of ITGB8 in PC/PTX cells. (A and B) The protein expression of ITGB8 was detected in si-NC, si-ITGB8, vector and ITGB8 groups of DU145/PTX and LNCaP/PTX cells with Western blot. (C and D) IC50 values of DU145/PTX and LNCaP/PTX cells were measured in si-NC, si-ITGB8, si-FEZF1-AS1 + vector, si-FEZF1-AS1 + si-ITGB8 groups. (E and F) Cell proliferations of DU145/PTX and LNCaP/PTX cells were measured in si-NC, si-ITGB8, si-FEZF1-AS1 + vector, si-FEZF1-AS1 + si-ITGB8 groups with CCK-8 assay. (G to I) The protein expression of LC3II/LC3I, ATG5 and Beclin-1 was measured in si-NC, si-ITGB8, si-FEZF1-AS1 + vector, si-FEZF1-AS1 + si-ITGB8 groups of DU145/PTX and LNCaP/PTX cells with Western blot. (J to L) The protein expression of E-cadherin, N-cadherin and Vimentin was measured in si-NC, si-ITGB8, si-FEZF1-AS1 + vector, si-FEZF1-AS1 + si-ITGB8 groups of DU145/PTX and LNCaP/PTX cells with Western blot. * $p < 0.05$.

evidence suggested that many lncRNAs are abnormally expressed in cancer cells, implying a close relationship with the characteristics of PC cells, including proliferation and metastasis³²⁻³⁴. Drug resistance is an important issue in the current treatment of PC^{35,36}. More than that, lncRNA is also involved in the chemoresistance mechanisms of various cancers, including colorectal cancer, gastric cancer, cervical cancer, breast cancer and prostate cancer³⁷⁻⁴⁰. Interestingly, increased MALAT1 expression could contribute to chemoresistance in gastric cancer,

lung cancer, and prostate cancer⁴¹⁻⁴³. Autophagy and EMT are important mechanisms of tumor cell development. Autophagy can improve the adaptability of cells, and abnormal autophagy mechanism will lead to the appearance of cancer cells⁴⁴. EMT is an important biological process for tumor cells to acquire the ability to migrate and invade. Moreover, lncRNA plays an important role in the regulation of autophagy and EMT⁴⁵. It has been observed that lncRNA CPS1-IT1 inhibited EMT and metastasis via suppressing hypoxia-induced autophagy by

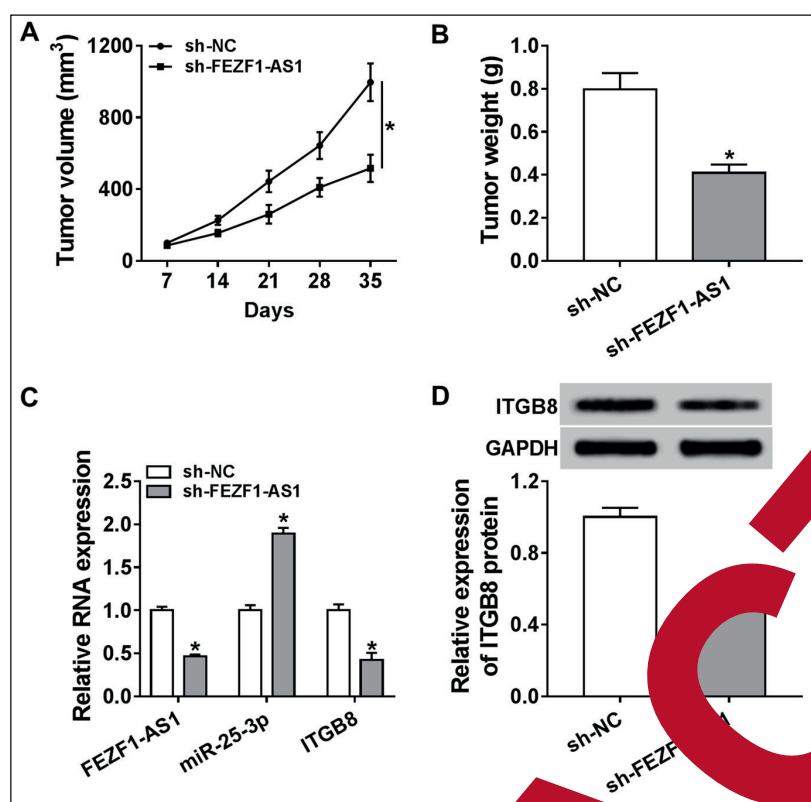


Figure 7. Knockdown of FEZF1-AS1 suppressed tumor growth and decreased PTX sensitivities of PC cells *in vivo*. (A) The tumor growth curves of sh-NC and sh-FEZF1-AS1 groups *in vivo*. (B) The mean tumor weight was measured in sh-NC and sh-FEZF1-AS1 groups *in vivo*. (C) The expression of FEZF1-AS1, miR-25-3p and ITGB8 was detected in sh-NC and sh-FEZF1-AS1 groups with qRT-PCR. (D) The protein expression of ITGB8 in sh-NC and sh-FEZF1-AS1 groups was measured using Western blot. * $p < 0.05$.

inactivation of HIF-1 α in colorectal cancer. In the present research we found that FEZF1-AS1 expression was increased in tumor tissues and PC/PTX cells, implying that FEZF1-AS1 was closely related to paclitaxel resistance in PC. Importantly, reduced FEZF1-AS1 expression inhibited proliferation, EMT as well as promoted autophagy in PC/PTX cells, revealing that the expression of FEZF1-AS1 contributed to paclitaxel resistance of PC/PTX cells. Decreasing the chemoresistance of PC cells. lncRNAs have been shown to play an important role in cancer drug resistance. Transcriptome analysis revealed that many miRNAs were differentially expressed in cancer and involved in the metastasis, proliferation and drug sensitivity of cancer cells to drugs, including paclitaxel⁴⁷⁻⁴⁹. Therefore, miR-25-3p significantly reduced the resistance of paclitaxel-refractory PC cells to paclitaxel in PC cells. However, we demonstrated that FEZF1-AS1 directly targeted miR-25-3p, and that reducing miR-25-3p significantly suppressed the resistance of PC/PTX cells transfected with si-FEZF1-AS1 and reduced the sensitivity of PC/PTX cells to paclitaxel. Thus, FEZF1-AS1

decreased the paclitaxel resistance of PC cells through downregulating miR-25-3p. Next, we found that the ITGB8 is the target gene of miR-25-3p. We thought that ITGB8 was involved in the regulatory mechanism of FEZF1-AS1/miR-25-3p. Many studies⁵⁰⁻⁵² demonstrated that ITGB8 also participated in chemoresistance of various cancers. Wang et al⁵³ showed that silencing ITGB8 reversed gefitinib resistance for hepatic cancer. In addition, ITGB8 also regulated cell progression in lung cancer and colorectal cancer^{51, 54}. In this paper, we found that the inhibition of ITGB8 expression significantly reduced drug-resistant cell proliferation, EMT and promoted autophagy and increased paclitaxel sensitivity in PC cells. Moreover, increasing the expression of ITGB8 reversed the inhibitory effect of si-FEZF1-AS1 on PC/PTX cells and reduced the resistance of PC cells.

lncRNAs have been reported to act as competing endogenous RNAs (ceRNAs) to modulate mRNA *via* sponging miRNA in the regulation mechanism of cell growth in cancers. Besides, in hepatocellular carcinoma, lncRNA CCAT1 promoted autophagy through regulating ATG7 expression by competitively binding miR-181⁵⁵.

In this study, we first found and verified that lncRNA FEZF1-AS1 promoted chemoresistance, autophagy and EMT through regulation of miR-25-3p/ITGB8 axis in PC, providing a new regulatory mechanism of paclitaxel-resistance of PC and a novel therapeutic target.

The Authors declare that they have no conflict of interests.

The present study was approved by the Ethical Review Committee of West China Hospital, Sichuan University.

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