

Targeted regulation of miR-195 on MAP2K1 for suppressing ADM drug resistance in prostate cancer cells

J.-Y. ZHANG, Y.-N. LI, X. MU, Z.-L. PAN, W.-B. LIU

Department of Urology, The Second Affiliated Hospital, Fujian Medical University, Fuzhou, Fujian, China

Abstract. – **OBJECTIVE:** Extra-cellular signal-regulated kinase/mitogen-activated protein kinase (ERK/MAPK) signaling pathway participates in cell proliferation, cycle and apoptosis. MAPK kinase 1 (MAP2K1) activates the ERK/MAPK pathway. The down-regulation of miR-195 is correlated with the onset and drug resistance of prostate cancer. Bioinformatics analysis identified complementary binding sites between miR-195 and MAP2K1. This study aimed to investigate the effect of miR-195 on the proliferation, apoptosis and adriamycin (ADM) resistance of prostate cancer cells.

MATERIALS AND METHODS: Dual-luciferase reporter gene assay confirmed targeted regulation between miR-195 and MAP2K1. Drug-resistant cell line DU145/ADM and PC-3/ADM were generated for comparing miR-195 and MAP2K1 expression. Apoptosis was measured by flow cytometry and caspase-3 activity was quantified. Cultured cells were treated with miR-195 mimic, followed by quantitative real-time PCR (qRT-PCR) was used to detect MAP2K1 expression. Western blot measured MAP2K1, ERK1/2 and phosphorylated ERK1/2 (p-ERK1/2) expression, and flow cytometry quantified cell apoptosis, followed by EdU staining for cell proliferation.

RESULTS: Targeted regulation existed between miR-195 and MAP2K1 in PC3A. Drug-resistant cells had lower miR-195 than parental cells, while MAP2K1 expression was higher. Under ADM treatment with IC50 concentration, drug resistance cells showed low apoptosis. The transfection of miR-195 decreased MAP2K1 expression and p-ERK1/2, elevated cell apoptosis and decreased EdU positive rate or cell proliferation.

CONCLUSIONS: The down-regulation of miR-195 is correlated with ADM resistance of prostate cancer cells. The over-expression of miR-195 weakens cancer cell proliferation, facilitates cell apoptosis and decreases ADM resistance via targeted inhibition on MAP2K1 expression and ERK/MAPK signal pathway.

Key Words:

Prostate cancer, miR-195, MAP2K1, ERK/MAPK, Drug resistance.

Introduction

Prostate carcinoma is a common malignant tumor in the urinal-reproductive system. Due to its relatively high incidence, it is the sixth most popular malignant tumor in males and has a higher frequency in male urinal reproductive tumors^{1,2}. Prostate cancer has relatively higher difficulty for treatment, unfavorable prognosis and higher mortality. Chemotherapy has become a critical treatment measure for prostate cancer, whilst the drug resistance is one major factor limiting clinical treatment efficiency^{3,4}. Mitogen-activated protein kinase kinase 1 (MAP2K1) is the upstream protein kinase of extra-cellular signal-regulated kinase (ERK) and can activate extra-cellular signal regulated kinase/mitogen activated protein kinase (ERK/MAPK) signal pathway. Abnormally elevated expression and function of MAP2K1 are correlated with onset, progression, metastasis and drug resistance of various tumors⁵⁻⁷. Previous studies showed the involvement of enhanced MAP2K1 expression or functional activity in malignant features including drug resistance and invasion of prostate cancer^{8,9}. MicroRNA (miR) is an endogenous non-coding small RNA molecule in eukaryotes and can regulate target gene expression by degrading mRNA or inhibiting mRNA translation by complementary binding with 3'-untranslated region (3'-UTR) of target gene mRNA, thus participating in the regulation of biological processes including cell survival, proliferation, apoptosis and migration.

Abnormal expression or function of microRNA has attracted increasing interests in tumor drug resistance^{10,11}. Previous studies¹²⁻¹⁴ showed the correlation between miR-195 down-regulation and the onset, progression, prognosis and drug resistance of prostate cancer. Bioinformatics analysis showed the existence of complementary binding sites between miR-195 and 3'-UTR of MAP2K1 mRNA, indicating possibly targeted regulatory roles. This work thus investigated the role of miR-195 in mediating MAP2K1 expression and affecting proliferation, apoptosis and ADM resistance of prostate cancer cells.

Materials and Methods

Major Reagents and Materials

Normal prostate gland epithelial cell RWPE-1 (CRL-11609), prostate cancer cell line DU145 (HTB-81) and PC-3 (CRL-1435) were purchased from the American Type Culture Collection (ATCC; Manassas, VA, USA). Dulbecco's Modified Eagle Medium (DMEM), keratinocyte-serum free medium (keratinocyte-SFM), minimum essential media (Opti-MEM) and fetal bovine serum (FBS) were purchased from Gibco (Grand Island, NY, USA). RNAiSelect was purchased from TaKaRa (Otsu, Shiga, Japan). Fluorescent quantitative PCR kit TranscripT Green One-Step quantitative Real Time PCR (qRT-PCR) SuperMix was purchased from Transgen (Beijing, China). miR-195 mimic and microRNA-normal control miR-NC were purchased from Ribobio (Guangzhou, China). Doxifluridine 2000 was purchased from Invitrogen (Carlsbad, CA, USA). Rabbit anti-human polyclonal antibody anti-MAP2K1, ERK1/2 and p-ERK1/2 were purchased from Abcam (Cambridge, MA, USA). Rabbit anti-human β -actin polyclonal antibody was purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Horseradish peroxidase (HRP) conjugated secondary antibody was purchased from Sangon Biotech (Shanghai, China). Annexin V/propidium iodide cell apoptosis test kit and BeyoECL chemiluminescence were purchased from Beyotime (Shanghai, China). Adriamycin (ADM) and cell counting kit were purchased from MedchemExpress (Monmouth Junction, NJ, USA). EdU cell proliferation flow cytometry kit was purchased from Molecular Probes (Eugene, OR, USA). Luciferase activity assay kit Dual-Glo Luciferase Assay System,

and Dual-Luciferase reporter plasmid pGL3 Luciferase vector were purchased from Promega Inc. (Austin, TX, USA).

Cell Culture

DU145 and PC-3 cells were incubated in DMEM medium containing 10% fetal bovine serum (FBS) and were kept in a 5% CO₂ incubator with 5% CO₂ model HERAcells (Thermo Fisher Scientific, Waltham, MA, USA). RWPE-1 cells were cultured in keratinocyte-SFM medium containing 5% epithelial growth factor (EGF) and 0.0 mg/ml bovine Pituitary Extract (BPE) in a 37°C incubator with 5% CO₂. Cells were seeded at 1:4 ratio, and the cells at log-growth phase were used for experiments.

Generation of Adriamycin (ADM) Resistant Cell Model

To generate an ADM drug-resistant cell model, DU145 and PC-3 cells at log-growth phase were incubated with 1 μ g/ml ADM into the culture medium. After 24 h, the fresh culture medium was replaced. After 2 weeks of stable growth, the ADM concentration was gradually elevated to 2 μ g/ml, 4 μ g/ml, 8 μ g/ml and 16 μ g/ml until stable growth for repeated passage within 16 μ g/ml ADM. ADM resistant prostate cancer cell lines DU145/ADM and PC-3/ADM were then generated. DU145, PC-3, DU145/ADM and PC-3/ADM cells were seeded into 96-well plate at 10000 cells per well density. After 24 h attached growth, cells were treated with 0, 5, 10, 20, 40, 80, 160 and 320 μ g/ml ADM, with 6 parallel replicates for each concentration. After 48 h incubation, each well was added with 10 μ l cell counting kit-8 (CCK-8) solution, and absorbance (A) values at 450 nm wavelength (A450) were measured after 4 h reaction. Inhibition rate = (1-A450 of treatment group)/A450 of treatment group \times 100%. Half maximal inhibitory concentration (IC₅₀) for the concentration of inhibiting 50% cell growth was calculated by SPSS 18.0 software (SPSS, Inc., Chicago, IL, USA). Resistance index (RI) = IC₅₀ of drug-resistant cells/IC₅₀ of parental cells.

Flow Cytometry for Cell Proliferation

Cells were re-suspended in DMEM medium containing 10% FBS. EdU Alexa Fluor 488 Flow Cytometry Assay Kit (Cat. No. 35002, Invitrogen, Carlsbad, CA, USA) was used to measure proliferation potency of cells. In brief, cells were incubated using 10 μ M EdU for 2 h and were continuously incubated for 48 h. Cells were digested

by trypsin and collected. After centrifugation, washing, fixation and permeabilization, cells were incubated in assay buffer containing Alex Fluor 488 labels for 30 min room temperature incubation. After washing and centrifugation, FC500 MCL flow cytometry (Beckman Coulter Inc., Brea, CA, USA) was used to measure cell proliferation.

Dual Luciferase Activity Assay

PCR products for full-length fragment or mutant form of MAP2K1 gene 3'-UTR were digested by dual restriction enzymes and were ligated to pLUC plasmid. After transforming competent bacteria, the cells with correct sequences were selected and named as pLUC-MAP2K1-WT or pLUC-MAP2K1-MUT. Lipofectamine 2000 was used to co-transfect pLUC-MAP2K1-WT (or pLUC-MAP2K1-MUT) and miR-195 mimic (or miR-NC) into HEK293T cells for 48h continuous incubation. Dual-Glo Luciferase Assay System kit was used for measuring Dual-Luciferase activity.

Cell Transfection and Grouping

Cultured DU145/ADM and PC-3/ADM cells were divided into two groups: miR-NC transfection group, and miR-195 mimic transfection group. In brief, 100 µl serum-free medium was used to dilute 10 µl Lipofectamine 2000, 100 nmol miR-NC, 30 nmol miR-195. After 5 min room temperature incubation, Lipofectamine 2000 was gently mixed with miR-NC and miR-195 for 20 min room temperature incubation. The transfection mixture was slowly added into the culture medium for general mixture and 82 h of continuous incubation for collecting cells. Cells from all groups were seeded into 6-well plate, and 50 µl/ml ADM was added when reaching 50% confluence. After 48 h of continuous incubation, cell apoptosis was measured by flow cytometry. Trypsin (Beyotime, Shanghai, China) was used to collect treated cells from all groups. After 24 h ADM incubation for 2 h, cells were

continuously incubated for 48 h as described in previous sections. Cell proliferation was measured by test kit.

qRT-PCR for Measuring Gene Expression

RNAiso Plus was used to extract cellular RNA. The relative expression of each gene was measured by qRT-PCR using TaqMan Green One-Step qRT-PCR SuperMix. In a 20 µl reaction system, one added 1 µl template RNA, 0.3 µM forward primer, 0.3 µM reverse primer, 10 µl 2× TransScript TipOne qPCR SuperMix, 0.4 µl One-Step RT Enzyme II, 0.1 µl Passive Reference Dye 1, and RNase-free water up to 20 µl. The qRT-PCR reaction conditions were: 45°C 5 min and 94°C 30 s, followed by 40 cycles each consisting of 95°C 30 s and 60°C 30 s. The gene expression was measured on Bio-Rad CFX96 Real-time fluorescent quantitative PCR cycler (Bio-Rad, Hercules, CA, USA). The primers for qRT-PCR were listed in Table I.

Western Blot

Protein was extracted by radioimmunoprecipitation assay (RIPA) lysis buffer (Tiangen Biotech Co. Ltd., Beijing, China), followed by protein quantification and quality control. Total of 40 µg protein samples were separated in sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) (10% separating gel, 4% condensing gel, 45 V, 150 min) and were transferred to polyvinylidene difluoride (PVDF, Amersham Biosciences, Little Chalfont, Buckinghamshire, UK) membrane (300 mA, 100 min). The membrane was blocked in 5% defatted milk powder at room temperature and was incubated in primary antibody (MAP2K1 at 1:2000, ERK1/2 at 1:2000, p-ERK1/2 at 1:1000, and β-actin at 1:10000) for 4°C overnight incubation. On the next day, the membrane was washed in Phosphate-Buffered Saline and Tween-20 (PBST) three times, and horseradish peroxidase (HRP) conjugated secondary antibody (1:15000 dilution) was added for 60 min room temperature incubation, followed

Table I. Primers for the qRT-PCR assay.

Gene		Sequences
miR-195	Forwards	5'-GAATCCGCCTCAAGAGAACAAGGTGGAG-3'
	Reverse	5'-AGATCTCCCATGGGGGCTCAGCCCCT-3'
MAP2K1	Forwards	5'-ATCTTCGGGAGAAGCACAAAG-3'
	Reverse	5'-CGAAGGAGTTGGCCATAGAG-3'
	Forwards	5'-TACCACATCCAAGAAGGCAG-3'
	Reverse	5'-TGCCCTCCAATGGATCCTC-3'

by washing in PBST three times. BeyoECL Plus working solution (prepared from an equal volume of solution A and B) was added onto the membrane. After 2-3 min of dark incubation, the membrane was exposed, and images were scanned for data processing.

Cell Apoptosis Assay

Cells were digested by Trypsin and collected for Phosphate-Buffered Saline (PBS, Tiangen Biotech Co. Ltd., Beijing, China) washing by centrifugation. Cells were resuspended in 100 μ l Annexin V Binding Buffer, and 5 μ l Annexin V-FITC and 5 μ l PI were sequentially added for staining. After 15 min of room temperature incubation, 400 μ l Annexin V Binding Buffer was added for measuring cell apoptosis on FC500 MCL flow cytometry (Beckman Coulter, Brea, CA, USA).

Statistical Analysis

SPSS 18.0 software (SPSS, Inc., Chicago, IL, USA) was used for data processing. Measurement data were presented as mean \pm standard deviation (SD). The Student's *t*-test was used to compare the differences between the two groups. The differences among groups were analyzed using Tukey's post-hoc test to validate the one-way analysis of variance (ANOVA). A statistical significance was defined when $p < 0.05$.

Result

Targeted Regulation between miR-195 and MAP2K1 mRNA

Bioinformatics analysis showed the existence of complementary binding sites between miR-195 and 3'-UTR of MAP2K1 mRNA (Figure 1A). Dual-Luciferase gene reporter assay showed that the transfection of miR-195 significantly decreased relative Luciferase activity in HEK293T cells transfected with pLUC-MAP2K1-WT plasmid. It had no significant effect on the relative Luciferase activity of HEK293T cells transfected with pLUC-MAP2K1-MUT plasmid (Figure 1B). These results suggest that miR-195 targets 3'-UTR of MAP2K1 mRNA to inhibit its expression.

Dose-Response of MiR-195 and Regulation of MAP2K1 in Drug-Resistant Prostate Cancer Cell Lines

In parental DU145 cells, IC_{50} reached 7.44 ± 0.82 μ g/ml, whilst drug-resistant DU145/ADM cells

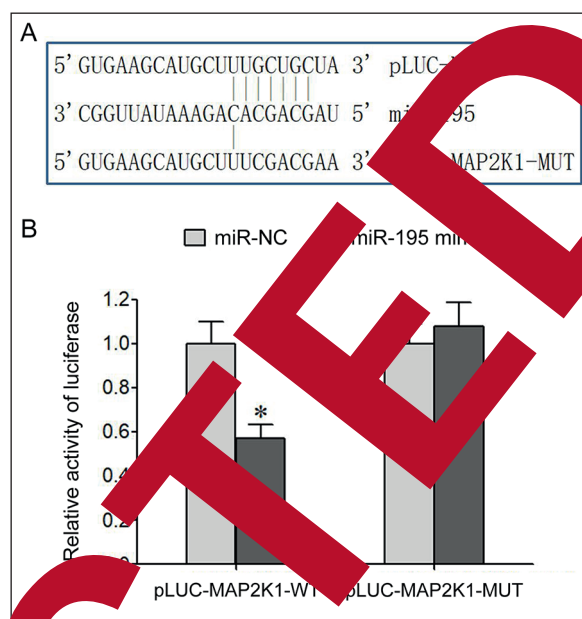


Figure 1. Targeted regulation between miR-195 and MAP2K1 mRNA. **A**, Functional site between miR-195 and 3'-UTR of MAP2K1 mRNA. **B**, Dual-Luciferase gene reporter assay. * $p < 0.05$ compared to the miR-NC group.

had IC_{50} values at 86.37 ± 7.11 μ g/ml. The relative drug-resistant index of DU145/ADM cells against parental DU145 cells was 11.61 (Table II). Parental PC-3 cells had IC_{50} values at 5.29 ± 0.47 μ g/ml, and drug-resistant cell line PC3/ADM had IC_{50} values at 57.71 ± 5.35 μ g/ml, with relative drug-resistant index at 10.91 against parental cells (Table I). The qRT-PCR results showed that, compared to normal prostate epithelial cell line RWPE-1, DU145 cells showed significantly decreased miR-195 expression, and even lower miR-195 expression occurred in drug-resistant cell line DU145/ADM cells. PC-3 cells had remarkably lower miR-195 expression compared to RWPE-1 cells, and drug-resistant cell PC-3/ADM had an even lower miR-195 expression (Figure 2A). Compared to RWPE-1 cells, DU145 cells had significantly higher MAP2K1 mRNA expression, which was further elevated in drug-resistant DU145/ADM cells. PC-3 cells also showed higher MAP2K1 mRNA expression compared to RWPE-1 cells, and MAP2K1 mRNA level was even higher in drug-resistant PC-3/ADM cells (Figure 2B). Western blot results showed that, compared to RWPE-1 cells, DU145 cells had remarkably higher MAP2K1 proteins, which was further elevated in drug-resistant DU145/ADM cells. Moreover, compared to RWPE-1 cells, PC-3 cells showed remarkably elevated MAP2K1 protein expres-

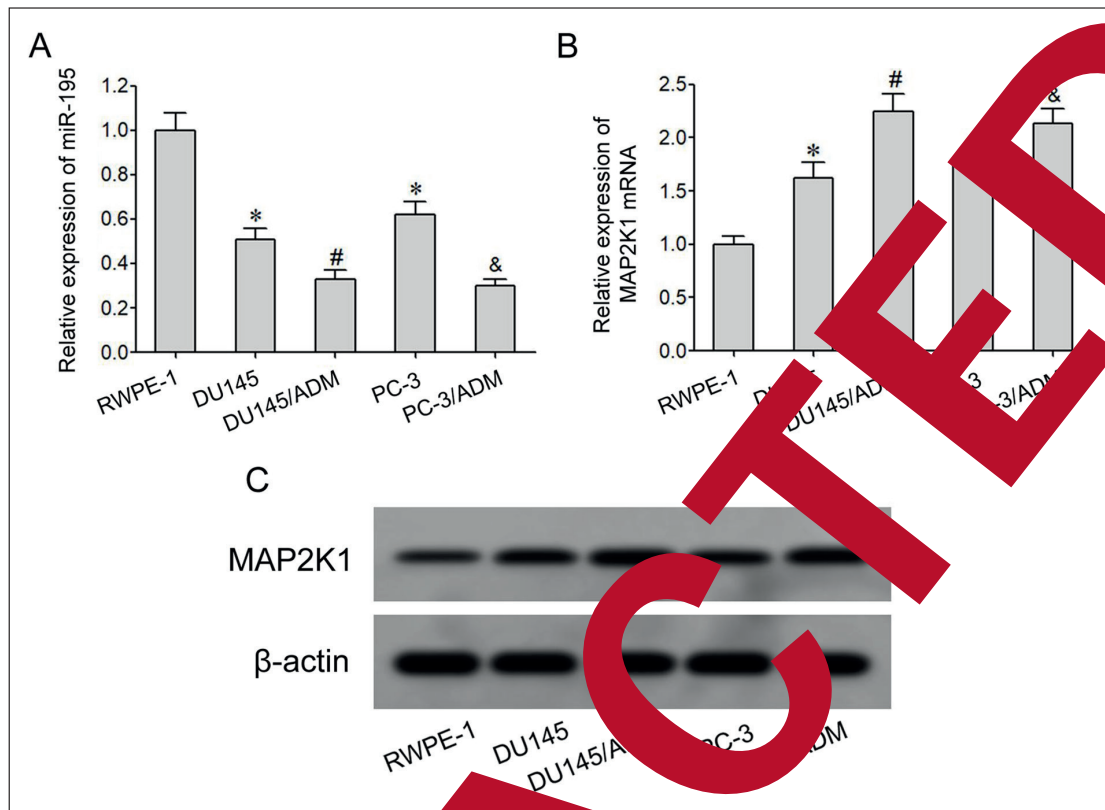


Figure 2. Down-regulation of miR-195 and up-regulation of MAP2K1 mRNA and protein in drug-resistant prostate cancer cell line. **A**, qRT-PCR for miR-195 expression. **B**, qRT-PCR measuring MAP2K1 mRNA. **C**, Western blot for protein expression. * $p < 0.05$ compared to RWPE-1 cells, # $p < 0.05$ compared to DU145 cells, & $p < 0.05$ compared to PC-3 cells.

sion, and drug-resistant cell PC-3/ADM had much higher MAP2K1 protein levels (Figure 2C).

Drug Resistant Cells Presented Resistance Against ADM Induced Cell Apoptosis

Under the treatment of ADM with concentrations equivalent to IC_{50} values of DU145 cells (7.51 μ g/ml), DU145 cells presented prominent cell apoptosis, whilst DU145/ADM cells had relatively lower apoptosis (Figure 3A) plus significantly depressed caspase-3 activity (Figure 3B). Using 5.29 μ g/ml ADM equivalent to IC_{50} , PC-3 cells presented prominent cell apoptosis, whilst PC-3/ADM cell had relatively fewer apoptosis (Figure 3C) plus lower caspase-3 activity (Figure 3D).

Overexpression of MiR-195 can Suppress Drug Resistance of Prostate Cancer Cells by Down-regulating MAP2K1

Under treatment of 16 μ g/ml ADM, both DU145/ADM and PC-3/ADM cells presented extremely lower apoptotic rates (Figure 4B)

whilst proliferation potency was fruitful (Figure 4C). The qRT-PCR results showed that the transfection of miR-195 mimic into DU145/ADM and PC-3/ADM cells remarkably decreased the MAP2K1 protein expression compared to miR-NC group, and p-ERK1/2 protein expression was remarkably suppressed (Figure 4B). Flow cytometry results showed that the transfection of miR-195 mimic remarkably elevated apoptosis of DU145/ADM and PC-3/ADM cells under 16 μ g/ml Adm treatment (Figure 4D), whilst cell proliferation potency was significantly depressed (Figure 4E, F).

Discussion

Prostate cancer is the malignant tumor originated from prostate gland epithelium. Having relatively higher malignancy and mortality rate, prostate cancer is the second popular cancer in males, only lower than the most deadly pulmonary carcinoma^{15,16}. Although China has a relatively lower incidence of prostate cancer than

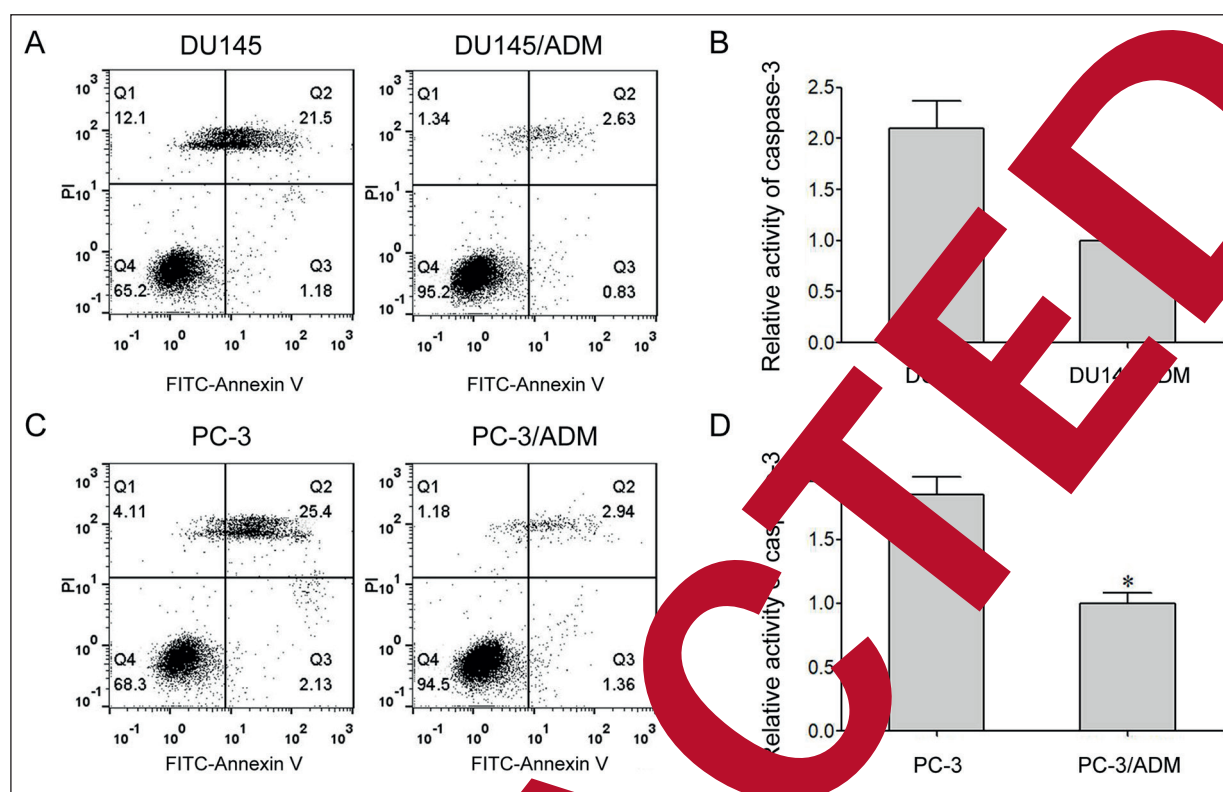


Figure 3. Drug-resistant cells presented resistance against induced cell apoptosis. **A**, Flow cytometry for DU145 and DU145/ADM cell apoptosis. **B**, Comparison of apoptosis rate between DU145 and DU145/ADM cells. **C**, Flow cytometry for apoptotic rate of PC-3 and PC-3/ADM cell. **D**, Statistical analysis of apoptotic rates between PC-3 and PC-3/ADM cells. * $p < 0.05$ comparing the two groups.

Western countries, its incidence is progressively increasing due to population aging and lifestyle transition^{17,18}. Chemotherapy is the main method for treating prostate carcinoma, but drug resistance has become the major factor affecting treatment efficiency, leading to chemotherapy failure and unfavorable prognosis. Therefore, an investigation for molecular candidates with abnormal change during chemotherapy resistance is of critical importance for revealing the mechanism for drug resistance, suppressing drug resistance, improving treatment efficiency, guiding individualized treatment and improving survival and prognosis. Jun N-terminal kinase (JNK) and p38 and Erk1/2, big MAPK kinase 1 (BMK1) are the major transduction pathways for MAPK signaling pathway. Among those, ERK-induced MAPK signaling transduction pathway is the classical MAPK signaling transduction pathway, and is the major transduction pathway through which the ERK signal pathway exerts its roles¹⁹⁻²¹. The ERK/MAPK signal transduction pathway is widely expressed in various tissues and cells and

can regulate various biological processes including cell proliferation, apoptosis and invasion²²⁻²⁵. It is closely correlated with tumor pathogenesis, progression and drug resistance²⁶⁻²⁸. MAP2K1 is a dual-specific protein kinase that exerts functions upstream of ERK protein and can phosphorylate tyrosine/threonine residues of substrate ERK protein, thus activating the ERK/MAPK signal pathway^{29,30}. Previous studies^{8,9} showed that enhanced expression or function of MAP2K1 are correlated with malignant properties of prostate cancer such as drug resistance and invasion. MiR-195 is a widely studied microRNA molecule, and its abnormal expression plays roles in the onset and progression of multiple tumors including lung cancer³¹, colorectal carcinoma³², and pancreatic cancer³³. Previous studies¹²⁻¹⁴ demonstrated the correlation between miR-195 down-regulation and the onset, progression, prognosis and drug resistance of prostate cancer. Bioinformatics analysis showed the complementary binding sites between miR-195 and 3'-UTR of MAP2K1 mRNA, suggesting potentially tar-

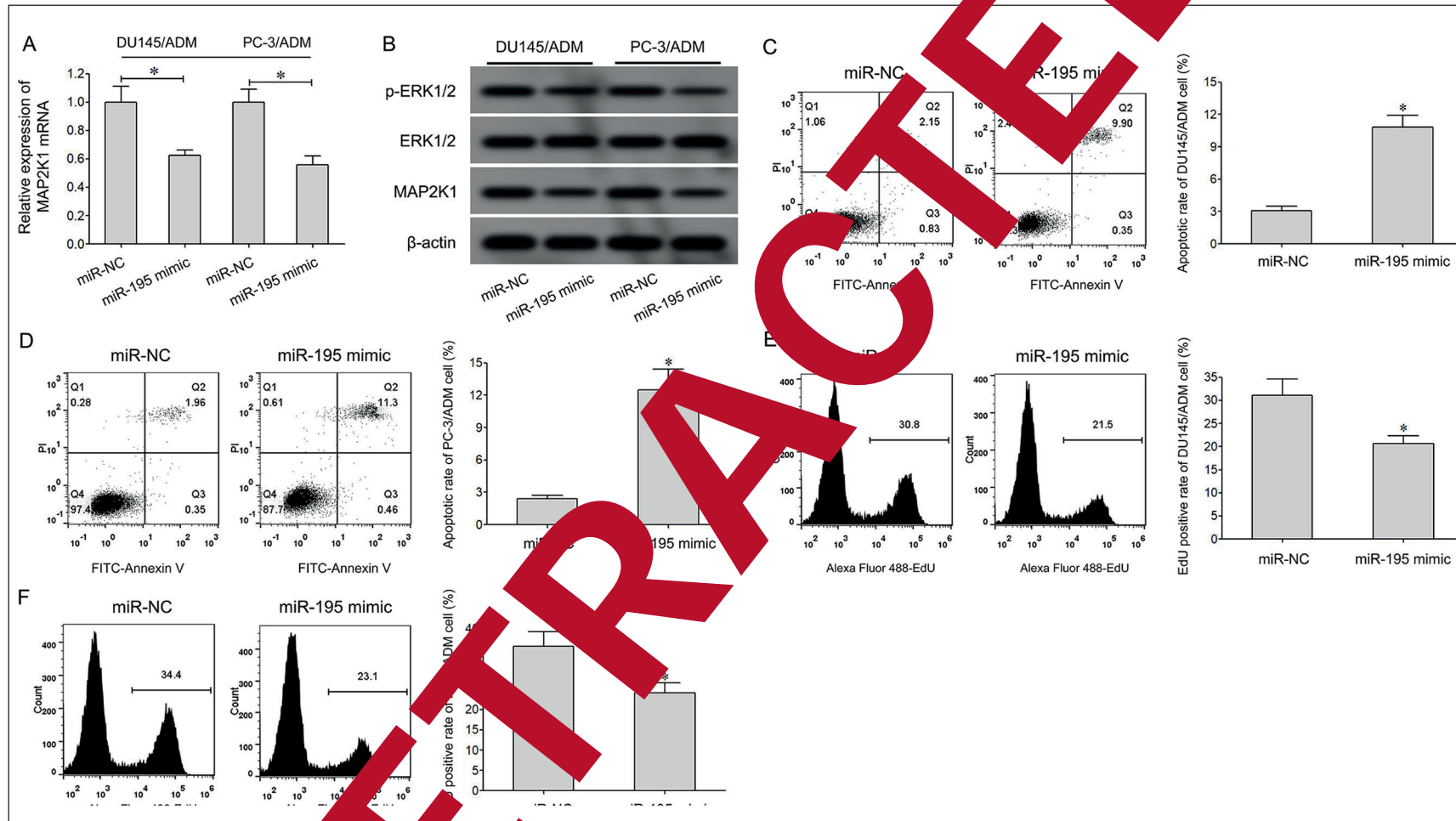


Figure 4. Over-expression of miR-195 decreases drug resistance of prostate cancer cell by down-regulating MAP2K1. **A**, qRT-PCR for measuring MAP2K1 mRNA expression. **B**, Western blot for protein expression. **C**, Flow cytometry for apoptosis of DU145/ADM cells. **D**, Flow cytometry for PC-3/ADM cell apoptosis. **E**, Flow cytometry for DU145/ADM cell proliferation. **F**, Flow cytometry for PC-3/ADM cell proliferation. * $p < 0.05$ comparing the two groups.

geted regulation between them. This work investigated whether miR-195 played a role in mediating MAP2K1 expression and affecting proliferation, apoptosis and ADM resistance of prostate cancer cells. Dual-Luciferase gene reporter assay showed that the transfection of miR-124 mimic remarkably decreased relative Luciferase activity of HEK293T cells transfected with pLUC-MAP2K1-WT plasmid, suggesting the targeted regulation between miR-195 and MAP2K1 mRNA. CCK-8 assay established that drug-resistant prostate cancer cell line DU145/ADM had remarkably higher IC_{50} values than DU145 cells. Similarly, PC-3/ADM cells also had remarkably higher IC_{50} values than parental PC-3 cells, suggesting the acquisition of ADM resistance in DU145/ADM and PC-3/ADM cells. Flow cytometry for apoptosis also revealed remarkably lower sensitivity of ADM-induced cell apoptosis in DU145/ADM and PC-3/ADM cells, producing apoptotic resistance. Both proliferation and apoptosis assay showed the successful generation of prostate cancer cell lines with ADM drug resistance that can satisfy the requirement of further experiments. Compared to normal prostate epithelial cell line RWPE-1, prostate cancer cell lines DU145 and PC-3 showed significantly decreased miR-195 expression, which was further down-regulated in drug-resistant cell lines DU145/ADM and PC-3/ADM. Prostate cancer cell line also showed significantly higher MAP2K1 mRNA and protein expression than normal prostate gland epithelial cells, and drug-resistant cells had higher miR-195 expression than parental normal cells. The results showed that the down-regulation of miR-195 played important roles in mediating MAP2K1 expression and its participation in the regulation of prostate cancer cell drug resistance in addition to prostate cancer. In a correlation study between miR-195 and prostate cancer, Guo et al³⁴ found significantly decreased miR-195 expression in tumor tissues, and its potency as the predictive factor for patient survival and prognosis. Zhang et al³³ shows that, compared to low-grade prostate cancer tissues, high-grade cancer tissues presented miR-195 down-regulation, while down-regulation is related to worse survival and prognosis. Cai et al¹⁴ demonstrated that, compared to normal prostate epithelial cell PrEC, prostate carcinoma cells PC-3, LNCaP and DU145 showed remarkably decreased miR-195 expression, plus miR-195 down-regulation in prostate cancer patients and the correlation between

miR-195 low expression and unfavorable survival or prognosis of patients ($p = 0.009$). This study indicated the participation of miR-195 down-regulation in the pathogenesis of prostate cancer, as our results also showed. Currently, a few studies have been performed regarding the regulation of prostate cancer cell's drug resistance by miR-195. Ma et al¹² found that, compared to drug-resistant cell DU15, docetaxel (DOC) resistant cell DU15/DOC had remarkably increased miR-195 expression, indicating that miR-195 down-regulation was a regulatory factor for drug resistance of prostate cancer cells, further supporting our study. To further investigate whether miR-195 played a role in mediating MAP2K1 expression and drug resistance of prostate cancer cells, this study over-expressed miR-195 in established prostate cancer cells with drug resistance to observe biological effects. The results showed that over-expression of miR-195 mimic remarkably increased the expression of the MAP2K1 protein in DU145/ADM and PC-3/ADM cells, and suppressed ERK1/2-MAPK pathway activity, leading to significantly higher apoptosis of those drug-resistant cells that originally have stable biological properties. In addition, over-expression of miR-195 inhibited cell growth in ADM-containing medium; in addition, it suppressed cell proliferation and malignant biological properties. Cai et al¹⁴ showed that the up-regulation of miR-195 can suppress migration or invasion potency of prostate cancer cells LNCaP and DU145 via targeted inhibition of RPS6KB1 expression to facilitate cell apoptosis, and can inhibit growth or tumorigenesis of tumor cells in BALB/c nude mice to suppress cell invasion or infiltration. Guo et al³⁴ found that the over-expression of miR-195 targets and inhibits BCOX1 gene expression, suppress proliferation, migration or invasion of *in vitro* cultured prostate cancer cells PC-3 or LNCaP, thus weakening *in vivo* growth, tumorigenesis and distal metastasis potency of those cells. Wu et al³⁵ found that the over-expression of miR-195 could target and regulate Fra-1 to weaken migration or invasion potency of prostate cancer cells PC-3 and DU145. It has been found that the over-expression of miR-195 could target FGF2 to suppress epithelial-mesenchymal transition (EMT) process of PC-3 or DU145 cells, thus weakening migration or invasion of cells. These studies all confirmed the role of miR-195 in weakening malignant biological properties of prostate cancer cells, supporting our results. In a study for the miR-195 regulation of prostate cancer cell's drug resistance, Ma et al¹² found that miR-195

played a role in targeted regulation of CLU, and over-expression of miR-195 can target and inhibit CLU expression to facilitate apoptosis of drug-resistant cells DU145/DOC and inhibit its clonal formation ability, thus suppressing DOC resistance of cells. In contrast to this study, we identified the role of miR-195 in targeted inhibition of the MAP2K1 expression, suppression of ERK/MAPK signal pathway transduction and weakening of prostate cancer cell's drug resistance, all of which have not been previously documented. However, this work has certain limitations as whether the regulation of MAP2K1 expression by miR-195 is correlated with drug resistance of prostate cancer patients is still unclear, which requires further assays for describing differential expression of miR-195 and MAP2K1 in tumor tissues from chemotherapy-sensitive and resistant patients.

Conclusions

We showed that the down-regulation of miR-195 was correlated with ADM resistance of prostate cancer cells. The up-regulation of miR-195 can weaken proliferation potency of drug-resistant prostate cancer cells *via* targeted inhibition on MAP2K1 expression and weakening of the ERK/AMPK signal pathway to facilitate cell apoptosis and suppress ADM.

Conflict of Interest

The Authors declare that they have no conflict of interest.

Acknowledgements

This work was supported by the Fujian Provincial Natural Science Foundation of China (2018J01282).

References

- 1) WANG W, WANG XX, FU H, TANG YC, MENG BQ, CHEN CL. Clinical diagnostic value of PSA combined miR-155 expression in prostate cancer. *Eur Rev Med Pharmacol* 2017; 22: 1615-1621.
- 2) COOPERBERG M, HAN JM. Epidemiology of prostate cancer. *World J Urol* 2017; 35: 849.
- 3) CHEN CA, KYRIANOU N. Profiling prostate cancer drug resistance. *Int J Mol Sci* 2018; 19: E904.
- 4) LIU Z. Molecular mechanisms of enzalutamide resistance in prostate cancer. *Curr Mol Biol Rep* 2017; 3: 230-235.
- 5) LI L, WANG J, ZHANG Y, MA L, WENG W, QIAO Y, XIAO W, WANG H, YU W, PAN Q. MEK1 protein expression and their interaction is critical for chemoresistance in liver cancer. *FEBS letters* 2013; 587: 3921-3927.
- 6) PENZVALTO Z, LANCZKY A, LENKAI Z, MEGGYESHAZI N, KRENACS T, SZOBOSZLAI N, DENKERTI Z, GYORFFY B. MEK1 is associated with carboplatin resistance and is a prognostic biomarker in epithelial ovarian cancer. *BMC Cancer* 2014; 14: 837.
- 7) GYSIN S, PAQUETTE J, MAHON M. Analysis of miRNA profiles after EGFR inhibition in human pancreatic cancer cells reveals pathways involved in drug sensitivity. *Cancers* 2012; 10: 1607-1619.
- 8) KATO T, FUJIMOTO Y, NAKANE K, MIYAMOTO T, TERAZAWA R, EHARA T, KAWANO Y, KOJIMA T, TERAZAWA Y, DEGUCHI T. C/EBP β /CCN3 interaction promotes invasion of taxane-resistant prostate cancer cells by increasing secretion of MMPs 2/9 and by activating PI3K and Rac signaling. *Cytokine* 2013; 64: 251-257.
- 9) LESCARBEAU RM, SEIB FP, PREWITZ M, WERNER C, KAPLAN DL. In vivo model of metastasis to bone marrow mediates prostate cancer castration resistant growth through paracrine and extracellular matrix interactions. *PLoS one* 2012; 7: e40372.
- 10) ANTONIOU MI, LIU C, LOU W, LOMBARD AP, EVANS CP, GAO AC. MicroRNA-181a promotes docetaxel resistance in prostate cancer cells. *Prostate* 2017; 67: 1020-1028.
- 11) LIU MAHATO RI. MicroRNAs and drug resistance in prostate cancers. *Mol Pharm* 2014; 11: 2539-2552.
- 12) MA X, ZOU L, LI X, CHEN Z, LIN Q, WU X. MicroRNA-195 regulates docetaxel resistance by targeting clusterin in prostate cancer. *Biomed Pharmacother* 2018; 99: 445-450.
- 13) ZHANG X, TAO T, LIU C, GUAN H, HUANG Y, XU B, CHEN M. Downregulation of miR-195 promotes prostate cancer progression by targeting HMGA1. *Oncol Rep* 2016; 36: 376-382.
- 14) CAI C, CHEN QB, HAN ZD, ZHANG YQ, HE HC, CHEN JH, CHEN YR, YANG SB, WU YD, ZENG YR. miR-195 inhibits tumor progression by targeting RPS6KB1 in human prostate cancer. *Clin Cancer Res* 2015; 21: 4922-4934.
- 15) ARCANGELI S, PINZI V, ARCANGELI G. Epidemiology of prostate cancer and treatment remarks. *World J Radiol* 2012; 4: 241-246.
- 16) BRAWLEY OW. Prostate cancer epidemiology in the United States. *World J Urol* 2012; 30: 195-200.
- 17) FENNER A. Prostate cancer: ERSPC calculator recalibrated for China. *Nat Rev Urol* 2017; 14: 66.
- 18) REN SC, CHEN R, SUN YH. Prostate cancer research in China. *Asian J Androl* 2013; 15: 350-353.
- 19) WANG DW, WANG YQ, SHU HS. MiR-16 inhibits pituitary adenoma cell proliferation via the suppression of ERK/MAPK signal pathway. *Eur Rev Med Pharmacol Sci* 2018; 22: 1241-1248.

- 20) LI B, LIU YH, SUN AG, HUAN LC, LI HD, LIU DM. MiR-130b functions as a tumor promoter in glioma via regulation of ERK/MAPK pathway. *Eur Rev Med Pharmacol Sci* 2017; 21: 2840-2846.
- 21) BUCHEGGER K, SILVA R, LOPEZ J, ILI C, ARAYA JC, LEAL P, BREBI P, RIOUELME I, ROA JC. The ERK/MAPK pathway is overexpressed and activated in gallbladder cancer. *Pathol Res Pract* 2017; 213: 476-482.
- 22) ZHANG G, CHENG Y, ZHANG Q, LI X, ZHOU J, WANG J, WEI L. ATXLPA axis facilitates estrogen-induced endometrial cancer cell proliferation via MAPK/ERK signaling pathway. *Mol Med Rep* 2018; 17: 4245-4252.
- 23) LIAO T, WEN D, MA B, HU JQ, QU N, SHI RL, LIU L, GUAN Q, LI DS, JI QH. Yes-associated protein 1 promotes papillary thyroid cancer cell proliferation by activating the ERK/MAPK signaling pathway. *Oncotarget* 2017; 8: 11719-11728.
- 24) LU Y, LI Y, XIANG M, ZHOU J, CHEN J. Khat promotes human breast cancer MDA-MB-231 cell apoptosis via mitochondria and MAPK-associated pathways. *Oncol Lett* 2017; 14: 3947-3952.
- 25) YANG XL, LIU KY, LIN FJ, SHI HM, OU ZL. CCL28 promotes breast cancer growth and metastasis through MAPK-mediated cellular anti-apoptosis and pro-metastasis. *Oncol Rep* 2017; 38: 1393-1401.
- 26) ZHUANG ST, CAI Y, GAO HC, QIU JF, ZENG J, WANG WJ. Study on the function and mechanism of non-coding RNA DMTF1v4 in the occurrence of colon cancer. *Eur Rev Med Pharmacol Sci* 2018; 22: 3779-3788.
- 27) MCGIVERN N, EL-HELALI A, MULLAN P, McNEISH I, PAUL HARKIN D, KENNEDY RD, MCCABE N. Activation of MAPK signalling results in resistance to sunitinib (AZD0530) in ovarian cancer. *Oncotarget* 2018; 9: 4722-4736.
- 28) HEW KE, MILLER PC, EL-ASHRY D, SUN J, BESSER AH, INCE TA, GU M, WEI Z, ZHANG G, BRAFFORD A, LU Y, MILLS GB, SLINGERLAND JM, SIMON T. MEK activation predicts poor outcome in the MEK inhibitor, selumetinib, reverses estrogen resistance in ER-positive high-grade serous ovarian cancer. *Clin Cancer Res* 2016; 22: 945-947.
- 29) ZHAO L. Hirudin inhibits cell growth and ERK/MAPK signaling in human glioma. *Int J Exp Med* 2015; 8: 20983-20987.
- 30) CHIBA T, SOENO Y, SHIBATA Y, SUDO M, YAGISHITA H, KAWA Y, KAWASHIRI S, OHTSUKI MAI K. EMT1 Inhibition of oral carcinoma cell growth and ERK/MAPK activation. *J Dent Res* 2016; 95: 446-451.
- 31) YU X, ZHANG C, CAVAZOS D, MONTANO DU L, PERTSEMLIDIS A. miR-195 targets cyclin D1 and survivin to modulate tumorigenesis of non-small cell lung cancer. *Cell Pathol Dis* 2018; 9: 193.
- 32) FENG C, ZHANG L, SUN X, ZHAN L, LOU Y, WANG Y, LIU L, ZHANG Y. GDPase is a target of miR-195-5p, is associated with metastasis and chemoresistance in colorectal cancer. *Biomed Pharmacother* 2018; 101: 945-952.
- 33) ZHOU B, SUN C, LIU X, ZHAN H, ZOU H, FENG Y, QIU F, ZHANG S, WU L, WANG B. MicroRNA-195 suppresses the progression of pancreatic cancer by targeting BCL2. *Cell Physiol Biochem* 2017; 44: 1867-1874.
- 34) GUO J, WANG M, LIU X. MicroRNA-195 suppresses tumor cell proliferation and metastasis by directly targeting BCOX1 in prostate carcinoma. *J Exp Clin Cancer Res* 2015; 34: 91.
- 35) WU J, JI A, WANG X, ZHU Y, YU Y, LIN Y, LIU Y, LI S, LIANG Z, XU X, ZHENG X, XIE L. MicroRNA-195-5p, a new regulator of Fra-1, suppresses the migration and invasion of prostate cancer cells. *J Transl Med* 2015; 13: 289.