

# Increased miR-142 and decreased DJ-1 enhance the sensitivity of pancreatic cancer cell to adriamycin

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**Abstract.** – **OBJECTIVE:** Phosphatidylinositol-3 kinase (PI3K)/protein kinase B (AKT) signaling pathway is related to tumorigenesis by up-regulating survivin. Phosphatase and tensin homologue deleted on chromosome ten (PTEN) can suppress PI3K/AKT signaling pathway, while DJ-1 is the negative regulator of PTEN. DJ-1 up-regulation is closely correlated with tumor occurrence, progression, and drug resistance in pancreatic cancer. MicroRNA-142 (MiR-142) is significantly declined in pancreatic cancer tissue. Bioinformatics analysis demonstrated complementary binding site exists between miR-142 and DJ-1. This investigation therefore, aimed to study the role of miR-142 in the regulation of DJ-1-PTEN/PI3K/AKT/Survivin signaling pathway as well as in pancreatic cancer cell proliferation, apoptosis, and adriamycin (ADM) resistance.

**MATERIALS AND METHODS:** Dual luciferase assay was performed to assess the targeted relationship between miR-142 and DJ-1. MiR-142, DJ-1, and PTEN expressions in SW1990 cells and drug resistance SW1990/ADM cells were compared. SW1990/ADM cells were divided into five groups, including mimic miR-142 mimic, si-DJ-1, si-DJ-1 + miR-142 mimic, si-DJ-1 + miR-142 mimic + si-DJ-1 groups. DJ-1, PTEN, phosphorylated-AKT (p-AKT), and Survivin expressions were tested. Cell apoptosis was determined by flow cytometry. Cell proliferation was evaluated by EdU staining.

**RESULTS:** MiR-142 targeted inhibited DJ-1 expression. MiR-142, PTEN, and cell apoptosis significantly down-regulated, while DJ-1, p-AKT, and Survivin expressions significantly elevated in SW1990/ADM cells compared with SW1990 cells. MiR-142 mimics and/or si-DJ-1 treatment markedly reduced DJ-1, p-AKT, and Survivin expressions enhanced PTEN level, attenuated cell proliferation, enhanced cell apoptosis, and weakened ADM resistance.

**CONCLUSIONS:** MiR-142 over-expression weakened ADM resistance in pancreatic cancer cells by targeting DJ-1 to enhance PTEN expression and attenuate PI3K/AKT signaling pathway activity.

**Key Words:** MiR-142, DJ-1, PTEN, PI3K/AKT, Adriamycin, Pancreatic cancer, Drug resistance.

## Introduction

Pancreatic carcinoma (PC) is a type of malignancy with poor curative effect and prognosis<sup>1,2</sup>. Chemotherapy is an important method in the treatment of PC. However, drug resistance is an adverse factor that affects the curative effect<sup>3</sup>. Therefore, investigating the mechanism of drug resistance in the PC is of clinical significance to increase the effect of chemotherapy as well as improve the prognosis.

Survivin is a critical anti-apoptotic factor that participates in regulating cell proliferation<sup>4,5</sup>. It was found that Survivin is one of the target genes of phosphatidylinositol-3 kinase (PI3K)/protein kinase B (AKT/PKB) signaling pathway. PI3K/AKT signaling pathway activation promotes Survivin transcription and expression, suppressed cell apoptosis, and facilitated cell proliferation. It is closely correlated with various tumor occurrence, progression, and drug resistance<sup>6-8</sup>. Phosphatase and tensin homologue deleted on chromosome ten (PTEN) is a tumor suppressor gene that negatively regulates PI3K/AKT signaling pathway and tumorigenesis<sup>9-11</sup>. DJ-1 is a negative regulator of PTEN

that can enhance the activity of the PI3K/AKT signaling pathway<sup>12,13</sup>. DJ-1 is found abnormally increased in several tumor tissues<sup>14-16</sup>. It was showed that DJ-1 significantly elevated in the tumor tissue<sup>17</sup> and peripheral blood of PC patients, suggesting the potential tumor suppressor gene role of DJ-1 in PC tumorigenesis<sup>16,18</sup>. MiRNA is a type of endogenous single-strand non-coding RNA with a length of 22-25 nt which has been discovered from an eukaryote. It plays a degrading or inhibiting role in the transcription of more than 1/3 mRNAs by binding with the 3'-UTR. Its relationship with tumorigenesis attracts more and more attention<sup>19</sup>. MiR-142 significantly down-regulated in PC tissue and cells, revealing that miR-142 might be a tumor suppressor gene in PC<sup>20,21</sup>. Bioinformatics analysis revealed a complementary binding site between miR-142 and DJ-1. This study aimed to evaluate the role of miR-142 in the regulation of DJ-1 as well as in PC cell proliferation, apoptosis, and adriamycin (ADM) resistance by establishing ADM resistant PC cell line.

## Materials and Methods

### Main Reagents and Materials

Human normal pancreatic cell line HPAC15 was purchased from Aolu Biotechnology (Shanghai, China). Human PC cell line SW1990 was purchased from Baili Biotechnology (Shanghai, China). Roswell park memorial institute 1640 (RPMI-1640) medium, optional minimum essential (Opti-MEM), penicillin, streptomycin and adriamycin were purchased from Gibco BRL Co. Ltd. (Grand Island, NY, USA). Fetal bovine serum (FBS) was purchased from Science Co. Ltd. (Carlsbad, CA, USA). RNA extraction kit SPLIT RNA Extraction Kit was provided by Lexogen (Vienna, Austria). TransScript Green One-Step qRT-PCR SuperMix was purchased from TransGen Biotech (Beijing, China). miR-NC, miR-142 mimic, miR-142 inhibitor and EdU flow cytometry detection reagent were purchased from Ribobio (Guangzhou, China). Rabbit anti-human DJ-1, p-PTEN, and β-actin antibodies were got from Abcam Biotechnology (Cambridge, MA, USA). Mouse anti-human PTEN and β-actin antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Horseradish peroxidase (HRP) conjugated secondary antibody was derived from Bio-Rad Laboratories (Hercules, CA, USA). Annexin-V/propidium iodide (PI) apoptosis detection kit and cell counting kit 8 (CCK-8)

cell viability detection kit were purchased from Dojindo Laboratories (Kumamoto, Japan). TransFast™ Transfection Reagent was purchased from Promega (Madison, WI, USA). Luciferase reporter plasmid pLUC was purchased from GeneBioVector (Beijing, China). Luciferase activity analysis kit LightSwitch was obtained from Switch Motif (Saranac Lake, NY, USA). Adriamycin was purchased from Meiji Seika (Chugoku, Tokyo, Japan).

### Cell Culture

HPC-Y5 and SW1990 cells were cultured in RPMI-1640 medium which contains 10% FBS and 100 U/ml penicillin and 100 U/ml streptomycin at 37°C and 5% CO<sub>2</sub>. The cells were passaged at 1:4. This investigation has been approved by the Ethics Committee of Hong Kong Hospital Affiliated to Mudanjiang Medical University, Mudanjiang, China.

### SW1990/ADM Model Establishment

SW1990 cells in logarithmic phase were treated by 0.1 μg/ml ADM for 24 h. Then, the medium was changed until the cells can grow and passaged in medium stably. Next, the concentration of ADM was gradually increased to 0.2 μg/ml, 0.4 μg/ml, 0.8 μg/ml, 1.6 μg/ml, and 3.2 μg/ml until the cell can grow stably in 3.2 μg/ml ADM to establish a cisplatin-resistant SW1990/ADM cell line. SW1990 and SW1990/ADM cells were treated by ADM at 0, 0.25, 0.5, 1, 2, 4, 8, 16, 32, and 64 μg/ml. After 48 h, 10 μl CCK-8 cells were added into the cells for 4 h and tested at 450 nm to obtain the absorbance value (A450). Inhibitory rate = (1- ADM group A450)/control A450 × 100%. IC<sub>50</sub> was calculated by SPSS software (SPSS, Inc., Chicago, IL, USA). Resistance index (RI) = IC<sub>50</sub> of SW1990/ADM/IC<sub>50</sub> of SW1990.

### Dual-Luciferase Assay

The PCR products containing the full length of DJ-1 gene 3'-UTR or mutant segment were cloned into pLUC followed by being transformed into DH5α competent cells and subsequent selection of the plasmid with the correct sequence. Then pLUC-DJ-1-wt (or pLUC-DJ-1-mut) was co-transfected into HEK293T cells using TransFast™ Transfection Reagent together with miR-142 mimic (or miR-142 inhibitor, or miR-NC). The luciferase activity was measured according to the Light Switch manual after cultured for 48 h.

### Cell Transfection and Grouping

SW1990/ADM cells were divided into five groups, including mimic-NC, miR-142 mimic,

**Table 1.** Primers for the PCR.

Genes		Primers
miR-142	Forward	5'-GCCACAAGGAGGGCTGGGG
	Reverse	5'-GAGCGCCGAGGAAGATGCGGC-3'
DJ-1	Forward	5'-CGGGGTGCAGGCTTCGAA-3'
	Reverse	5'-TCCGGTTTTCTGCTGCTTC-3'
PTEN	Forward	5'-CACACGACGGGAAGACGATTC-3'
	Reverse	5'-CCTCTGGTCTGCTGCTATGAAATGG-3'
GAPDH	Forward	5'-ACACCCACTCCTCCACCTTCTT-3'
	Reverse	5'-TCCACCACCGGTTGCTGTAG-3'

si-NC, si-DJ-1, and miR-142 mimic + si-DJ-1 groups. Nucleotide fragments and TransFast™ Transfection Reagent were added to Opti-MEM for incubation at room temperature for 20 min, respectively. After that, they were added to the cells and cultured in Opti-MEM. After incubation for 6 h, the medium was changed to RPMI-1640 medium containing 10% FBS and 1% penicillin-streptomycin and further cultured for 48 h followed by detection.

#### Quantitative Real Time-PCR (qRT-PCR)

Total RNA was extracted using SPLIT™ RNA Extraction Kit for PCR analysis using TaqMan Script Green One-Step qRT-PCR SuperMix. The reaction system consisted of 1 µg RNA template, 0.3 µM primers, 10 µl 2×TransStart Tip Green qPCR SuperMix, 0.4 µl RT Enzyme Mix, 0.4 µl Dye II, and ddH<sub>2</sub>O. The reverse transcription condition was 37°C for 15 min and 98°C for 5 min. The PCR reaction was performed on ABI 7500 (Applied Biosystems, Foster City, CA, USA) at the condition as follows: 45°C reverse transcription for 5 min, 94°C pre-denaturation for 30 s, followed by 40 cycles of 94°C for 5 s, and 60°C for 30 s. The primers were listed in Table 1.

#### Western blot

Total protein was extracted using radioimmuno-precipitation assay (RIPA) lysis buffer. A total of 50 µg protein was separated by 12% Sodium Dodecyl Sulfate-Polyacrylamide Gel Electrophoresis (SDS-PAGE) for 3 h and transferred to a nitrocellulose membrane. After that, the membrane was blocked and incubated with primary antibody at 4°C overnight (DJ-1, PTEN, p-AKT, Survivin, and p-ERK1/2) at dilutions of 1:100, 1:300, 1:200, 1:300, and 1:800, respectively) followed by incubation with secondary antibody (1:10000) for 60 min after washed by Phosphate Buffered Saline Tween-20 (PBST) for three times. At last, the protein expression

was detected using enhanced chemiluminescence (ECL) (Amersham Biosciences, Little Chalfont, Buckinghamshire, England).

#### Flow Cytometry

Cells were re-suspended in 100 µl binding buffer and incubated with 5 µl Annexin V-FITC and 5 µl PI for 15 min under dark. Then, 400 µl Annexin V Binding Buffer was added into the cells and analyzed by analysis of cell apoptosis by FC 500 MCL/MPL flow cytometer (Beckman Coulter Inc., Brea, CA, USA).

#### EdU Staining

15 µM EdU solution was added into cells and incubated at 37°C for 60 min. After incubated for 48 h, cells were digested by trypsin and fixed in 4% paraformaldehyde (Sigma-Aldrich, St. Louis, MO, USA), followed by incubation in 1% saponin and suspension in phosphate buffered saline (PBS). At last, cells were stained with 500 µl 6-FAM Azide at room temperature under dark for 30 min and tested on FC 500 MCL/MPL flow cytometer r (Beckman Coulter Inc., Brea, CA, USA).

#### Statistical Analysis

All data analyses were performed by SPSS 18.0 software (SPSS, Inc., Chicago, IL, USA). The measurement data were presented as mean ± standard deviation (SD). The Student's *t*-test was used to compare the differences between the two groups. Tukey's post hoc test was used to validate the ANOVA for comparing measurement data among groups. *p*<0.05 was considered as statistical significance.

## Results

### MiR-142 Regulates DJ-1 Expression

MicroRNA.org online prediction revealed a targeted binding site between miR-142 and 3'-

**Table II.** IC50 of SW1990/ADM cells

Group	IC50 (µg/ml)
miR-NC	22.71±1.13
miR-142 mimic	9.14±0.76a
si-NC	24.28±1.09
si-DJ-1	8.65±0.62b
miR-142 mimic+ si-DJ-1	5.75±0.45cd

<sup>a</sup>*p*<0.05, compared with miR-NC, <sup>b</sup>*p*<0.05, compared with si-NC, <sup>c</sup>*p*<0.05, compared with miR-142 mimic, <sup>d</sup>*p*<0.05, compared with si-DJ-1 group.

UTR of DJ-1 mRNA (Figure 1A). Dual luciferase assay showed that miR-142 mimics or inhibitor transfection significantly decreased or increased the relative luciferase activity of HEK293 cells transfected by pLUC-DJ-1-wt, while it showed no statistical effect on the luciferase activity in HEK293 cells which were transfected by pLUC-DJ-1-mut (Figure 1B), suggesting a regulatory relationship between miR-142 and DJ-1 mRNA.

**MiR-142 Down-Regulated, While DJ-1 Over-Expressed in SW1990/ADM Cells**

The IC<sub>50</sub> of SW1990 cells was 0.86 µg/ml, while it was 23.75 µg/ml in SW1990/ADM cells. The RI was 27.61. The qRT-PCR showed that miR-142 and PTEN mRNA levels were significantly lower, while DJ-1 mRNA level was markedly higher in SW1990 cells compared with HPC-Y5

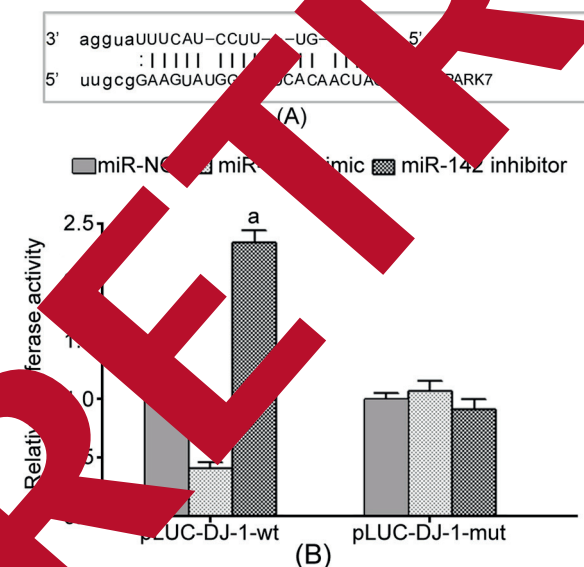
cells. Similar results were observed in SW1990/ADM cells compared with SW1990 (Figure 2A). Western blot showed that DJ-1 protein expression was markedly higher, while PTEN protein expression was significantly lower in SW1990/ADM cells compared with SW1990 cells and HPC-Y5 cells (Figure 2B). Flow cytometry demonstrated that the cell apoptotic rate was significantly lower (Figure 2C), while cell proliferation (Figure 2D) was significantly stronger in SW1990/ADM cells than that in SW1990 cells treated by 0.86 µg/ml ADM.

**MiR-142 Over-Expression Attenuates ADM Resistance in SW1990/ADM Cells**

3.2 µg/ml ADM was used to evaluate the impact of miR-142 regulation and/or DJ-1 reduction on the ADM sensitivity of SW1990/ADM cells. MiR-142 mimic and/or si-DJ-1 transfection significantly enhanced PI3K level, reduced p-AKT and Survivin expressions (Figure 3A and B), attenuated cell proliferation (Figure 3C), enhanced cell apoptosis (Figure 3D), and weakened ADM resistance (Table II).

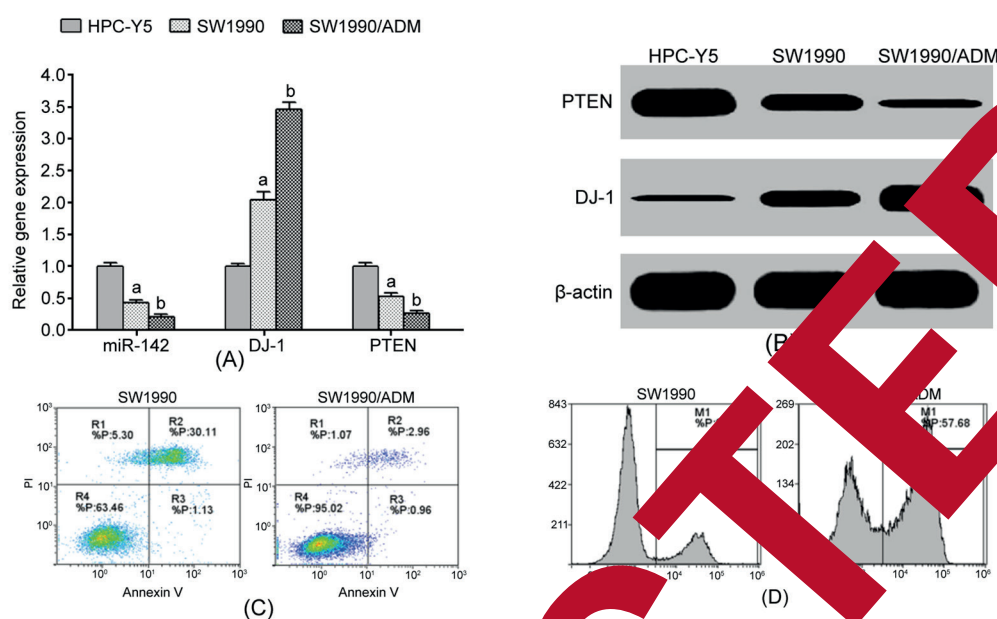
**Discussion**

The PI3K/AKT signaling pathway is involved in multiple biological behaviors, such as cell proliferation, apoptosis, cell cycle, etc. It is closely associated with embryonic development, angiogenesis, and tumorigenesis. The kinase activity of PI3K phosphorylates PIP2 to PIP3, which phosphorylates AKT at Ser473 and Thr308 with the help of 3-phosphoinositide-dependent protein kinase 1 (PDK1) and 3-phosphoinositide-dependent protein kinase 2 (PDK2). Phosphorylated AKT further participates in gene transcription and translation that regulates cell proliferation, cell cycle, apoptosis, and invasion<sup>22</sup>. As the strongest factor in inhibitor of apoptosis protein (IAPs) family, Survivin antagonizes cell apoptosis by suppressing Caspase-3 and Caspase-7 activities<sup>4</sup>. In addition, Survivin up-regulation is related to the enhancement of cell proliferation<sup>5</sup>. Survivin is one of the important target genes of PI3K/AKT signaling pathway. PI3K/AKT pathway promotes Survivin gene transcription and expression, and inhibits cell apoptosis, and accelerates cell proliferation. It is related to multiple cancers occurrence, progression, and drug-resistance, including gastric cancer<sup>6</sup>, endometrial cancer<sup>7</sup>, and colorectal cancer<sup>8</sup>. PTEN is the only discovered tumor suppressor gene with the dual activity of



**Figure 1.** MiR-142 targeted regulated DJ-1 expression. **A**, The binding site between miR-142 the 3'-UTR of DJ-1 mRNA. **B**, Dual luciferase assay. \**p*<0.05, compared with miR-NC.

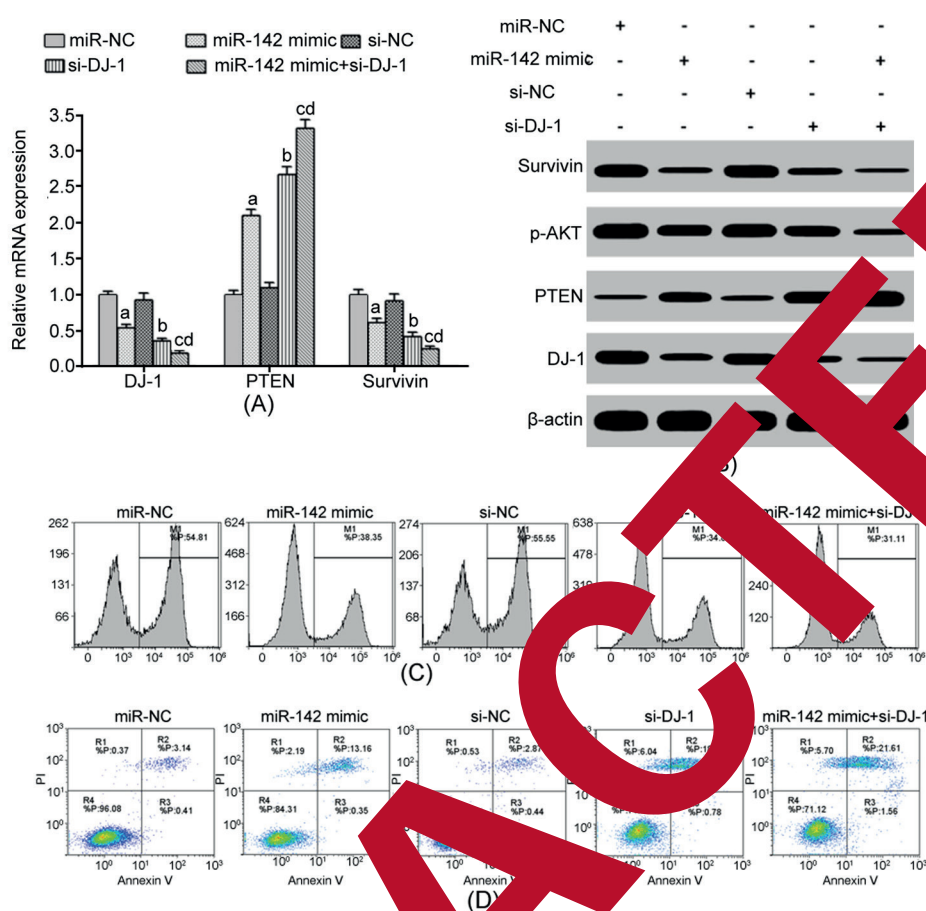




**Figure 2.** MiR-142 down-regulated, while DJ-1 over-expressed in SW1990/ADM cells. **A**, qRT-PCR detection of gene expression. **B**, Western blot detection of protein expression. **C**, Flow cytometry detection of cell apoptosis. **D**, EdU staining detection of cell proliferation. <sup>a</sup> $p < 0.05$ , compared with HPC-Y5 cells, <sup>b</sup> $p < 0.05$ , compared with SW1990 cells.

protease and phosphatase. It plays an important role in tumorigenesis through negatively regulating PI3K/AKT signaling pathway, such as in bladder carcinoma<sup>9</sup>, breast cancer<sup>10</sup>, and prostate cancer<sup>11</sup>. DJ-1, also known as Parkinson gene (PARK7), is related to Human autosomal recessive inheritance early-onset Parkinson's disease (PD)<sup>23,24</sup>. DJ-1 involves multiple biological processes, such as anti-apoptosis, chaperone, cell proliferation, apoptosis, transformation, and androgen receptor signaling transduction<sup>25,26</sup>. DJ-1 enhances the activity of the PI3K/AKT signaling pathway by attenuating the inhibitory effect of PTEN on PI3K/AKT<sup>12,13</sup>. DJ-1 up-regulation is found in many tumor tissues, such as breast cancer<sup>14</sup> and lung cancer<sup>15</sup>. It was shown that DJ-1 significantly elevated in the tumor cells and peripheral blood<sup>16,18</sup> of PC patients, suggesting the potential oncogene role of DJ-1 in PC tumorigenesis. MiR-142 significantly down-regulated in PC tissue and cells, revealing that miR-142 might be a tumor suppressor gene in PC. Bioinformatics analysis revealed a complementary binding site between miR-142 and DJ-1. Establishment of drug-resistant cancer cell lines *in vitro* is of great significance in exploring the mechanism of cancer resistance and screening chemotherapy drugs. This study investigated the role of miR-142 in the regulation of DJ-1, PC cell

proliferation, apoptosis, as well as ADM resistance. Establishing ADM resistant PC cell line. Dual luciferase assay showed that miR-142 mimics or inhibitor transfection significantly reduced or increased the relative luciferase activity of HEK293 cells transfected by pLUC-DJ-1-wt, while it did not exhibit any statistical effect on the luciferase activity in HEK293 cells transfected by pLUC-DJ-1-mut, suggesting a regulatory relationship between miR-142 and DJ-1 mRNA. MiR-142 and PTEN mRNA levels were significantly lower, while DJ-1 mRNA expression was significantly higher in SW1990 cells compared with HPC-Y5 cells. Similar results were observed in SW1990/ADM cells compared with SW1990. It indicated that miR-142 and PTEN reduction, and DJ-1 elevation are related not only to PC tumorigenesis, but also ADM resistance enhancement. Lu et al<sup>20</sup> reported that miR-142 significantly declined in PC cell line PANC-1, SW1990, Hup, and CFPAC-1 compared with normal pancreatic cell HPC-Y5. Moreover, miR-142 significantly declined in PC tumor tissue compared with adjacent normal control. In this study, the miR-142 level was markedly lower in PC cells than normal pancreatic cells, suggesting that miR-142 down-regulation was a tumor-promoting factor in PC, which was similar with Lu et al<sup>20</sup> findings. Ohuchida et al<sup>21</sup> established



**Figure 3.** MiR-142 over-expression attenuates ADM resistance in SW1990/ADM cells. **A**, qRT-PCR detection of gene expression. **B**, Western blot detection of protein expression. **C**, Flow cytometry detection of cell apoptosis. **D**, EdU staining detection of cell proliferation. <sup>a</sup> $p < 0.05$ , compared with miR-NC, <sup>b</sup> $p < 0.05$ , compared with si-NC, <sup>c</sup> $p < 0.05$ , compared with miR-142 mimic, <sup>d</sup> $p < 0.05$ , compared with si-DJ-1 group.

Gemcitabine resistant cell line by using Gemcitabine to treat SW1990, SUIT-2 and PANC-1. It was found that miR-142 significantly decreased in Gemcitabine resistant cell line compared with parental drug-sensitive cells. In this work, miR-142 over-expression in SW1990 cells was markedly lower than HPC-Y5, revealing that miR-142 over-expression may be associated with PC drug resistance, which was in accordance with Ohuchida et al<sup>21</sup> findings. Chen et al<sup>18</sup> discovered that DJ-1 was apparently up-regulated in the peripheral blood of PC patients compared with healthy control. Tsiaousidou et al<sup>17</sup> demonstrated that DJ-1 was normally elevated in PC tumor tissue and was negatively correlated with chemotherapy sensitivity, indicating that DJ-1 over-expression was associated with PC tumorigenesis and drug resistance. In this study, DJ-1 increased in PC cells, while its increasing amplitude was larger

in drug-resistant cells, suggesting that DJ-1 was a promoting factor in PC occurrence and drug resistance, which was confirmed by Chen et al<sup>18</sup> and Tsiaousidou et al<sup>17</sup>. Further investigation showed that miR-142 mimics and/or si-DJ-1 transfection markedly enhanced PTEN level, reduced p-AKT and Survivin expressions, attenuated cell proliferation, enhanced cell apoptosis, and weakened ADM resistance. Lu et al<sup>20</sup> showed that miR-142 declined, while PC cell proliferation and invasion enhanced under hypoxia condition. MiR-142 over-expression attenuated PC cells SW1990 and PANC-1 proliferation, EMT, and invasion by restraining hypoxia-inducible factor 1 (HIF-1 $\alpha$ ). MacKenzie et al<sup>27</sup> exhibited that miR-142 up-regulation weakened PC cells MIA PaCa-2 and Capan-1 proliferation through targeting HSP-70 expression. Ohuchida et al<sup>21</sup> also showed that miR-142 level was significantly higher in patients

with better survival and prognosis after Gemcitabine treatment, suggesting that miR-142 may be related to drug resistance. This research found that miR-142 elevation reduced PC malignancy and attenuated drug resistance, which was supported by MacKenzie et al<sup>27</sup> and Ohuchida et al<sup>21</sup>. Chen et al [18] reported that DJ-1 level was higher in Gemcitabine resistant MIA PaCa-2 cells compared with parental cells, while down-regulation of DJ-1 enhanced Gemcitabine sensitivity and apoptosis in MIA PaCa-2 cells, which was similar with our results. We revealed that miR-142 reduction plays a role in down-regulating PTEN, enhancing PI3K/AKT signaling pathway, and promoting PC drug resistance.

## Conclusions

We found that miR-142 down-regulation and DJ-1 over-expression are associated with DM resistance in PC cells. MiR-142 over-expression weakened ADM resistance in pancreatic cancer cells by targeting DJ-1 to enhance PTEN expression and attenuate PI3K/AKT signaling pathway activity.

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## Conflict of Interest

The Authors declare that they have no conflict of interest.

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RETRACTED