

## The Effect of Gemcitabine on Cell Cycle Arrest and microRNA Signatures in Pancreatic Cancer Cells

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**Abstract.** *Background/Aim: Gemcitabine, an inhibitor of DNA synthesis, is the gold standard chemotherapeutic agent for pancreatic ductal adenocarcinoma (PDAC). MicroRNAs (miRNAs) play critical roles in cancers, including PDAC. However, less is known about the effect of gemcitabine on PDAC cells and miRNA expression in PDAC. We evaluated the effect of gemcitabine on the cell cycle of PDAC cells in vitro and in vivo and on the miRNA expression profile. Materials and Methods: Effects of gemcitabine on PK-1 and PK-9 cell growth were evaluated using a cell counting kit-8 assay. Xenografted mouse models were used to assess gemcitabine effects in vivo. Results: Gemcitabine inhibited the proliferation and tumour growth of PK-1 cells, and induced S phase cell cycle arrest. Numerous miRNAs were altered upon gemcitabine treatment of PK-1 cells and xenograft models. Conclusion: Altered miRNAs may serve as potential therapeutic targets for improving the efficacy of gemcitabine in PDAC.*

Pancreatic ductal adenocarcinoma (PDAC) is the seventh leading cause of cancer-related death in both men and women according to the 2017 Global Burden of Disease study (1). Most exocrine pancreatic cancers are adenocarcinomas arising from the ductal epithelium. Surgical resection is the only potential curative treatment for PDAC;

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however, only 15% to 20% of patients have resectable disease at initial diagnosis (2, 3). The majority of patients have locally advanced or metastatic cancer, and chemotherapy is mainly performed for unresectable PDAC. Gemcitabine monotherapy, a more intensive first-line chemotherapy regimen, has been approved for first-line treatment of patients with metastatic PDAC (4). Gemcitabine is a cell cycle-specific inhibitor of DNA synthesis and ribonucleotide reductase that has become the gold standard chemotherapeutic agent for pancreatic cancer (5, 6). However, the response rate to single administration of gemcitabine is not high, and thus, improvement of the therapeutic effect is required. Furthermore, no reliable molecular targets have been identified that can predict or influence the success of gemcitabine in PDAC.

MicroRNAs (miRNAs) are endogenous, single-stranded noncoding RNAs (~22 nucleotides in length) that control gene expression at the post-transcriptional level (7). miRNAs have been shown to function in cell proliferation, differentiation, cell cycle control, and chemotherapy resistance in cancer (8, 9). Aberrant expression of miRNAs has been observed in tissue and serum samples from PDAC patients as well as PDAC cell lines, indicating that miRNAs may participate in the pathogenesis of PDAC (10-12). We have previously reported on miRNA signatures for chemosensitivity and chemoresistance in hepatocellular and cholangiocellular carcinomas (13-14). We hypothesized, that currently unidentified miRNAs may influence gemcitabine treatment of PDAC and present targets for novel diagnostic and therapeutic options. Better understanding of the effects of gemcitabine on PDAC cells and potentially related miRNAs may provide insights for improving treatment efficacy.

In this experimental study, we analysed the effect of gemcitabine on PDAC cells and examined miRNA signatures of gemcitabine-treated PDAC in *in vitro* and *in vivo* models.

## Materials and Methods

**Drugs.** Gemcitabine was purchased from Eli Lilly Japan (Hyogo, Japan). Gemcitabine was prepared as a 10 mM stock solution in dimethyl sulfoxide (DMSO) and stored at  $-20^{\circ}\text{C}$ .

**Cell lines and culture.** The human pancreatic cancer cell lines PK-1 and PK-9 were provided by the Japanese Cancer Research Resources Bank (JCRB, Tokyo, Japan). The cell lines were maintained at  $37^{\circ}\text{C}$  with 5%  $\text{CO}_2$  in RPMI-1640 (Gibco, Grand Island, NY, USA) supplemented with 10% foetal bovine serum (Wako Pure Chemical Industries, Osaka, Japan), 20 U/ml penicillin, and 100  $\mu\text{g}/\text{ml}$  streptomycin (Invitrogen, Carlsbad, CA, USA).

**Cell proliferation assay.** The cell proliferation assay was performed as described previously (13). PK-1 and PK-9 cells were seeded into 96-well plates ( $5.0 \times 10^3$  cells) and treated as indicated for 48 h. Cell proliferation was assayed using the CCK-8 cell counting kit (Dojindo Laboratories, Kumamoto, Japan) according to the manufacturer's instructions. The absorbance was measured at 450 nm using a microplate reader (Multiskan FC, Thermo Scientific Corp., Waltham, MA, USA).

**Cell cycle analyses.** PK-1 cells were collected, seeded into 100-mm culture dishes at  $1.0 \times 10^6$  per dish, and cultured for 24 h. Cells were then treated with 30 nM gemcitabine for 24–48 h. Cell cycle was determined by flow cytometry using the Cytomics FC 500 flow cytometer (Beckman Coulter, Indianapolis, IN, USA), according to the manufacturer's protocol. Data were analysed using Kaluza software (Beckman Coulter). All experiments were performed in triplicate.

**Western blotting.** Gel electrophoresis and western blotting were performed in accordance with our previously described methods (13). Briefly, proteins were subjected to sodium dodecyl sulphate polyacrylamide gel electrophoresis and then transferred to nitrocellulose membranes. After blocking in 5% dry skim milk in 0.05% Tween-20/TBS buffer [25 mM Tris, 0.15 M NaCl, 0.05% (v/v) Tween-20, pH 7.5], the membranes were incubated with primary antibodies, followed by incubation with peroxidase-conjugated secondary antibodies in 5% dry skim milk in 0.05% Tween-20/TBS buffer. The proteins were visualized on an X-ray film using an enhanced chemiluminescence detection system (PerkinElmer Co., Waltham, MA, USA). Band intensities were quantified using ImageJ software 1.52q (NIH, Bethesda, MD, USA) and normalized to  $\beta$ -actin.

The following primary antibodies were used in this study:  $\beta$ -actin (1:10,000; Sigma-Aldrich, St. Louis, MO, USA), cyclin D1 (1:1,000; Thermo Fisher Scientific, Waltham, MA, USA), Cdk4 (1:1,000; Santa Cruz Biotechnology, Santa Cruz, CA, USA), Cdk6 (1:1,000, Santa Cruz Biotechnology), and Cdk2 (1:2,000; Santa Cruz Biotechnology). The secondary antibodies included horseradish peroxidase (HRP)-linked anti-mouse and anti-rabbit IgG (1:2,000; Cell Signaling Technology, Boston, MA, USA).

**MiRNA array and data analysis.** MiRNA array analysis was performed as described in our previous studies (13–14). Briefly, total RNA was isolated from PK-1 cells and tumour tissues using a miRNeasy Mini Kit (Qiagen, Hilden, Germany) according to the manufacturer's instructions. miRNA expression analysis was performed using the miRCURYHy3/Hy5 Power Labeling Kit and

human miRNA Oligo chip (v. 21.0; Toray Industries, Tokyo, Japan). The arrays were scanned in a 3D-Gene Scanner 3000 (Toray Industries), and fluorescence images were analysed using the 3D-Gene extraction version 1.2 software (Toray Industries).

**Xenografted tumour model.** All animals were treated in accordance with the guidelines of the Kagawa University Committee on Experimental Animals. The Kagawa University Animal Care Committee approved our animal protocols including animal ethics. Six-week-old BALB/c-nu/nu mice ( $n=18$ ) were obtained from Japan SLC (Shizuoka, Japan) and housed and maintained in specific pathogen-free conditions at  $20\text{--}26^{\circ}\text{C}$ . A standard sterilized laboratory diet and water were available *ad libitum*. A total of  $1 \times 10^6$  PK-1 cells were injected subcutaneously into the left flank of nude mice. After approximately 2 weeks, when tumours reached a maximal diameter of  $>3$  mm, 18 mice were randomly assigned into three groups. Mice were administered PBS only (control) ( $n=6$ ), 40 mg/kg gemcitabine ( $n=6$ ), or 80 mg/kg gemcitabine ( $n=6$ ) five times per week intraperitoneally (*i.p.*). The body weight and tumour volume were monitored every 3 days. Tumour volumes were calculated using the formula  $V = \text{length} \times \text{width}^2/2$  (15). The animals were sacrificed on day 42 after treatments.

**Immunohistochemistry.** We prepared 5- $\mu\text{m}$ -thick sections from formalin-fixed, paraffin-embedded tissue blocks. To retrieve antigens, the sections were boiled in 10 mmol/l citrate buffer (pH 6.0) using a microwave oven (MR-M201 Microwave Processor; Hitachi, Tokyo, Japan) at 500 W for 5 min. The sections were then dewaxed in xylene, rehydrated in graded alcohol solutions, and then mixed with a solution containing 0.5% hydrogen peroxidase to block endogenous peroxidase activity. After washing with phosphate-buffered saline, the sections were immunostained. Expression of cyclin D1 was detected by incubation with a mouse mAb (1:200 dilution) and rabbit pAb (1:50 dilution) overnight. The sections were then incubated for 30 min with appropriate peroxidase-conjugated secondary antibodies (PK-6102, PK-6101, Vectastain Elite ABC kit, Vector Laboratories, CA, USA).

The standard avidin-biotin-peroxidase complex method (Funakoshi Chemical Co., Tokyo, Japan) with diaminobenzidine (Sigma Chemical Co., Tokyo, Japan) as the substrate was employed for detection. Sections were counterstained by Meyer's hematoxylin. Samples treated with phosphate buffer served as the negative controls. Sections were examined microscopically, and nuclei with a brown colour (regardless of staining intensity) were regarded as positive. Cyclin D1-positivity was calculated for both at  $40\times$  magnification by dividing the number of positive cells by the total number of cells counted in five random fields and expressed as a percentage. Images were obtained with a digital image capture system (Olympus, Tokyo, Japan).

**Statistical analysis.** Data are expressed as the mean  $\pm$  standard deviation (SD). All statistical analyses were performed using GraphPad Prism 6 software (GraphPad Software, La Jolla, CA, USA). Nonparametric Wilcoxon/Mann-Whitney *U*-test was used to examine statistical significance between the two groups. A *p*-value less than 0.05 was considered significant.

## Results

**Gemcitabine treatment suppresses human PDAC cell growth in vitro.** Two human PDAC cell lines, PK-1 and PK-9, were

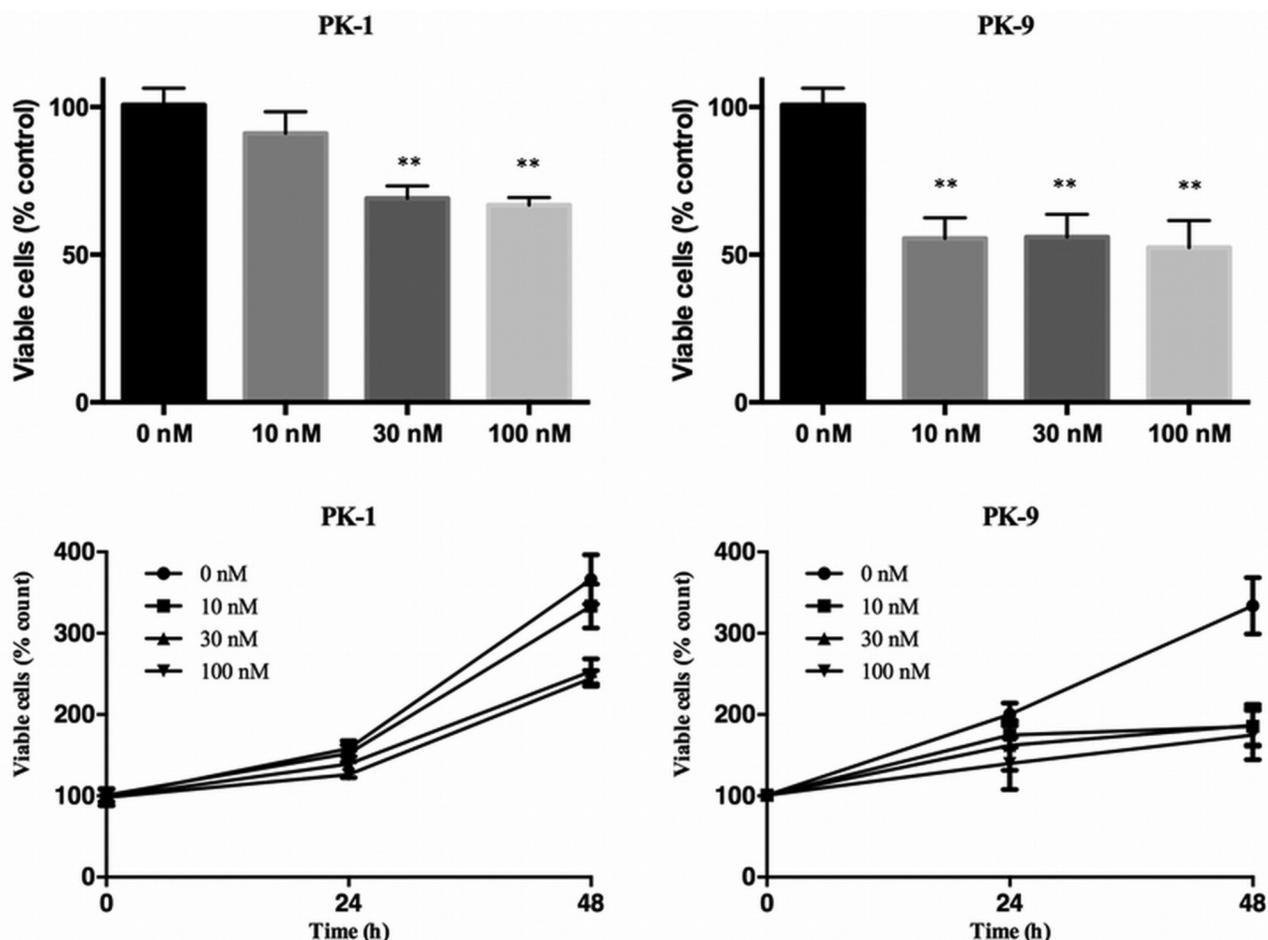


Figure 1. Gemcitabine inhibits the proliferation of PDAC cells. PDAC cells were incubated with 0, 10, 30, or 100 nmol/l gemcitabine for 48 h. Cell proliferation was evaluated by the CCK-8 assay. The results are expressed as the percentage of viable cells compared with the control (0 nmol/l). All treatments were significantly different from the control, based on Student's t-test (\* $p < 0.05$  and \*\* $p < 0.01$  vs. control).

treated with various concentrations of gemcitabine (0, 10, 30, or 100 nM) for 48 h. Treatment of PK-1 and PK-9 cells with gemcitabine for 48 h reduced their proliferation in a dose-dependent manner (Figure 1).

*Gemcitabine induces S phase cell cycle arrest and regulates cell cycle-related proteins in PK-1 cells.* We further investigated the effects of gemcitabine on the cell cycle of PK-1 cells by flow cytometric analysis. After treatment with 30 nM gemcitabine for 24 or 48 h, the percentage of PK-1 cells in the G<sub>0</sub>/G<sub>1</sub> phase was increased compared with control cells (56.48% vs. 34.14% after 24 h, 61.23% vs. 38.76% after 48 h) (Figure 2A). As the proportion of cells in the G<sub>0</sub>/G<sub>1</sub> phase increased, the proportion of cells in the S phase was decreased (29.57% vs. 36.67% after 24 h, 38.76% vs. 41.20% after 48 h). This accumulation was associated with a reduction of G<sub>2</sub>/M phase cells. We further found that the expression levels of cyclin D1 as well as CDK4 and

CDK6, the catalytic partners of cyclin D1, were decreased in PK-1 cells after gemcitabine treatment (Figure 2B and C). Together these results indicated that the growth of PDAC cells was suppressed after gemcitabine treatment by impairing cell cycle progression.

*Gemcitabine treatment suppresses growth of tumours derived from human PDAC cells.* We next investigated the anti-tumour effect of gemcitabine *in vivo*. Xenografted mice were intraperitoneally administrated gemcitabine (40 or 80 mg/kg) or PBS after subcutaneous implantation of PK-1 cells. Gemcitabine effectively suppressed the tumour growth from PK-1 xenografts (Figure 3A). The tumour size was significantly smaller in gemcitabine-treated mice compared with the controls (Figure 3B). We also analysed tumour sections by immunohistochemistry to determine whether gemcitabine also affected cyclin D1 *in vivo*. Gemcitabine treatment resulted in decreased expression levels of cyclin

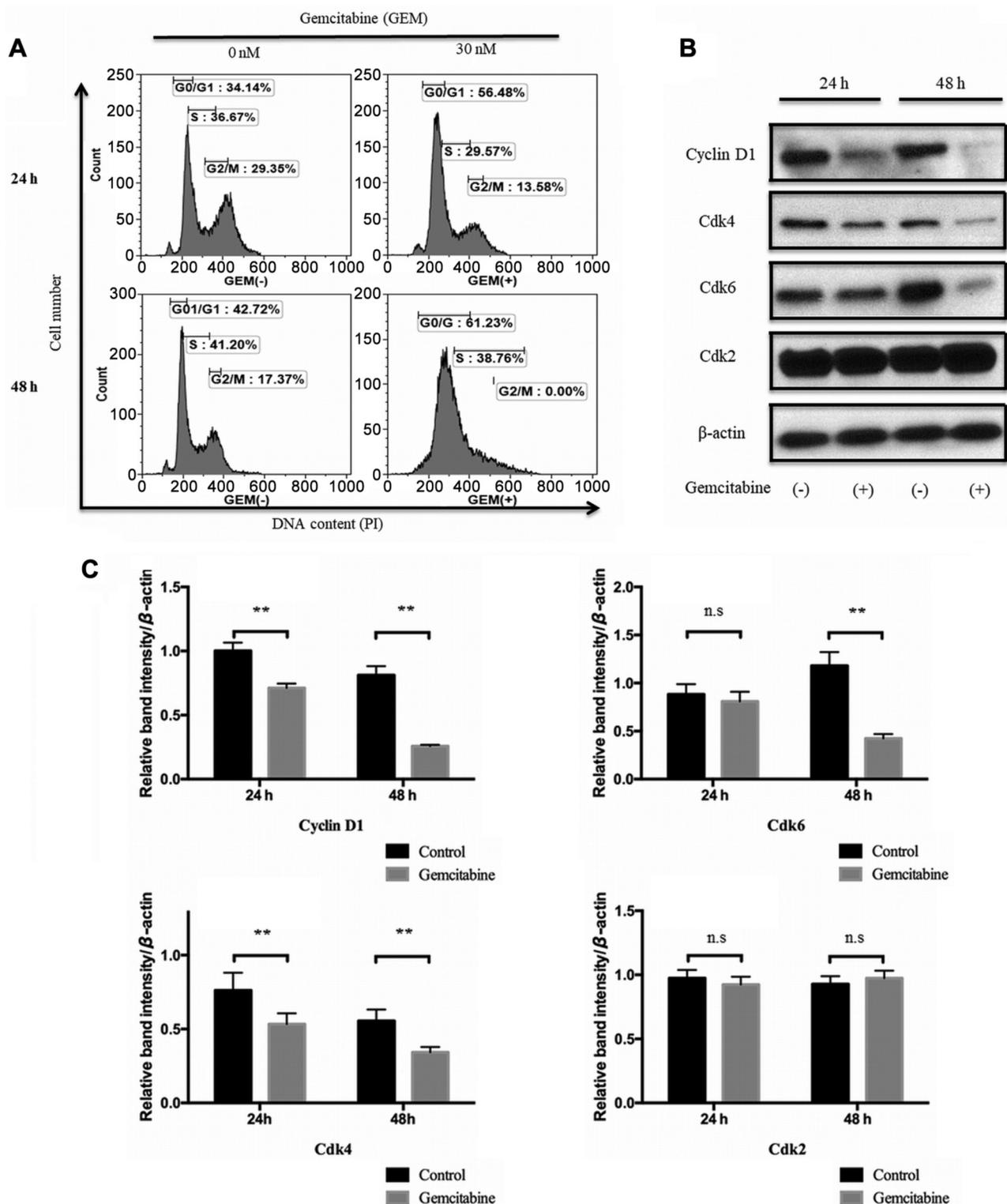


Figure 2. Gemcitabine induces S phase arrest in PK-1 cells (A) Representative results showing the distribution of PK-1 cells in G0/G1, S, and G2/M phases following treatment with 30 nM gemcitabine for 24 and 48 h. (B) Western blot showing expression of cyclin D1, CDK4, and CDK6 in PK-1 cells following treatment with 30 nM gemcitabine for 24 and 48 h. (C) Cyclin D1, CDK4, and CDK6 expression were decreased by 30 nM gemcitabine treatment compared with the control. The images are representative of three independent experiments, and protein levels were normalized to  $\beta$ -actin. Values represent mean $\pm$ SD; \* $p$ <0.05 and \*\* $p$ <0.01 vs. control. n.s.: Not significant.

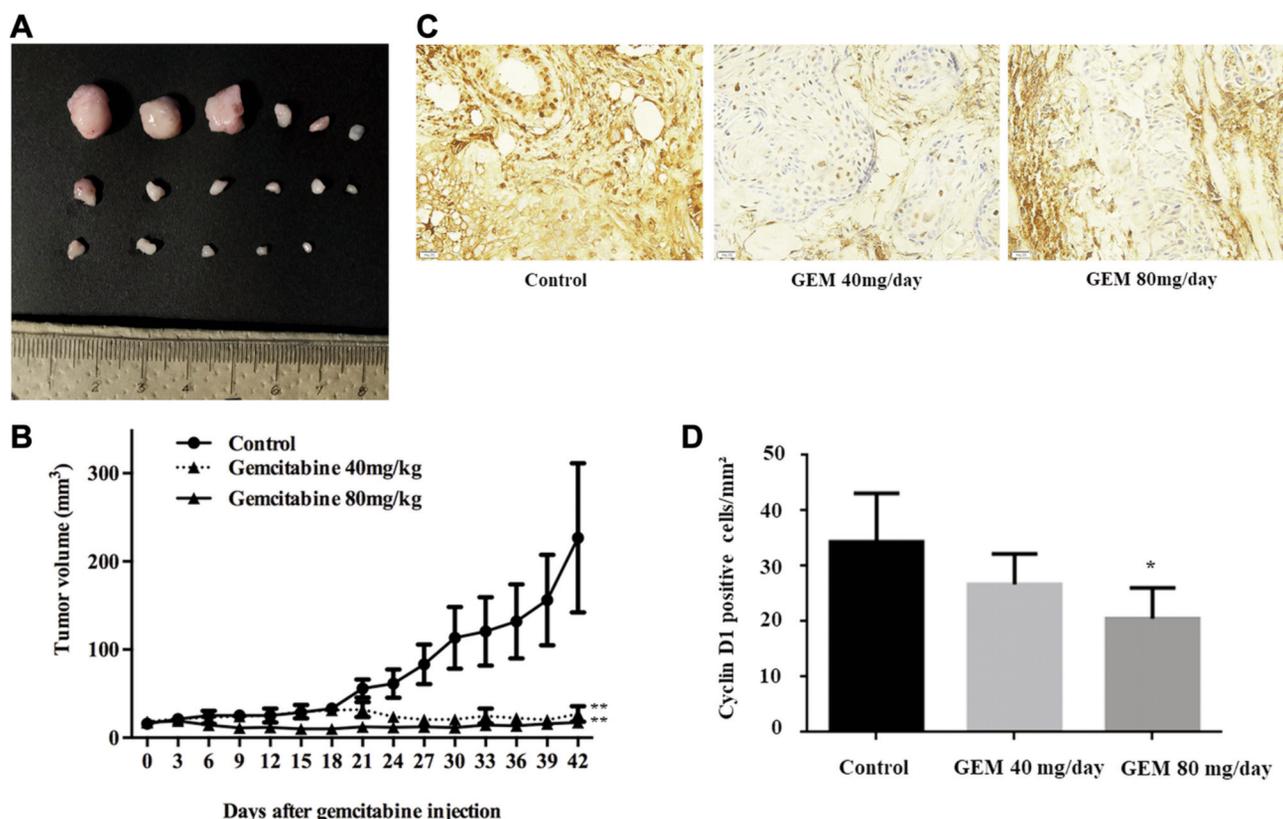


Figure 3. Gemcitabine inhibits the growth of PK-1 cell xenografts in nude mice. PK-1 cells were subcutaneously implanted into the flanks of nude mice. When the tumours became palpable, 0, 40 or 80 mg/kg gemcitabine were intraperitoneally injected five times per week for 42 days. (A) Tumour growth curves of control and gemcitabine treatment groups. (B) Tumours were significantly smaller in gemcitabine-treated mice compared with vehicle-treated mice. Each point represents the mean±standard deviation of eight animals (\* $p < 0.05$  and \*\* $p < 0.01$  vs. control). (C) Immunohistochemical staining of cyclin D1 in cancerous tissues from gemcitabine-treated and control groups of xenografted mice. Cyclin D1-positive cells (black arrows) were decreased in mice treated with gemcitabine. (D) Cyclin D1-positive cells in gemcitabine-treated mice were reduced compared with untreated mice.

D1 compared to controls (Figure 3C). The labelling index of gemcitabine-treated cell was also lower compared to controls (Figure 3D). Gemcitabine treatment had no significant effect on the body weight of animals during the treatments.

*Gemcitabine treatment alters the miRNA profile in gemcitabine-treated PK-1 cells and tumour tissues.* Heat maps generated by miRNA microarray analysis revealed dysregulated expression of several miRNAs in gemcitabine-treated PK-1 cells and tumour tissues (Figure 4A). We examined the expression levels of 2555 miRNAs in PK-1 cells with or without 30 nM gemcitabine treatment. After normalization and removing miRNAs with missing values, 241 miRNAs were used to perform hierarchical clustering. Finally, 14 significantly altered miRNAs were identified. Of these, 11 were significantly up-regulated miRNAs and 3 were significantly down-regulated miRNAs. The miRNAs that were up-regulated included miR-338-3p, miR-132-3p,

miR-135b-5p, miR-455-3p, miR-345-5p, miR-4506, miR-3653-3p, miR-191-5p, miR-424-5p, miR-34a-3p, and miR-34a-5p (Table I). Those that showed significantly increased expression levels were miR-5100, miR-19b-1-5p, and miR-205-3p (Table I).

In addition, 29 significantly altered miRNAs were identified in xenograft tumour tissues with or without gemcitabine treatment (Figure 4B), including 7 significantly up-regulated and 22 significantly down-regulated miRNAs (Table II).

## Discussion

In the present study, we examined the effects of gemcitabine in PDAC. Gemcitabine induced cell cycle arrest at the S phase through the modulation of the cell cycle-regulating protein cyclin D1 in PK-1 cells. We also identified the miRNA signature of gemcitabine in PDAC cells and a

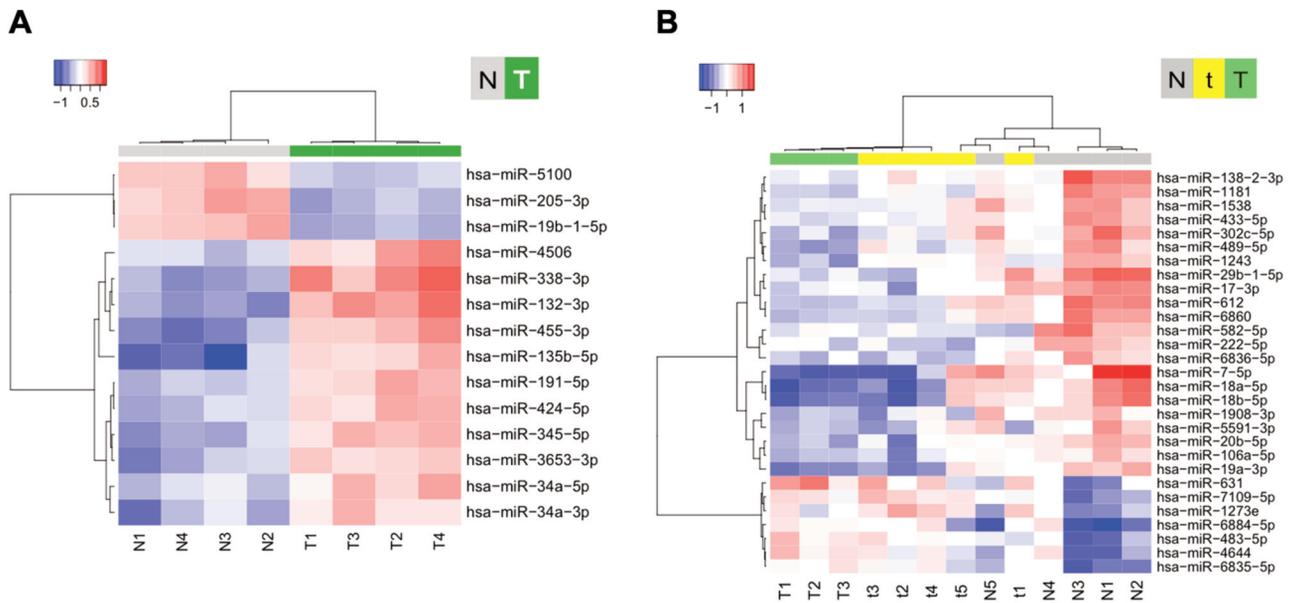


Figure 4. Hierarchical clustering of miRNAs in (A) PK-1 cells and (B) tumour tissues treated with or without gemcitabine. The analysed samples are shown in the columns, and the miRNAs are presented in the rows. The miRNA clustering colour scale presented at the top indicates the relative miRNA expression levels; red and blue represent high and low expression levels, respectively. Gray scale indicates no treatment. Yellow scale indicates 40 mg/kg gemcitabine, and light green scale indicates 80 mg/kg gemcitabine.

xenograft model. To the best of our knowledge, this is the first study to show that the miRNA signature of the anti-tumour effect of gemcitabine associated with cell proliferation and tumour growth *in vitro* and *in vivo*.

Gemcitabine is a deoxycytidine analogue and its cytotoxic activity is based on several activities in DNA synthesis (16). As a cytidine analogue, gemcitabine is incorporated into DNA during replication in the S phase of cell cycle. Gemcitabine is used as chemotherapy for pancreas, breast, ovarian, and non-small-cell lung cancer (16), and it is the standard of care for patients with these tumours (17). Gemcitabine exerts its activity primarily by inducing cell cycle arrest and cell death (18, 19). Our results showed that gemcitabine induced a time- and concentration-dependent cell cycle arrest at the S phase, which is consistent with previous results (20, 21). The gemcitabine-induced S phase arrest was detected as both an increase in the transition rate from G<sub>0</sub>/G<sub>1</sub> and an inhibition of the transition rate from S phase to G<sub>2</sub>/M phase. The S phase arrest was increased with gemcitabine concentration (22). In the present study, gemcitabine treatment in PK-1 and PK-9 cells led to a strong, dose-dependent inhibition of cell proliferation. However, gemcitabine did not exert an anti-proliferative effect on PANC-1 cells, which are considered relatively resistant to many chemotherapeutic regimens (23). Additionally, the expression levels of cyclin D1 was decreased in PK-1 cells after gemcitabine treatment. Cyclin D1 is an important interacting partner for CDK4 and CDK6 and regulates the

Table I. Statistical results and chromosomal locations of miRNAs in PK-1 cells treated with or without gemcitabine. Fold changes are gemcitabine-treated/non-treated. Fold change (FC)>1.5, FC<0.4, p<0.005.

miRNA	Fold change (T/N)	Chromosomal localization	p-Value
Up-regulated			
hsa-miR-338-3p	2.237	17q25.3	0.000099
hsa-miR-132-3p	2.206	17p13.3	0.000046
hsa-miR-135b-5p	2.099	1q32.1	0.002882
hsa-miR-455-3p	2.016	9q32	0.000275
hsa-miR-345-5p	1.711	14q32.2	0.000433
hsa-miR-4506	1.639	14q32.12	0.003404
hsa-miR-3653-3p	1.617	22q12.2	0.000703
hsa-miR-191-5p	1.614	3p21.31	0.00018
hsa-miR-424-5p	1.593	Xq26.3	0.000601
hsa-miR-34a-3p	1.58	1p36.22	0.00453
hsa-miR-34a-5p	1.525	1p36.22	0.002963
Down-regulated			
hsa-miR-5100	0.631	10q11.21	0.000035
hsa-miR-19b-1-5p	0.571	13q31.3	0.000025
hsa-miR-205-3p	0.559	1q32.2	0.000064

cell cycle transition from G<sub>1</sub> to S phase (24). Increased cyclin D1 expression is associated poor prognosis (25) and decreased postoperative survival (26) of PDAC patients. Therefore, suppression of cyclin D1 with gemcitabine treatment was associated with the growth-inhibitory effect of

Table II. Statistical results and chromosomal locations of miRNAs in PK-1 cell-derived tumour tissues treated with or without gemcitabine. Fold changes are gemcitabine-treated/non-treated (T/N). Fold change (FC)>1.5, FC<0.67, p<0.01.

miRNA	Fold change (T/N)	Chromosomal localization	p-Value
<b>Up-regulated</b>			
hsa-miR-631	1.88	15q24.2	0.0061
hsa-miR-6884-5p	1.77	17q21.1	0.007
hsa-miR-1273e	1.75	17q23.3	0.0014
hsa-miR-6835-5p	1.61	6p21.31	0.0025
hsa-miR-4644	1.6	6q27	0.0087
hsa-miR-7109-5p	1.6	22q12.2	0.0029
hsa-miR-483-5p	1.57	11p15.5	0.0089
<b>Down-regulated</b>			
hsa-miR-7-5p	0.35	9q21.32	0.0059
hsa-miR-18a-5p	0.45	13q31.3	0.0058
hsa-miR-18b-5p	0.46	Xq26.2	0.0043
hsa-miR-612	0.53	11q13.1	0.0019
hsa-miR-29b-1-5p	0.54	7q32.3	0.0062
hsa-miR-19a-3p	0.55	13q31.3	0.0026
hsa-miR-302c-5p	0.55	4q25	0.0028
hsa-miR-6860	0.55	11q13.2	0.0017
hsa-miR-582-5p	0.57	5q12.1	0.009
hsa-miR-138-2-3p	0.58	16q13	0.0095
hsa-miR-1181	0.59	19p13.2	0.0022
hsa-miR-1908-3p	0.59	11q12.2	0.0011
hsa-miR-17-3p	0.6	13q31.3	0.0068
hsa-miR-489-5p	0.6	7q21.3	0.0065
hsa-miR-5591-3p	0.6	4p14	0.0036
hsa-miR-222-5p	0.62	Xp11.3	0.0011
hsa-miR-1538	0.63	16q22.1	0.0018
hsa-miR-20b-5p	0.63	Xq26.2	0.0067
hsa-miR-106a-5p	0.66	Xq26.2	0.0036
hsa-miR-1243	0.66	4q25	0.0099
hsa-miR-433-5p	0.66	14q32.2	0.0038
hsa-miR-6836-5p	0.66	7p22.3	0.0073

other chemotherapeutic agents and decreased expression of multiple chemoresistance genes in human PDAC cells (27).

In addition to its effects upon the cell cycle, gemcitabine induces caspase-3-mediated apoptosis (28). Hamed *et al* have revealed that the apoptotic effect of gemcitabine was more evident at later time points of treatment (72 and 96 h) (29). These results suggest that gemcitabine induces cell cycle arrest in the early phase (24-48 h after gemcitabine treatment) and that the apoptotic effects of gemcitabine take place at later time points (72 and 96 h).

Cell cycle regulation is crucial for cell growth and many miRNAs are involved in the negative control of cell cycle genes, which are essential during oncogenesis of PDAC (30). miRNAs are endogenous mediators of gene expression through their site-specific binding at the 3' untranslated region or other sites of mRNAs to either degrade their target or inactivate protein synthesis (31). miRNAs regulate many biological

processes such as cancer cell proliferation, tumour growth, differentiation, apoptosis, and energy metabolism (32). We identified miRNAs associated with the anti-tumour effects of gemcitabine in PK-1 cells using miRNA expression arrays. Several miRNAs were significantly altered following gemcitabine treatment of PK-1 cells. Among these, miR-338-3p has been shown to be up-regulated in pancreatic intraepithelial neoplasms, which is a precursor lesion of PDAC (33). However, the role of miR-338-3p in advanced PDAC remains unknown. Overexpression of miR-338-3p has been shown to promote cell proliferation, migration, and invasion in gastric cancer by targeting metastasis-associated in colon cancer-1 (MACC1) (34). Our results showed that miR-338-3p expression levels were significantly increased in gemcitabine-treated PK-1 cells. Overexpression of miR-582-5p has been shown to induce colorectal cancer cell proliferation by increasing cyclin D1 expression (35). Gemcitabine significantly down-regulate miR-582-5p in PK-1 xenograft tissues, and down-regulation of miR-582-5p may contribute to suppressed cancer proliferation via modulating cell cycle-related proteins (35). Although several miRNAs were significantly altered in PK-1 xenograft tumour tissues from mice following 40 mg and 80mg gemcitabine treatment, we did not identify any overlapping significantly altered miRNAs from cultured cells and tumour tissues following gemcitabine treatment.

In conclusion, our results revealed that gemcitabine inhibits human PDAC cell proliferation by inducing cell cycle arrest. Additionally, we identified numerous miRNAs associated with the anti-tumour effects of gemcitabine in PDAC cells and tumour tissues using miRNA expression arrays. This aberrant miRNA expression signature may provide insights into PDAC development and therapeutic opportunities to improve gemcitabine monotherapy.

### Conflicts of Interest

The Authors disclose no potential conflicts of interest related to this study.

### Authors' Contributions

DN and TM designed experiments. SF, HI, KF, TM, MN, MO, MH, TK, NF, HY, KK, HKa, AM, HKo, and KT conducted the experiments, data analysis, and final drafting and writing of the manuscript. SF contributed to drafting of the manuscript. TM was involved in the research design and drafting of the final manuscript. All Authors have read and approved the final version of the manuscript.

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