

学 位 論 文

**MicroRNA profiles during  
galectin-9-induced apoptosis of  
pancreatic cancer cells**

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# MicroRNA profiles during galectin-9-induced apoptosis of pancreatic cancer cells

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**Abstract.** Pancreatic cancer is the eighth-leading cause of cancer-associated mortality in males and the ninth-leading cause in females worldwide. Even when diagnosed early enough to be potentially resectable, the prognosis of invasive pancreatic cancer is poor. Galectin-9 (Gal-9) is a tandem-repeat type galectin that has recently been demonstrated to possess an anti-proliferative effect on cancer cells. Therefore, the present study evaluated the effects of Gal-9 on the proliferation of human pancreatic cancer cells and examined the microRNAs that are associated with the antitumor effects of Gal-9. Gal-9 suppressed the proliferation of multiple pancreatic cancer cell lines. In addition, Gal-9 treatment increased the levels of caspase-cleaved keratin 18 and the expression of cytochrome c in pancreatic cancer cell lines. This data suggests that Gal-9 induces intrinsic apoptosis in pancreatic cancer cell lines through the caspase-dependent and caspase-independent pathways. In addition, Gal-9 reduced the expression levels of phosphorylated epidermal growth factor receptor and numerous

receptor tyrosine kinases (RTKs). In conclusion, Gal-9 may suppress the growth of human pancreatic cancer cells *in vitro*. These findings suggest that Gal-9 may be a new therapeutic agent for the treatment of pancreatic cancer.

## Introduction

Pancreatic cancer is the eighth-leading cause of cancer associated mortality in males and the ninth-leading cause in females worldwide (1). Pancreatic cancer has a rapid disease progression that is accompanied by the absence of specific symptoms, which largely precludes early diagnosis and curative treatment (2,3). The vast majority of patients with pancreatic cancer are not diagnosed until their cancer has metastasized, and so surgical treatment is the only curative treatment. However, due to this late diagnosis the majority of patients present in an advanced stage, and only a minority (10-20%) of these patients are amenable to surgical intervention (4,5). Even when diagnosed early enough to be potentially resectable, the prognosis of invasive pancreatic cancer is poor. Due to its high recurrence rate, post-operative patients with pancreatic cancer require adjuvant chemotherapy with or without radiotherapy, which provides a 5-year patient survival rate of 15-25% (6).

Galectin-9 (Gal-9) is a tandem-repeat type galectin with 2 carbohydrate-recognition domains (CRDs), and it was first identified as an eosinophil chemoattractant and activation factor (7-10). Similarly to other galectins, Gal-9 performs a role in cell aggregation and adhesion and in the apoptosis of tumor cells (10,11). Gal-9 can enhance antitumor immunity through the initial CRD-independent maturation of dendritic cells and the subsequent induction of Th1-mediated antitumor immunity (12). In addition, treatment with recombinant Gal-9 prolonged survival in a murine melanoma model, not only by increasing the numbers of CD8 cytotoxic T cells (CTLs) but also by increasing the number of natural killer (NK) cells and macrophages (13).

Previous studies have uncovered additional mechanisms by which T cell immunoglobulin mucin-3, a receptor for Gal-9, negatively regulates T cell responses by promoting CD8+ T cell exhaustion and inducing the expansion of myeloid-derived suppressor cells (14,15). Therefore, Gal-9 augments antitumor

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**Abbreviations:** Gal-9, Galectin-9; CRDs, carbohydrate-recognition domains; CTLs, cytotoxic T cells; NK cells, natural killer cells; miRNAs, microRNAs; CCK-8, cell counting kit; CCK18, Caspase-cleaved keratin 18; phospho-RTK, phosphorylated receptor tyrosine kinases; EGFR, epidermal growth factor receptor; CRDs, carbohydrate-recognition domains; Tie-2, tyrosine kinase with immunoglobulin-like and EGF-like domains 2; TrKC, tropomyosin receptor kinase C; MuSK, muscle-Specific Kinase; ALK, anaplastic lymphoma kinase; Eph, erythropoietin-producing human hepatocellular carcinoma cell

**Key words:** pancreatic cancer, galectin-9, apoptosis, caspase-cleaved keratin 18, cell cycle, microRNAs, cytochrome c

immunity by inducing a subset of macrophages and dendritic cells and activating tumor-specific CTLs and NK cells.

Recombinant Gal-9 induced apoptosis in various T cell leukemic cell lines in a dose-dependent and functional CRD-dependent manner (16,17). Additionally, several *in vitro* and *in vivo* studies have also indicated that Gal-9 inhibits the growth of multiple myeloma (18) and chronic myeloid leukemia (19). In hematologic malignancies, Gal-9 suppresses cellular proliferation and tumor growth *in vitro* and *in vivo*. However, in solid malignancies, the antitumor effect of Gal-9 remains unknown. Previously, it was reported that Gal-9 inhibits the growth of hepatocellular carcinoma and cholangiocarcinoma via apoptosis *in vitro* and *in vivo* (20,21).

However, less is known regarding the antitumor effects of Gal-9 on pancreatic cancer cells or the microRNAs (miRNAs) that are associated with these effects. The present study therefore evaluated the effects of Gal-9 on the growth of two pancreatic cancer cell lines, its mechanism of action, and the miRNAs that are associated with the antitumor effect of Gal-9 on pancreatic cancer cells.

## Materials and methods

**Cell lines and culture.** The human pancreatic cancer PK-1 and PK-9 cell lines were obtained from the RIKEN cell bank (Ibaraki, Japan) and passaged in our laboratory for <6 months. The 2 cell lines were authenticated by the cell bank using short tandem repeat polymerase chain reaction. PK-1 and PK-9 cells were grown in RPMI 1640 supplemented with 10% fetal bovine serum and 100 mg/l of penicillin-streptomycin in a humidified atmosphere with 5% CO<sub>2</sub> at 37°C.

**Chemicals and reagents.** Recombinant mutant forms of human Gal-9 that lack linker peptides were expressed and purified as described in one of our previous studies (22). A cell counting kit (CCK-8) was purchased from Dojindo Molecular Technologies, Inc. (Kumamoto, Japan), and all other chemicals were obtained from Sigma-Aldrich (Merck KGaA, Darmstadt, Germany).

**Cell proliferation assay.** A total of 5 × 10<sup>3</sup> cells were seeded in 100 μl of culture medium supplemented with 10% FBS into each well of a 96-well microplate and incubated at 37°C overnight. Following 24 h, cells were treated with 0.1, 0.3 or 1.0 μM of Gal-9 and the cells were cultured at 37°C for an additional 48 h. A total of 10 μl CCK-8 reagent was added to each well and the plates were incubated at 37°C for 3 h. The absorbance of each well was measured at 450 nm using an auto-microplate reader.

**ELISA assay for apoptosis.** Caspase-cleaved keratin 18 (CCK18) was evaluated using M30 Apoptosense ELISA kits obtained from PEVIVA AB (Bromma, Sweden) (23). Each cell line (cell density, 5 × 10<sup>3</sup> cells) was seeded into 96-well plates and cultured in 100 μl of culture medium for 24 h. Cells were then treated with 0.3 μM of Gal-9. The procedures of the assays were performed according to the manufacturer's protocol. The amounts of antigen in the control and treated samples were calculated by interpolation of a standard curve.

**Gel electrophoresis and western blotting.** PK-1 cells (1.0 × 10<sup>6</sup>/dish) were seeded in 100-mm culture dishes and

cultured at 37°C for 24 h. Gal-9 was subsequently added, and the cells were cultured at 37°C for an additional 48 h. Cells were washed twice in PBS and lysed using a protease inhibitor cocktail (Proprep, Complete protease inhibitor mixture; Intron Biotechnology, Inc., Seongnam, Korea). Protein concentration was quantified using a NanoDrop 2000 fluorospectrometer (Thermo Fisher Scientific, Inc., Waltham, MA, USA). Whole-cell lysates (10 μg) were separated using SDS-PAGE on 10% Tris-glycine gradient gels (24), and the proteins were transferred to nitrocellulose membranes. Subsequent to blocking with a solution containing 5% non-fat milk powder in TBS/Tween-20 at 37°C for 1 h, the membranes were incubated with primary antibodies at 4°C overnight and incubated with horseradish peroxidase (HRP) -conjugated secondary antibodies at 4°C for 1 h (25). The antibodies used were: Anti-β-actin monoclonal antibody (dilution, 1:3,000; cat. no. A5441; Sigma-Aldrich; Merck KGaA), cyclin D1 (dilution, 1:1,000; cat. no. RB-9041; Thermo Fisher Scientific Inc.), cyclin E (dilution, 1:1,000; cat. no. 870P1605B; Thermo Fisher Scientific, Inc.), Cdk (cyclin dependent kinase) 6 (dilution, 1:1,000; cat. no. sc-177; Santa Cruz Biotechnology, Inc., Dallas, TX, USA), Cdk4 (dilution, 1:1,000; cat. no. sc-749; Santa Cruz Biotechnology, Inc.), Cdk2 (dilution, 1:2,000; cat. no. sc-163; Santa Cruz Biotechnology, Inc.) and secondary horseradish peroxidase-linked anti-mouse (dilution, 1:2,000; cat. no. 7076; Cell Signaling Technology, Inc., Danvers, MA, USA) and anti-rabbit Immunoglobulin G antibodies (dilution, 1:2,000; cat. no. 7074; Cell Signaling Technology, Inc.).

Immunoreactive proteins were visualized using an enhanced chemiluminescence detection system (Perkin Elmer Inc., Waltham, MA, USA) on X-ray film.

**Antibody arrays of apoptosis-related proteins.** A Human Apoptosis Antibody Array kit (R&D Systems, Inc., Minneapolis, MN, USA) was used according to the manufacturer's protocol.

**Antibody arrays of phosphorylated receptor tyrosine kinase (p-RTK).** Human phospho-RTK was assayed using Human Phospho-RTK Array kits (R&D Systems, Inc.), according to the manufacturer's protocol. Each array membrane was exposed to X-ray film using a chemiluminescence detection system (Perkin-Elmer Co.).

**Analysis of miRNA arrays.** Total RNA was extracted from tumor samples using miRNeasy Mini kits (Qiagen GmbH, Hilden, Germany) according to the manufacturer's protocol. RNA samples typically demonstrated A<sub>260/280</sub> ratios between 1.9 and 2.1 when assayed on an Agilent 2100 Bioanalyzer (Agilent Technologies, Inc., Santa Clara, CA, USA). PK-1 cells were cultured at 37°C with or without 0.3 μM Gal-9 for 24 h, with 5 replicates per sample. Following RNA measurement with an RNA 6000 Nano kit (Agilent Technologies, Inc.), the samples were labeled using a miRCURY LNMä microRNA Array Hi-Power Labeling kit (Exiqon A/S, Vedbaek, Denmark) and were hybridized to a human miRNA Oligo chip (v.21.0; Toray Industries). The chips were scanned with a 3-D Gene Scanner 3000 (Toray Industries, Inc., Tokyo, Japan), and the results were analyzed with 3D-Gene extraction version 1.2 software (Toray Industries, Inc., Tokyo, Japan).

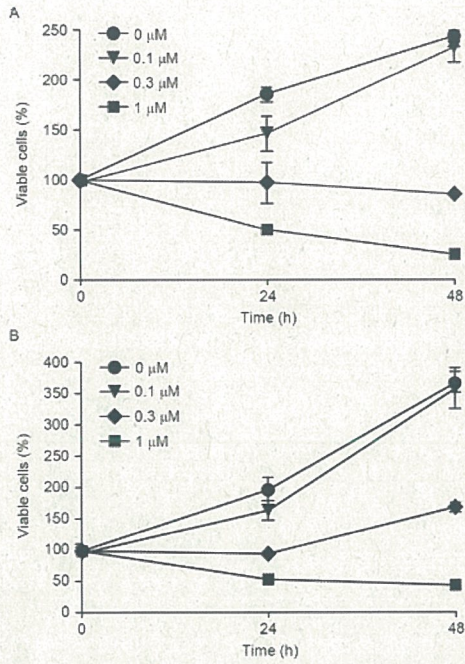


Figure 1. Gal-9 suppresses the proliferation of pancreatic cancer cell lines. (A) PK-1 and (B) PK-9 cells were seeded in 96-well plates. After 24 h, Gal-9 (0.1, 0.3 and 1 μM) was added to the culture medium. Subsequently, 24 h after the addition of these agents, a CCK-8 assay was conducted. PK-1 and PK-9 cells were seeded at 5,000 cells per well in a 96-well plate, and the agents were added at the 0-h time point. The data points represent mean cell numbers from 2 independent clusters, and the error bars represent the standard deviation. For PK-1 and PK-9 cells, the conditions at 24 and 48 h were significantly different from those of the control. Gal-9, galectin-9.

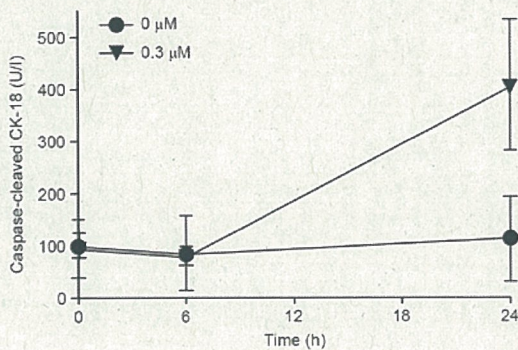


Figure 2. CCK18 produced specifically during apoptosis was identified by ELISA. Cells were incubated with or without 0.3 μM Gal-9. Gal-9 increased the levels of CCK18 in PK-1 cells, suggesting that the apoptotic process following phosphatidylserine exposure proceeds to cut the intermediate filaments of cells. Fold-change in CCK18 is shown as the mean ± standard deviation, with analysis by a paired t-test. For each cell line, the condition at 24 h was significantly different from that of the control. CCK18, caspase-cleaved cytokeratin-18; Gal-9, galectin-9.

Differences in miRNA expression between Gal-9-treated and control samples were assessed by analyzing the raw data using GeneSpringGX v. 10.0 (Agilent Technologies, Inc). On the raw data that were above the background level, quantile normalization was performed. Differentially expressed miRNAs were determined using the Mann-Whitney U test. False discovery rate was computed with Benjamini-Hochberg method (26) as the correction for multiple testing. Hierarchical clustering was performed using the farthest neighbor method with the

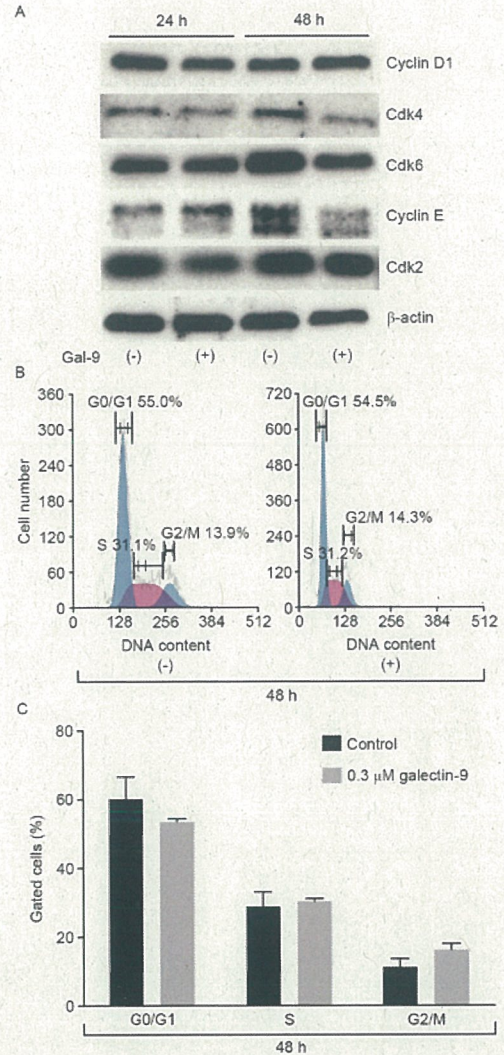


Figure 3. Western blot analysis and flow cytometric analysis of the cell cycle in PK-1 cells. (A) Expression levels of cell cycle-related proteins (cyclin D1, Cdk4, Cdk6, cyclin E and Cdk2) in PK-1 cells subsequent to the addition of 0.3 μM Gal-9 did not change compared with the β-actin control. (B) Flow cytometric analysis of proliferating PK-1 cells 48 h after the addition of 0.3 μM Gal-9. (C) PK-1 cells treated with Gal-9 demonstrated no change in the cell cycle. Cdk, cyclin dependent kinase; Gal-9, galectin-9.

absolute uncentered Pearson's correlation coefficient as a metric. A heat map was produced with the relative expression intensity for each miRNA, in which the base-2 logarithm of the intensity was median-centered for each row.

**Statistical analysis.** All statistical analyses were performed using JMP 9.0 software (SAS Institute, Cary, NC, USA). Paired analysis between groups was performed using t-tests. P<0.05 was considered to indicate a statistically significant difference.

**Results**

**Gal-9 suppresses the proliferation of human pancreatic cancer cells.** To evaluate the effect of Gal-9 on the growth activity of human pancreatic cells *in vitro*, the present study examined the effect of Gal-9 on the proliferation of the pancreatic cancer PK-1 and PK-9 cell lines. Cells were grown in 10% FBS and treated with 0.1, 0.3 or 1.0 μM Gal-9 or without Gal-9 (as a

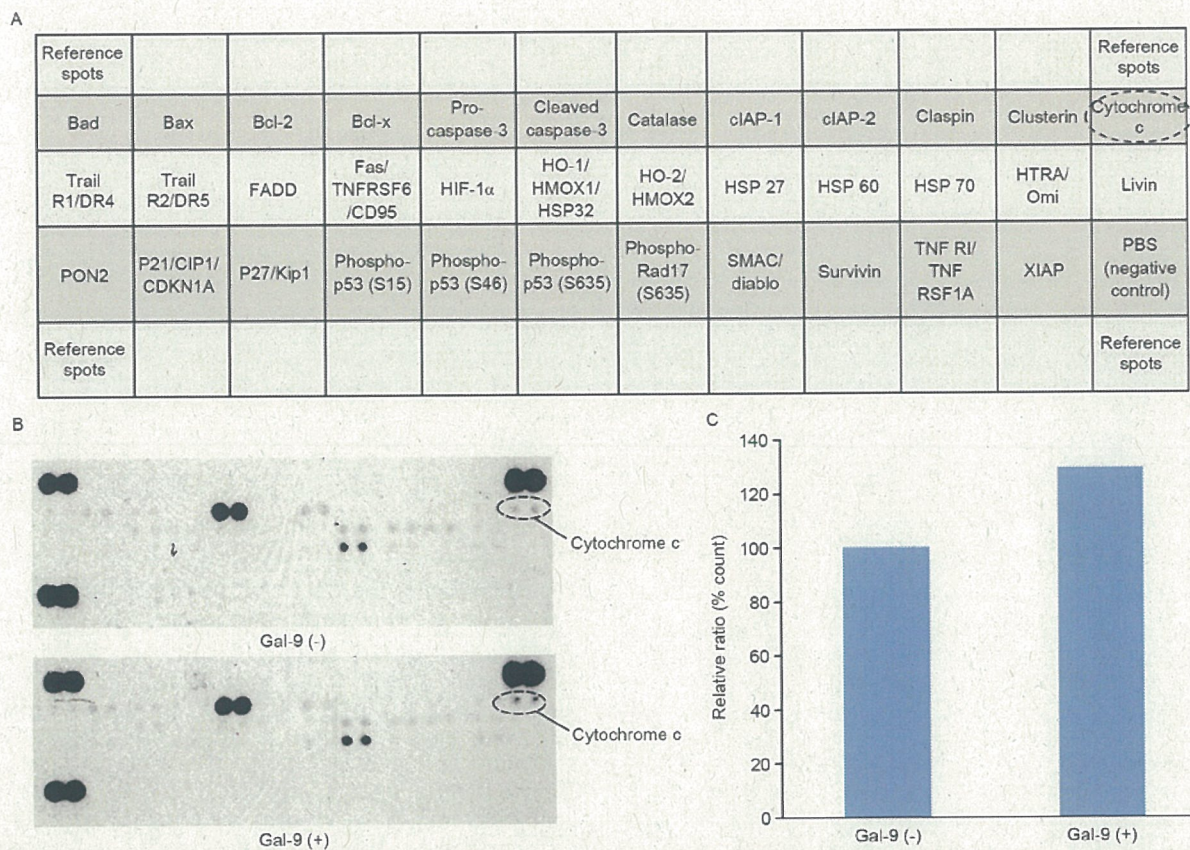


Figure 4. The effects of Gal-9 on apoptosis-related proteins in PK-1 cells. (A) Template showing the locations of tyrosine kinase antibodies spotted onto a human apoptosis array. (B) Representative expression of various apoptosis-related proteins in PK-1 cells treated with or without Gal-9. (C) Densitometry analysis demonstrated that the intensity of cytochrome c spots from Gal-9-treated cells relative to untreated cells was 129.7%. Gal-9, galectin-9; Bad, Bcl-2 associated death promotor; Bax, Bcl-2 associated x protein; Bcl-2, B-cell lymphoma 2; Bcl-x, B cell lymphoma-extra-large; clAP, cellular inhibitor of apoptosis protein; Trail R1/DR4, Trail receptor 1/death receptor 4; FADD, Fas associated via death domain; TNFRSF6/CD95, tumor necrosis factor receptor superfamily member 6/cluster of differentiation 95; HIF-1 $\alpha$ , hypoxia-inducible factor-1 $\alpha$ ; HO-1/HMOX1/HSP32, heme oxygenase-1/heme oxygenase (decycling) 1/heat shock protein 32; PON2, paraoxonase 2; P21/CIP1/CDKN1A, cyclin-dependent kinase inhibitor 1A; SMAC, second mitochondria-derived activator of caspases; TNF R1, tumor necrosis factor receptor 1; TNFRSF1A, TNF receptor superfamily member 1A; XIAP, X-linked inhibitor of apoptosis protein.

control) for 48 h. Gal-9 demonstrated strong, dose-dependent inhibition of cellular proliferation the two pancreatic cancer cell lines (Fig. 1).

*Gal-9 induces apoptosis to suppresses cellular proliferation of pancreatic cancer.* To clarify the mechanism of the inhibitory effect of Gal-9 on cancer cell growth, the present study first examined the induction of apoptosis by Gal-9. CCK18 was evaluated by ELISA to confirm whether apoptosis was involved in Gal-9-induced cell death. Gal-9 increased the levels of CCK18 in the two pancreatic cancer cell lines (Fig. 2).

*The effects of Gal-9 on the cell cycle in PK-1.* The effects of Gal-9 on the expression of various cell-cycle-related molecules in PK-1 cells were evaluated using western blot analysis. Cells were treated with 0 or 0.3  $\mu$ M Gal-9 for 24-48 h. Assays for the expression of proteins associated with the G<sub>0</sub>-to-G<sub>1</sub> transition demonstrated that cyclin E and Cdk4 were slightly decreased 48 h after the addition of 0.3  $\mu$ M of Gal-9 (Fig. 3A). The expression of other proteins, including cyclin D1, Cdk2 and Cdk6 were not changed compared with the control.

Next, a flow cytometric analysis of the cell cycle was performed to evaluate the contribution of Gal-9 to cell cycle arrest during the suppression of pancreatic cancer cellular

proliferation. PK-1 cells were treated with 0.3  $\mu$ M Gal-9. In contrast to the expression levels of cyclin E and Cdk4, which decreased, Gal-9 did not alter the cell cycle of PK-1 cells (Fig. 3B and C). These results suggest that Gal-9 suppresses cell growth through tumor cell apoptosis but not through cell cycle arrest.

*The effects of Gal-9 on apoptosis-associated proteins in PK-1 cells.* The present study used an apoptosis array system to identify apoptosis proteins that were associated with the antitumor effect of Gal-9. An antibody array enabled the screening of 35 apoptosis-associated proteins in PK-1 cells in the presence and absence of Gal-9 (Fig. 4A). Gal-9 increased the expression levels of cytochrome c (Fig. 4B). Densitometry analysis demonstrated that the intensity of cytochrome c spots from Gal-9-treated cells relative to untreated cells was 129.7% (Fig. 4C).

*Effects of Gal-9 on p-RTKs in PK-1.* The present study used a p-RTK array system to identify key RTKs that were associated with the antitumor effect of Gal-9. An antibody array enabled the screening of 49 activated RTKs in PK-1 cells in the presence and absence of Gal-9 (Fig. 5A). Gal-9 reduced the expression levels of phosphorylated epidermal growth

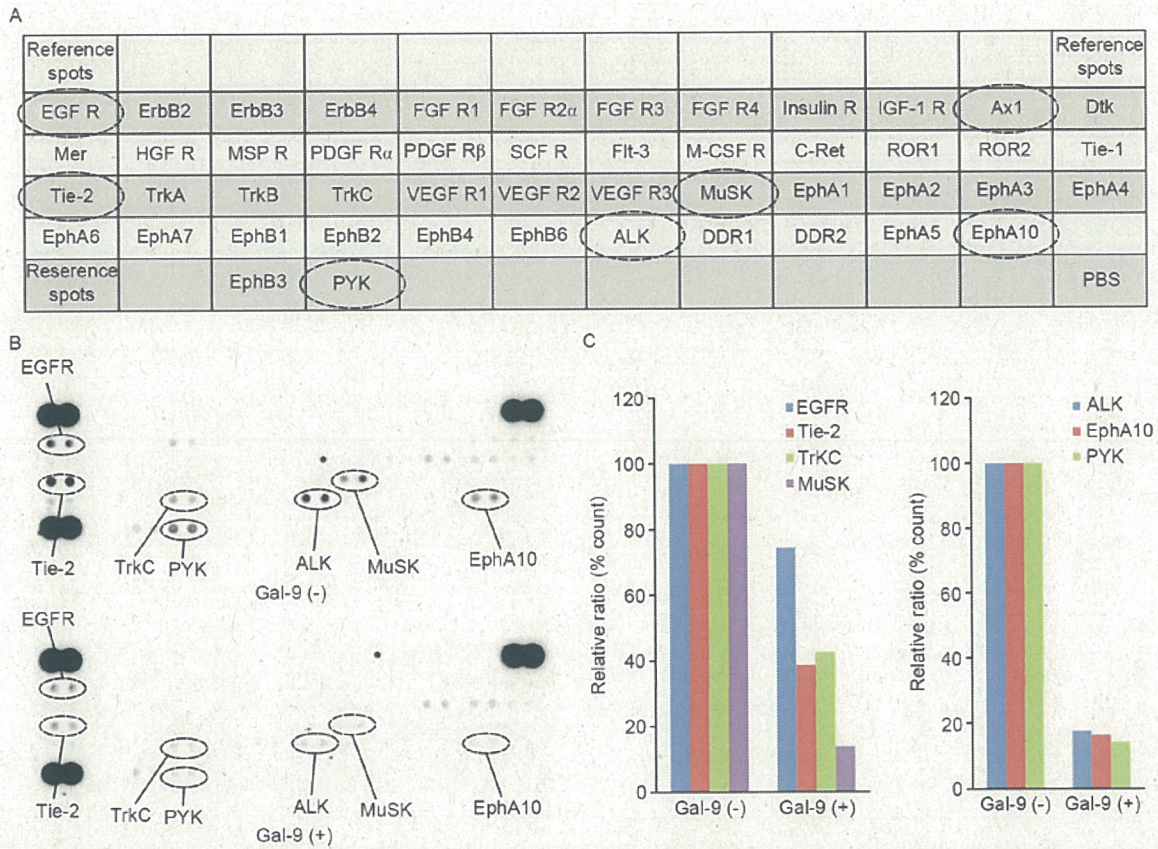


Figure 5. The effects of Gal-9 on pRTKs in PK-1 cells. (A) Template showing the locations of tyrosine kinase antibodies spotted onto a human pRTK array. (B) Representative expression of various phosphorylated tyrosine kinase receptors in PK-1 cells treated with or without Gal-9. (C) Densitometry analysis demonstrated that the intensities of p-EGFR, p-Tie-2, p-TrkC, p-MuSK, p-ALK, p-EphA10 and p-RYK spots from Gal-9-treated cells relative to untreated cells were 74.4, 38.6, 42.5, 13.8, 17.7, 16.5 and 14.3%, respectively. Gal-9, galectin-9; pRTK, phospho-receptor tyrosine kinase; p-EGFR, phospho-epidermal growth factor; p-Tie-2, phospho-tyrosine kinase with immunoglobulin-like and EGF-like domains-2; p-TrkC, phospho-tropomyosin receptor kinase C; p-MuSK, phospho-muscle-specific kinase; p-ALK, phospho-anaplastic tyrosine kinase; p-EphA10, phospho-erythropoietin-producing human hepatocellular receptors 10; p-RYK, phospho-receptor-like tyrosine kinase; ErbB2, human epidermal growth factor 2; FGF R1, fibroblast growth factor receptor 1; IGF-1R, insulin-like growth factor 1 receptor; Ax1, tyrosine-protein kinase receptor UFO; Dtk, Dtk receptor tyrosine kinase; Mer, Mer receptor tyrosine kinase; HGF R, hepatocellular growth factor receptor; MSP R, macrophage-stimulating protein receptor; PDGF R $\alpha$ , platelet-derived growth factor receptor  $\alpha$ ; SCF R, stem cell factor receptor; Flt-3, Fms-like tyrosine kinase 3; M-CSF R, monocyte colony stimulating factor; C-Ret, C-Te receptor tyrosine kinase; ROR1, receptor tyrosine kinase-like orphan receptor 1; Tie-1, tyrosine kinase with immunoglobulin-like and EGF-like domains-1; Trk, tropomyosin receptor kinase; VEGF R1, vascular endothelial growth factor receptor 1; Musk, muscle-specific kinase; EphA1, erythropoietin-producing human hepatocellular receptor A1; ALK, anaplastic tyrosine kinase; DDR1, discoidin domain receptor 1.

factor receptor (p-EGFR) and phosphorylated tyrosine kinase with immunoglobulin-like and EGF-like domains 2 (p-Tie-2), and it also reduced the expression of tropomyosin receptor kinase C (TrkC), muscle-Specific Kinase (MuSK), anaplastic lymphoma kinase (ALK), erythropoietin-producing human hepatocellular carcinoma cell (EphA10) and receptor-like tyrosine kinase (RYK) (Fig. 5B).

Densitometry analysis demonstrated that the intensities of p-EGFR, p-Tie-2, p-TrkC, p-MuSK, p-ALK, p-EphA10 and p-RYK spots from Gal-9-treated cells relative to untreated cells were 74.4, 38.6, 42.5, 13.8, 17.7, 16.5 and 14.3%, respectively (Fig. 5C).

*The effects of Gal-9 on the in vitro miRNA expression of cells treated with and without Gal-9.* Using a custom microarray platform, the present study analyzed the *in vitro* expression levels of 2,555 miRNA probes in tumorous tissues that were treated with and without Gal-9. In PK-1 cells treated with Gal-9, there were 13 upregulated and 6 downregulated miRNAs among the 2,555 miRNAs (Table I).

Unsupervised hierarchical clustering analysis using Pearson's correlation demonstrated that *in vivo*, tumorous tissues treated with Gal-9 clustered together and separated from untreated cell lines (Fig. 6).

**Discussion**

Pancreatic cancer has the worst 5-year survival rate of all malignancies due to its aggressive progression and resistance to therapy. Worldwide, the incidence of all types of pancreatic cancer ranges between 1 and 10 cases per 100,000 individuals, and it is generally higher in developed countries and among men (1). However, there is no effective screening tool to detect asymptomatic premalignant or early malignant tumors, and >90% of patients who receive a diagnosis of pancreatic cancer die from the disease (27). Thus, there is strong demand for new curative approaches to pancreatic cancer therapy.

The present data revealed that Gal-9 suppressed the cellular proliferation and tumor growth of human pancreatic cancer cell lines *in vitro*. The antitumor effect of Gal-9 in

Table I. Statistical results and chromosomal locations of miRNAs in PK-1 cells treated with and without Gal-9.

miRNA	Fold-change (treated/non-treated)	P-value	FDR	Chromosomal localization
<b>Upregulated</b>				
hsa-miR-6807-5p	2.27	0.00016	0.0045	19
hsa-miR-6784-5p	2.25	0.00016	0.0045	17
hsa-miR-4739	2.18	0.00016	0.0045	17
hsa-miR-6845-5p	2.04	0.00016	0.0045	8
hsa-miR-3196	1.91	0.00016	0.0045	20
hsa-miR-3649	1.88	0.00016	0.0045	12
hsa-miR-1202	1.88	0.00016	0.0045	6
hsa-miR-762	1.72	0.00016	0.0045	16
hsa-miR-204-3p	1.67	0.00016	0.0045	9q21.12
hsa-miR-4689	1.63	0.00016	0.0045	1
hsa-miR-575	1.55	0.00031	0.0071	4q21.22
hsa-miR-6765-5p	1.46	0.00031	0.0071	14
hsa-miR-4655-5p	1.33	0.00016	0.0045	7
<b>Downregulated</b>				
hsa-miR-301a-3p	0.46	0.00031	0.0071	17q22
hsa-miR-503-5p	0.56	0.00016	0.0045	Xq26.3
hsa-miR-422a	0.63	0.00016	0.0045	15q22.31
hsa-miR-30a-3p	0.66	0.00031	0.0071	6q13
hsa-miR-1913	0.68	0.00016	0.0045	6
hsa-miR-1285-3p	0.69	0.00016	0.0045	7q21-q22

miRNA, microRNA; Gal-9, galectin-9; FDR, false discovery rate; hsa-miR, human microRNA.

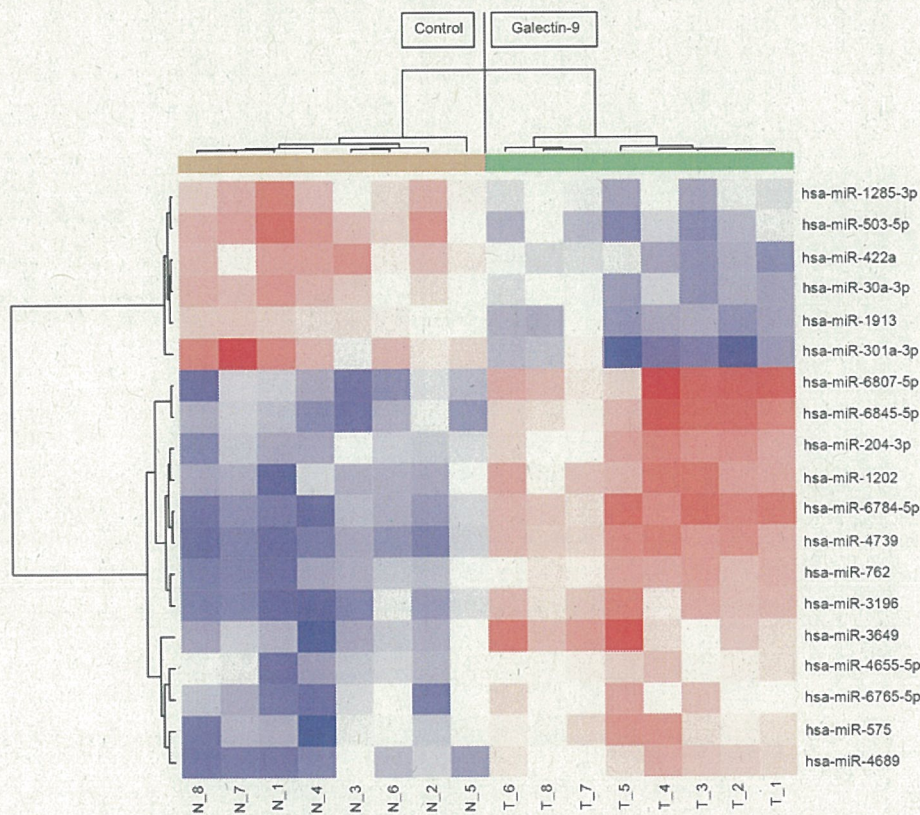


Figure 6. Hierarchical clustering of PK-1 cell tumor tissue with and without Gal-9. Tumor tissues clustered according to the expression profiles of 49 miRNAs that were differentially expressed in PK-1 cells with and without Gal-9. The analyzed samples are shown in the columns, and the miRNAs are presented in the rows. The miRNA clustering color scale shown at the top indicates the relative expression levels of miRNAs, with red and blue representing high and low expression levels, respectively. Gal-9, galectin-9; hsa-miR, human microRNA.

T cell hemostasis, cell aggregation and metastasis is well known (14,15). Previous findings suggest that Gal-9 inhibits the proliferation of hematologic malignancies, including multiple myeloma (18) and chronic myeloid leukemia (19), and significantly retards the tumor growth of myeloma xenografts in mice (18). Cell surface-associated Gal-9 triggered the aggregation of melanoma cells, indicative of Gal-9-mediated cellular adhesion and inhibition of cell detachment (28). In hematologic malignancies, Gal-9 may suppress cellular proliferation and tumor growth *in vitro* and *in vivo*. On the other hand, in solid malignancies, breast cancer cell lines with high levels of endogenous Gal-9 had a strong tendency to aggregate, whereas cells with low levels of Gal-9 did not (29). Importantly, ectopic expression of endogenous Gal-9 and treatment with recombinant Gal-9 triggered the formation of tight cellular clusters (28,29). Therefore, Gal-9 directly suppresses cellular proliferation and tumor growth and has therapeutic potential for several solid tumors.

Recombinant Gal-9 induces apoptosis and cell death through an apoptotic signaling pathway (18,19). Such apoptotic signaling was caspase-dependent and was induced by activation of the mitogen activated protein kinases, c-Jun N-terminal kinases and p38 in multiple myeloma cells (18). In addition, Gal-9 induced the proapoptotic Bcl-2 family member Noxa via activation of transcription factor 3, leading to the death of chronic myeloma cells (19). Various hematological malignancies are sensitive to apoptotic elimination by recombinant Gal-9. Cleavage of cytokeratin 18 (CK18) occurs as an early event during apoptosis following activation of apoptosis executioners, particularly effector caspases (30). However, CK18 remains intact during other types of cell death, including autophagy or necrosis (30). Several studies have made use of this phenomenon to detect cellular apoptosis at its early phase (31-33). Our data suggested that Gal-9 increases the levels of CCK18 in human pancreatic cancer cell lines. Additionally, using an apoptosis array, the present study revealed that the expression of cytochrome c was increased in Gal-9-treated pancreatic cancer cell lines. Cytochrome c release from damaged mitochondria is an early event in the intrinsic apoptosis pathway and contributes to caspase-9 activation. The present data suggest that Gal-9 may induce the apoptosis of pancreatic cancer cell lines in the intrinsic apoptosis pathway through caspase-dependent and caspase-independent pathways.

A previous study demonstrated that Gal-9 suppresses cellular proliferation and tumor growth in hepatocellular carcinoma and cholangiocarcinoma by inducing apoptosis but not cell cycle arrest (20,21). Although the expression levels of certain cell cycle-related proteins (Cdk4 and cyclin E) decreased 48 h after the addition of Gal-9, flow cytometry demonstrated that Gal-9 did not affect PK-1 pancreatic cancer cells at the G0-to-G1 transition *in vitro*. These data suggest that the antitumor effect of Gal-9 may not be associated with the reduction of various cell cycle-related proteins.

Since the discovery of these proteins, RTKs have been investigated as key regulators of the proliferation, differentiation and metastasis of cancer cells (34). Gal-9 reduced the expression levels of p-EGFR and phosphorylated Tie-2, TrkC, MuSK, ALK, EphA10 and RYK, according to the pRTK array. The EGFR family and their ligands are frequently

overexpressed in pancreatic cancer (35), and EGFR activity correlates with the prognosis of the patient (36).

The miRNAs that are associated with the antitumor effects of Gal-9 were assessed using miRNA expression arrays. miRNAs are small, endogenous, non-coding siRNAs that are 21-30 nucleotides in length and that modulate the expression of various target genes at the post-transcriptional and translational levels (37). miRNAs take part in fundamental molecular processes associated with pancreatic cancer initiation and progression, including the cell cycle, DNA repair, apoptosis, invasion and metastasis (38). Cluster analyses clearly demonstrated that Gal-9 treatment affected the extent of miRNA expression in pancreatic cancer cell lines. The present study identified 19 miRNAs that were differentially expressed in the cluster. These miRNAs are potential candidates to gauge the effectiveness of Gal-9 treatment, and provide clues regarding the molecular basis of the anti-cancer effects of Gal-9, particularly those mediated by miRNAs. Notably, miR301a was downregulated in a Gal-9-treated pancreatic cancer cell line. The overexpression of miR-301a has been previously demonstrated in pancreatic tumor tissue (39), hepatocellular carcinoma (40) and breast cancer (41). Additionally, miR-301a acts as a nuclear factor- $\kappa$ B activator in pancreatic cancer (42) and promotes proliferation and invasion in breast cancer. Thus, the present data suggest that miR-301a may be a candidate target for new therapeutic approaches to pancreatic cancer.

In conclusion, our results reveal that Gal-9 inhibits human pancreatic cancer cellular proliferation, possibly by inducing apoptosis through cytochrome c release, which is associated with the alteration of miRNAs. These findings suggest that Gal-9 may be a new therapeutic agent for the treatment of pancreatic cancer.

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