

Reversion of trichostatin A resistance via inhibition of the Wnt signaling pathway in human pancreatic cancer cells

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Received June 16, 2014; Accepted August 22, 2014

DOI: 10.3892/or.2014.3476

Abstract. Drug resistance is a major impediment to successful chemotherapy in pancreatic cancer (PC) patients. We investigated the effect of Wnt/ β -catenin signaling inhibition by wnt-c59 on chemoresistance in a trichostatin A-resistant Panc-1 cell line (Panc-1/TSA). Panc-1/TSA cells were treated with the Wnt/ β -catenin signaling inhibitor wnt-c59 ($10 \mu\text{mol} \cdot \text{l}^{-1}$) and/or trichostatin A (TSA; $10 \mu\text{mol} \cdot \text{l}^{-1}$) for 24 h. CCK-8 assay was utilized to analyze the interactive effect of TSA and wnt-c59 on induction of apoptosis of the Panc-1/TSA cells. Cell apoptosis was measured by flow cytometry. Real-time PCR and western blotting were used to assess Wnt/ β -catenin signaling, epithelial-mesenchymal transition (EMT) and multidrug resistance (MDR). Real-time cell analysis (RTCA) was used to detect the cell migration ability. After wnt-c59 treatment for 24 h, relative genes and transcriptional targets of Wnt/ β -catenin signaling were downregulated ($P < 0.05$). CCK-8 assay indicated that the combination of TSA and wnt-c59 had a synergistic effect on induction of Panc-1/TSA cell apoptosis. As detected by FACS, cell apoptosis rates increased significantly ($P < 0.05$). The results of RTCA showed that the cell indices of the control group, wnt-c59 group, TSA group and TSA+wnt-c59 combination group were 1.2842 ± 0.0257 , 1.2155 ± 0.0282 , 1.2533 ± 0.0194 and 0.8541 ± 0.0250 , respectively. In accordance, MMP-9 protein in the wnt-c59 treatment groups was decreased compared to the non-wnt-c59 treatment

groups. Meanwhile, E-cadherin protein was upregulated and vimentin protein was downregulated, both of which are characteristic markers of EMT. Chemoresistant gene MDR1 and P-glycoprotein (P-gp) in the wnt-c59 treatment groups had a reduced expression compared to the non-wnt-c59 treatment groups. This study revealed that TSA sensitivity, migration ability, and the EMT phenotype in Panc-1/TSA cells were reversed following Wnt/ β -catenin signaling inhibition.

Introduction

Over the past three decades, pancreatic cancer (PC) has remained one of the most lethal cancers worldwide, leading to an estimated 227,000 deaths annually (1). Despite some efforts to improve diagnosis and treatment strategies (including surgery, chemotherapy, radiation therapy and immunotherapy), the overall 5-year survival rate of all PC patients is dismal, at approximately 5%. At present, less than 20% of PC patients are indicated for surgery. Chemotherapy has become the standard treatment of locally advanced and metastatic PC patients when disregarding surgical resection as the optimal therapeutic option (2,3). However, patients undergoing continued chemotherapy ultimately develop chemoresistance (4). Hence, effective therapeutic strategies for increasing the sensitivity of chemotherapy or reducing the multidrug resistance (MDR) of PC cells are urgently needed to improve patient survival.

To date, the principal mechanisms of chemoresistance may include increased efflux, activation of detoxifying systems, DNA repair, and failure to undergo apoptosis (5). Numerous studies have indicated that many genes, proteins and cell-autonomous signaling pathways have important roles in the chemoresistance of PC patients (6-9). Further analysis of these signaling pathways revealed that Wnt/ β -catenin signaling played important roles in drug exclusion from cells, changes in the enzymes metabolizing drugs, or the increased cell resistance to stress and apoptosis (4). The activation of Wnt/ β -catenin signaling was found to stimulate angiogenesis and regulate the cell cycle and apoptosis (10).

Among the multiple genetic and cytogenetic alterations that characterize human tumors, great emphasis has been recently given to epigenetic events, such as DNA methylation or histone acetylation (11). Alterations in histone acetyltrans-

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Abbreviations: APC, adenomatous polyposis coli; PC, pancreatic cancer; MDR, multidrug resistance; P-gp, P-glycoprotein; GSK3 β , glycogen synthase kinase 3 β ; TCF/LEF1, T cell factor/lymphoid enhancer-binding factor 1

Key words: drug resistance, epithelial-mesenchymal transition, pancreatic cancer, trichostatin A, Wnt/ β -catenin signaling pathway

ferase (HAT) or histone deacetylase (HDAC) activity occur in numerous cancers and have prompted the search for pharmacological agents capable of inhibiting these enzymes. HDAC inhibitors show a very low toxicity *in vivo* and have become a new class of chemotherapeutic agents against cancer (12,13). Trichostatin A (TSA) is one of the HDAC inhibitors (HDACi) and promotes histone hyperacetylation and strongly induces apoptosis by altering the expression of various apoptotic genes (14). However, in our previous study, gradually increasing concentrations of TSA in the PC cell line Panc-1 induced apoptosis. We found that MDR of the Panc-1 cells increased and the apoptotic effects of TSA decreased. Meanwhile, Wnt/ β -catenin signaling became highly activated. The precise mechanisms of HDACi-mediated chemosensitization, with regard to specific signal pathways/networks, remain somewhat speculative.

Based on these findings, we used a series of approaches to ascertain whether wnt-c59 inhibits Wnt/ β -catenin signaling. In conclusion, TSA-resistant Panc-1 cells (Panc-1/TSA) were re-sensitized to TSA.

Materials and methods

Reagents. TSA, dimethyl sulfoxide (DMSO), propidium iodide (PI) and Annexin V-FITC apoptosis detection kits were purchased from Sigma-Aldrich (St. Louis, MO, USA). Dulbecco's modified Eagle's medium (DMEM), fetal bovine serum (FBS), trypsin-EDTA, penicillin, streptomycin and TRIzol were obtained from Gibco-BRL (Invitrogen, Grand Island, NY, USA). TCF, E-cadherin, and MMP-9 antibodies were obtained from Cell Signaling Technology, Inc. (Beverly, MA, USA). Antibodies against Frizzled 7, β -catenin and phospho- β -catenin were purchased from Abcam (Cambridge, UK). GAPDH, c-myc and horseradish peroxidase (HRP)-conjugated goat anti-rabbit secondary antibodies were obtained from Bio-World Co. (Nanjing, China). Reagents were used at the following working concentrations: $10 \mu\text{mol} \cdot \text{l}^{-1}$ of TSA; $10 \mu\text{mol} \cdot \text{l}^{-1}$ of Wnt/ β -catenin signaling inhibitor wnt-c59 (Gene Operation Inc., St. Joseph, MI, USA).

Cell lines and culture conditions. The cell line Panc-1/TSA was obtained from Dr Liu Biao (Wenzhou Medical University). Monolayer cultures of Panc-1/TSA cells were maintained in DMEM supplemented with 10% FBS, 100 U/ml penicillin, and 100 $\mu\text{g}/\text{ml}$ streptomycin at 37°C in a humidified chamber of 95% air and 5% CO_2 , and the medium was replaced every 2-3 days. When reaching subconfluency, the cells were harvested by trypsinization, and recultured according to the experimental requirements. Cells were treated with TSA ($10 \mu\text{mol} \cdot \text{l}^{-1}$) or wnt-c59 ($10 \mu\text{mol} \cdot \text{l}^{-1}$) in 2 ml of medium for 24 h.

Real-time PCR analysis. Total cellular RNA was extracted from the different groups using TRIzol reagent (Invitrogen). Subsequently, mRNA was reverse transcribed to single-stranded cDNA using the ReverTra Ace[®] RT kit (Toyobo Co., Ltd., Osaka, Japan) according to the manufacturer's instructions. The cDNAs obtained were amplified by PCR using ABI 7500 Thermal Cycler Dice[™] real-time system (Applied Biosystems, USA). The thermal cycler protocols included 3 min at 95°C and 40 cycles of 5 sec at 95°C and 30 sec at 60°C .

The primer sequences are listed in Table I. For each sample, relative mRNA expression levels were derived from the ratio of their expression to GAPDH expression as an internal standard using the $2^{-\Delta\Delta\text{Ct}}$ method (15).

Western blot analysis. Cells were harvested and lysed in lysis buffer containing protease inhibitors after the treatment with wnt-c59 and/or TSA, as required. The lysates were centrifuged at $12,000 \times g$ for 15 min at 4°C . The supernatant was collected, and protein concentrations were determined by BCA protein assay according to the manufacturer's instructions. Primary antibodies were diluted 1:500-1:1,000 in 5% skimmed milk with TBS-T [10 mM Tris-HCl (pH 8.0), 150 mM NaCl and 0.1% Tween-20]. Protein samples were solubilized and boiled in SDS sample buffer for 5 min and then separated using 12% SDS-PAGE at 200 V for 1 h. Subsequently, the separated proteins were transferred to a polyvinylidene difluoride membrane. Following incubation in blocking solution consisting of 5% skimmed milk in TBS-T for 1 h at room temperature and overnight incubation at 4°C with the primary antibodies including Frizzled 7, β -catenin, phospho- β -catenin, c-myc, TCF and GAPDH, the membrane was washed and then probed with respective horseradish peroxidase-conjugated secondary antibodies for 1.5 h at room temperature. After washing three times with TBS-T, hybridization was visualized using the ECL chemiluminescence detection system (Kodak). Expression levels of the proteins were compared to the control based on the relative intensities of the bands.

Immunofluorescent staining for assessment of cell morphology. Panc-1/TSA cells were incubated for 24 h using the desired method. After three washings with PBS, cells were fixed in 4% paraformaldehyde for 10 min at room temperature. The fixed cells were permeabilized with PBS containing 0.3% Triton X-100 for 10 min. After soaking the cells in 10% goat serum (Solarbio, Beijing, China) with PBS for 30 min, they were treated with the anti-Frizzled 7 antibody (1:200) in PBS overnight. The cells were then washed with PBS three times followed by incubation at 37°C for 30 min. The secondary antibody solution mixed with PBS was incubated for 1 h in a dark room. After washing cells three times with PBS, they were covered with an anti-fade medium containing the nuclear stain 4',6-diamidino-2-phenylindole (MultiSciences Biotech, Hangzhou, China). Subsequently, they were examined using a DM4000B automated upright microscope system (Leica, Wetzlar, Germany).

Drug combination studies. Panc-1/TSA cells were trypsinized and seeded in 96-well cell culture plates (1×10^4 cells/well in 100 μl culture medium) for overnight adhesion. They were then treated with wnt-c59 and/or TSA, as required, for an additional 24 h. Next, cells treated in each well were maintained with 10 μl of CCK-8 (Dojin, Japan) and 100 μl DMEM at 37°C in 5% CO_2 for 2 h. The optical density (OD) of each well was then measured using the Varioskan microplate reader (Thermo, USA) at a 450-nm wavelength. The cell inhibitory rate was calculated according to the following equation: The cell inhibitory rate = $1 - [(\text{OD experiment} - \text{OD blank}) / (\text{OD control} - \text{OD blank})] \times 100\%$. We used the Zhengjun Jin method to determine whether the two drugs interact (16). The general

Table I. The primers used in the real-time PCR.

Gene	Sense (5'-3')	Antisense (5'-3')
β -catenin	GCCAAGTGGGTGGTATAGAGG	GCGGGACAAAGGGCAAGA
Frizzled 7	AGATCTGCGTGGGCCAGAAC	AGCGCCTCTCCTCCTCTTA
GSK3 β	AGACGCTCCCTGTGATTTATG	AACAAGAGGTTCTGCGGTTT
TCF	AGGCAGGGAGAATGCGACTC	GCGTTGAGGTTGGTGCCTTC
cyclin D1	AGGCGGAGGAGAACAAACAG	TGAGGCGGTAGTAGGACAGGA
c-myc	CCACCACCAGCAGCGACT	CTTGACCCTTTGGCAGCA
MDR1	AGGTTCTGGGAAGATCGC	CCTCAAAGAGTTTCTGTATGGTA
GAPDH	TCCCATCACCATCTTCCAGG	GATGACCCTTTTGGCTCCC

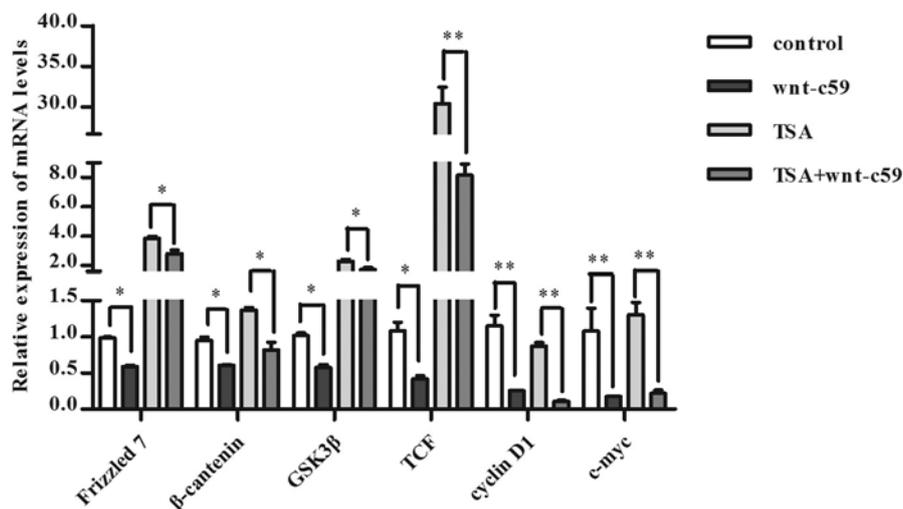


Figure 1. Expression of genes associated with the Wnt/ β -catenin signaling pathway detected by real-time RT-PCR. The mRNA expression levels of Frizzled 7, β -catenin, GSK3 β , TCF, cyclin D1 and c-myc were significantly decreased after wnt-c59 and/or TSA treatment for 24 h. The data are represented as the mean \pm SD of at least three independent experiments. * P <0.05; ** P <0.01.

equation is given by $q = E_{a+b}/(E_a + E_b - E_a \times E_b)$, where E_a and E_b in the denominator are the cell inhibition rates of TSA and wnt-c59 alone, respectively, whereas E_{a+b} in the numerator is the cell inhibitor rate of TSA and wnt-c59 in combination. $q < 0.85$, $q \sim 0.85$ -1.15, or $q > 1.15$ generally indicates an antagonistic, an additive, or a synergistic effect, respectively.

Flow cytometry. We used a monoclonal P-glycoprotein (P-gp) antibody (CD243, antibodies-online; Ebioscience, San Diego, CA, USA; 1:50) and the Annexin V apoptosis detection kit (Sigma-Aldrich). We added wnt-c59 and/or TSA into the Panc-1/TSA cells in the desired groups, collected 1.5×10^5 cells by centrifugation, and resuspended cells in $500 \mu\text{l}$ of 1X binding buffer. A single-cell suspension was mixed with anti-CD243 at a ratio of 50:1 and stained for 30 min at 4°C or with $5 \mu\text{l}$ of Annexin V-FITC and $10 \mu\text{l}$ of propidium iodide for 5 min in the dark. Cells were washed with PBS. CD243-positive cells and apoptotic cells were detected by flow cytometry. All experiments were repeated three times.

Real-time cell analysis (RTCA) experiments. The migratory ability of the Panc-1/TSA cells was assessed by xCELLigence real-time cell analyzer (RTCA) DP device, which was placed in

a humidified incubator at 37°C with 5% CO_2 . Panc-1/TSA cells (Roche Applied Science, Basel, Switzerland) were seeded into the upper chamber of a two-chamber instrument separated by a porous membrane (CIM-plate 16). The cells either attached or migrated directly through the pores to the bottom side of the membrane. In either device, this increased the electrical impedance of integrated gold microelectrodes. The electrical impedance was displayed as a dimensionless parameter termed cell index. The cell index represents the capacity of cell migration, and the slope of the curve can be related to the migration velocity of the Panc-1/TSA cells. The cell index thus reflects the tumor cell migratory capacity. Prior to cell seeding the bottom side of the wells from the upper chamber of the CIM-plate 16 was coated with 30 ml of collagen I. Furthermore, 40,000 cells/well suspended in culture medium containing 10% FBS were then seeded into the upper chamber according to the manufacturer's manual (17). The cell index was determined every 15 min for up to 24 h using RTCA software (ver. 1.2; Roche Diagnostics).

Statistical analysis. All data are expressed as mean values \pm SD. Statistical analysis was conducted using SPSS 13.0 software. The analysis of variance (ANOVA) was used to test

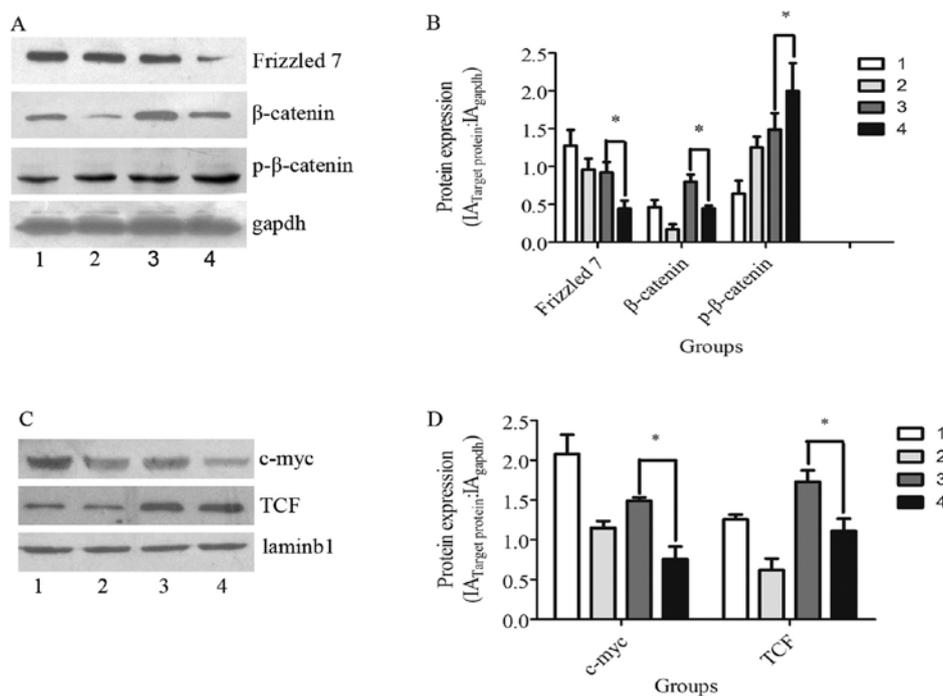


Figure 2. Protein expression of Frizzled 7, phosphorylated- and non-phosphorylated- β -catenin, TCF and c-myc measured by western blotting in Panc-1/TSA cells. (B and D) were the semiquantitative results of A and C, respectively. Lane 1: control group; lane 2: wnt-c59 group; lane 3: TSA group; lane 4: TSA+wnt-c59 group. The data was represented by the mean \pm SD of at least three independent experiments * $P < 0.05$, TSA group compared to TSA+ wnt-c59 group.

the difference in the means between groups. Differences were considered statistically significant at $P < 0.05$.

Results

Downregulation of the expression levels of mediators of Wnt/ β -catenin signaling following wnt-c59 treatment. To determine the effects of TSA on mRNA expression of the mediators in Panc-1/TSA cells in the presence or absence of wnt-c59, we analyzed the level of mRNAs by real-time PCR. The results revealed that treatment with $10 \mu\text{mol} \cdot \text{l}^{-1}$ wnt-c59 significantly decreased mRNA expression levels of Frizzled 7, β -catenin, downstream glycogen synthase kinase 3β (GSK3 β) and T cell factor/lymphoid enhancer-binding factor 1 (TCF/LEF1). Consistently, transcriptional targets downstream of this pathway, such as c-myc and cyclin D1, were downregulated (Fig. 1). We confirmed these changes at the protein level by western blotting. As shown in Fig. 2, the protein levels of Frizzled 7, β -catenin, TCF and c-myc were decreased, correlating with the trend found for mRNA levels following wnt-c59 treatment. Immunofluorescence staining was performed to detect the cellular localization of the ligand-receptor Frizzled 7 protein, which revealed a membranous accumulation in the non-wnt-c59 treatment groups compared to the wnt-c59 treatment groups (Fig. 3). Our findings suggest that the inhibition of Wnt/ β -catenin activity in the Panc-1/TSA cells was due to downregulation of the expression of Wnt ligands, their receptors and modulators of Wnt signaling.

Interaction of TSA and wnt-c59 in the Panc-1/TSA cells. The results of the CCK-8 assay showed that the cell inhibitory rates of the wnt-c59 group, TSA group and TSA+wnt-c59 group were

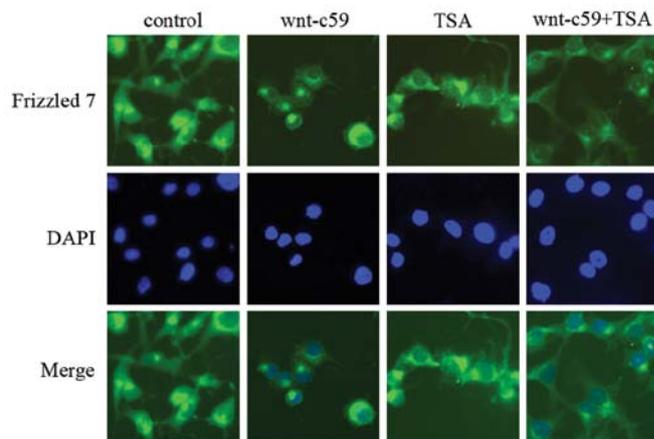


Figure 3. Expression of the ligand-receptor Frizzled 7 detected by immunofluorescence analysis. Minimal staining (membranous expression of Frizzled 7) was observed in Panc-1/TSA cells of wnt-c59 treated groups, whereas cells of non-wnt-c59 treated groups showed high levels of fluorescent staining readily distinguished from background. Zoom, $\times 400$.

0.0028 ± 0.0455 , 0.1960 ± 0.0455 and 0.2541 ± 0.0059 , respectively. According to a method previously reported (16), $q = 1.282 > 1.15$, indicated that the combination of TSA and wnt-c59 had a synergistic effect on inducing Panc-1/TSA cell apoptosis.

Effect of wnt-c59 on the apoptosis of Panc-1/TSA cells. To investigate the effect of wnt-c59 on the apoptosis of Panc-1/TSA cells, we detected cell apoptosis rates using flow cytometry. The cell apoptosis rates of the control group, wnt-c59 group, TSA group, and TSA+wnt-c59 group were 13.86 ± 1.47 , 11.94 ± 3.32 , 20.97 ± 5.62 and $37.24 \pm 8.2\%$, respec-

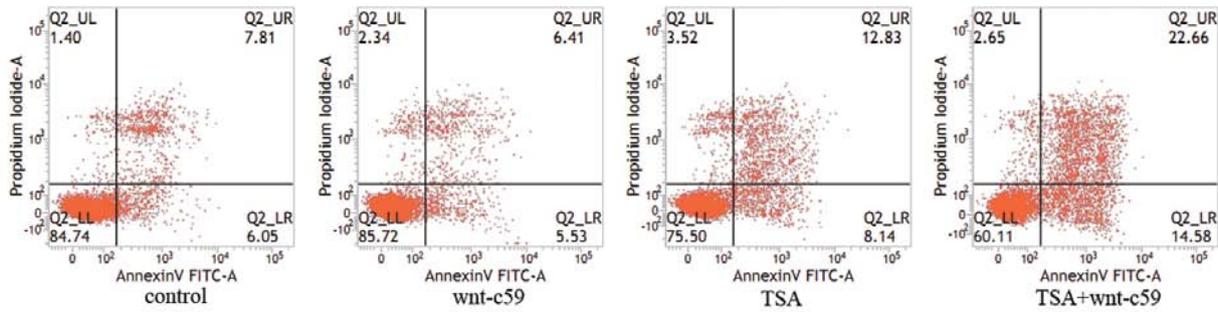


Figure 4. Induction of apoptosis by wnt-c59 and/or TSA treatment. Panc-1/TSA cells were treated with wnt-c59 and/or TSA for 24 h. The percentage of apoptosis was calculated by Annexin V⁺ (early apoptosis) and Annexin V⁺/propidium iodide⁺ staining (late apoptosis).

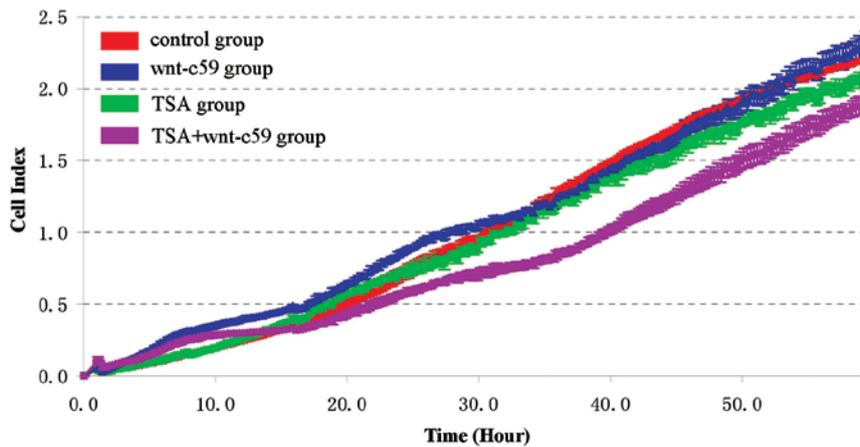


Figure 5. Real-time measurement of the cell migration in the Panc-1/TSA cells using the xCELLigence DP system. Panc-1/TSA cells were stimulated with wnt-c59 and/or TSA in serum-reduced medium during the assay as indicated by the color code. At 24, 36 and 48 h time points, differences between the TSA+wnt-c59 (purple curve) and the TSA treatment group (green curve) were significant (P<0.05) but not between the control and the wnt-c59 group.

Table II. Migration data of various groups measured by RTCA.

Time-interval (h)	Control group			Wnt-c59 group			TSA group			TSA+Wnt-c59 group		
	Mean CI	SD	CV (%)	Mean CI	SD	CV (%)	Mean CI	SD	CV (%)	Mean CI	SD	CV (%)
24 ^a	0.7075	0.0393	5.5	0.7144	0.031	4.3	0.8579	0.0172	2	0.5694 ^a	0.0207	3.6
36 ^a	1.2842	0.0257	2	1.2155	0.0282	2.3	1.2533	0.0194	1.5	0.8541	0.025	2.9
48 ^a	1.8633	0.0105	0.5	1.6994	0.045	2.6	1.8315	0.0485	2.6	1.4153	0.0642	4.5

Mean cell index (CI) represents the capacity for cell migration. CV represents the well-to-well variations. SD, standard deviation. ^aP<0.05.

tively (P<0.05). The results of FACS detection showed that the apoptosis induced by the combination of TSA and wnt-c59 was significantly increased compared to the treatment with TSA or wnt-c59 alone (Fig. 4).

Reduced metastatic ability and reversal of EMT of Panc-1/TSA cells after treatment with the combined chemotherapeutics.

We investigated the effect of inhibition of the Wnt/β-catenin pathway on migratory ability. The results of RTCA showed that the cell indices of the control group, wnt-c59 group, TSA group, and TSA+wnt-c59 combination group were 1.2842±0.0257, 1.2155±0.0282, 1.2533±0.0194 and 0.8541±0.0250, respec-

tively (P<0.05) (Fig. 5; Table II). In accordance, MMP-9 protein in the wnt-c59 treatment groups decreased compared to the non-wnt-c59 treatment groups. Meanwhile, the results of the western blotting indicated that the level of E-cadherin and vimentin protein, characteristic markers of EMT, was upregulated and downregulated, respectively (Fig. 6).

Inhibition of the Wnt/β-catenin pathway decreases the expression of the MDR gene and protein. Various mechanisms contribute to MDR. One of the best-studied mechanisms is the increased expression of P-gp protein, which acts as a drug efflux pump lowering the intracellular concentration

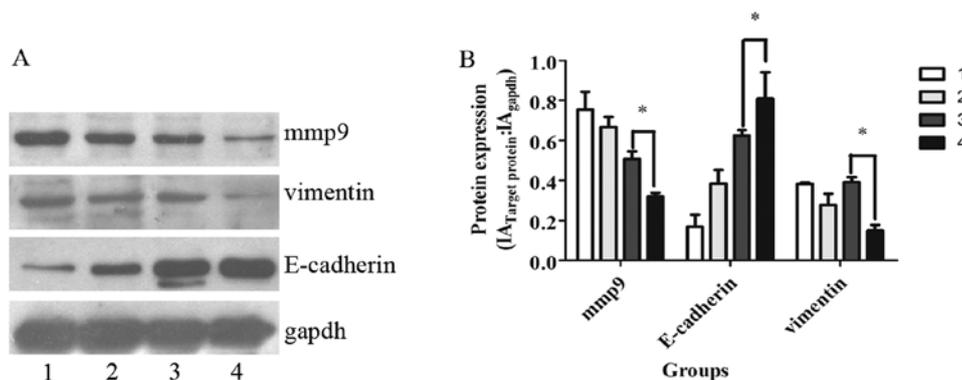


Figure 6. Effect of the inhibition of Wnt/ β -catenin on markers of EMT and cell migration. Panc-1/TSA cells in the wnt-c59 treatment groups exhibited lower expression levels of MMP-9 and vimentin proteins and higher expression levels of E-cadherin protein associated with EMT compared to the non-wnt-c59 treatment groups. A representative western blot is shown. (B) Semiquantitative results of A. Lane 1, control group; lane 2, wnt-c59 group; lane 3, TSA group; lane 4, TSA+wnt-c59 group. The data are represented as the mean \pm SD of at least three independent experiments. * P <0.05, TSA group compared to the TSA+wnt-c59 group.

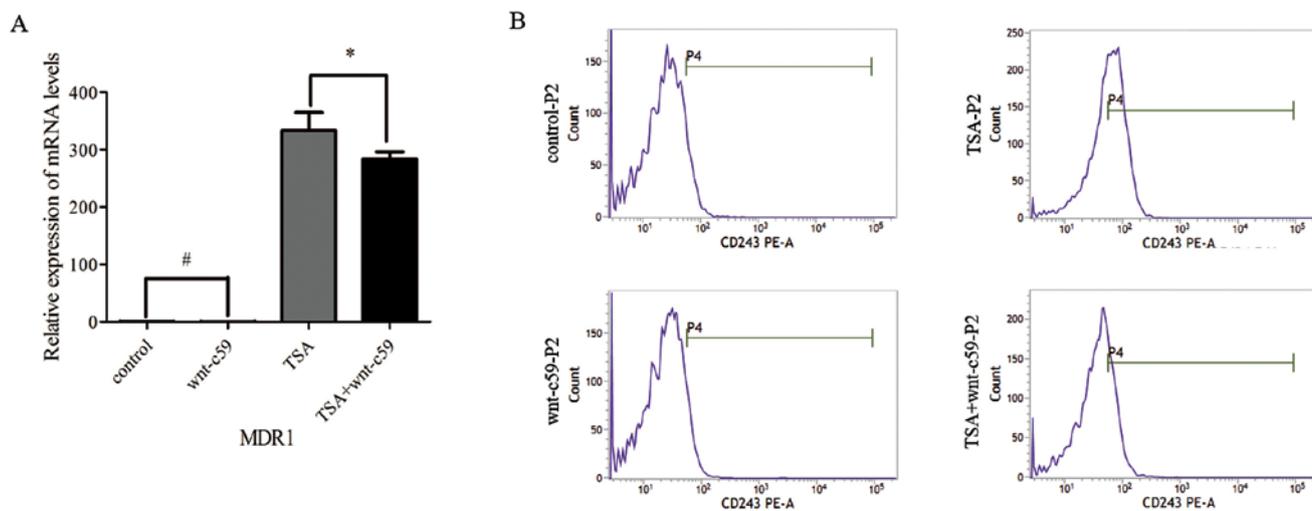


Figure 7. Downregulation of the MDR1 gene and P-gp protein in the Panc-1/TSA cells. (A) Real-time PCR demonstrated that the MDR1 transcript level was decreased after wnt-c59 treatment compared to that in the non-wnt-c59 treatment group. (B) FACS demonstrated that P-gp protein was downregulated after wnt-c59 treatment compared to that without wnt-c59 treatment.

Table III. Data of the P-gp protein results detected using FACS.

Groups	Events	% Parent	% Grandparent	% Total	FSC-A mean	SSC-A mean
Control	720	8.53	7.20	7.20	158,412	139,773
Wnt-c59	704	7.96	7.04	7.04	131,582	128,787
TSA	4,429	55.15	44.29	44.29	116,918	128,910
TSA+Wnt-c59	1,890	24.56	18.9	18.9 ^a	138,996	147,827

^a P <0.05, TSA+wnt-c59 group compared with the TSA group.

of cytotoxic drugs (18). We detected expression of the MDR gene (MDR1) by real-time PCR. The results revealed that chemoresistant gene MDR1 in the wnt-c59 treatment groups had a lower expression compared to the levels in the non-wnt-c59 treatment groups (P <0.05, Fig. 7A). We confirmed a similar change in the expression of chemoresistant protein P-gp using FACS (Fig. 7B; Table III).

Discussion

The aggressive and deadly biological characteristics of pancreatic cancer (PC) commonly result in the inability of the patients to undergo curative resection after diagnosis and thus chemotherapy becomes preferred treatment (19,20). Unfortunately, PC often develops intrinsic resistance to continued chemo-

therapy (21). Great advances in studying the mechanisms of tumor chemoresistance have been made. Studies have revealed that the Wnt/ β -catenin signaling pathway plays an important role in cell proliferation, differentiation, invasion and migration in many types of cancers. The Wnt/ β -catenin signaling pathway has become a frequently studied mechanism of cancer chemoresistance including that in PC (22,23). However, very few studies have used chemical inhibitors of Wnt/ β -catenin signaling to overcome chemoresistance in PC therapy. We assumed that wnt-c59, one of the Wnt/ β -catenin signaling inhibitors, might inhibit Wnt/ β -catenin signaling, decreasing the expression of drug-resistant genes and proteins and the capacity of migration, reversing the EMT phenotype, and increasing the apoptosis induced by chemotherapeutic drugs in PC. Consequently, the drug susceptibility of Panc-1/TSA cells may be enhanced.

The Wnt/ β -catenin signaling pathway represents the cross-link between a series of signaling pathways. Classical Wnt/ β -catenin includes mainly extracellular factor wnt protein, transmembrane receptor protein β -catenin, and nuclear transcription factor TCF protein (24). β -catenin, called the CTNNB1 gene, is a type of adhesion molecule. β -catenin mediates intercellular adhesion and maintains cell morphology by binding to the intracellular domain of E-cadherin. β -catenin is also involved in Wnt signaling transduction. In the absence of Wnt signaling, β -catenin in the cytosol binds mostly to the intracellular domain of E-cadherin and is attached to the cytoskeleton protein actin by α -cadherin to mediate intercellular adhesion. A part of β -catenin, which is complexed with adenomatous polyposis coli (APC) and Axin, is phosphorylated by casein kinase1 (CK1) as well as glycogen synthase kinase 3 β (GSK3 β) and is finally degraded. T cell factor/lymphoid enhancer-binding factor 1 (TCF/LEF1), due to the low β -catenin in the cytosol, is not activated since it binds to the transcription suppressor (25). In the presence of Wnt signaling, the signaling is initiated by Wnt proteins binding to the receptors of the Frizzled family on the cell surface. The activation of cytoplasmic degradation complex, comprising Axin, APC, GSK3 β and CK1, is inhibited, leading to the accumulation of cytosolic β -catenin. Cytosolic β -catenin enters the nucleus and forms a complex with TCF/LEF1 to activate the transcription of Wnt target genes, such as MDR1, cyclin D1, c-myc, MMPs and IL-8 (10,26-29).

The downstream target genes of the Wnt/ β -catenin signaling pathway, such as MDR1, cyclin D1, c-myc, MMPs and IL-8, are related to chemoresistance. Overexpression of P-gp protein and the MDR1 gene, markers of MDR, is broadly known to limit the efficacy of chemotherapeutic drugs in inducing cancer cell apoptosis (30). Accumulating data indicate that several signaling pathways associated with the progression of tumor chemotherapy, such as Hedgehog, K-ras, TGF- β , Notch and Wnt/ β -catenin pathway signaling, play a pivotal role in the induction of MDR (31). Previous studies have demonstrated that the activation of Wnt/ β -catenin signaling contributes to the chemoresistance of human neuroblastomas or rat brain endothelial cells through upregulation of P-gp expression (32,33). In a study of colon cancer the product of the MDR1 gene was upregulated with high expression of β -catenin (34). The data suggest that the role of Wnt/ β -catenin signaling is also involved in chemoresistance in PC (22). Our

study results showed that the MDR1 gene and P-gp protein were downregulated after Wnt/ β -catenin signaling was inhibited by wnt-c59. With Wnt signaling activation, increasing nuclear β -catenin could promote high expression of c-myc and cyclin D1, which are involved in cell proliferation (35). Moreover, Kornmann *et al* and Qiao *et al* showed that chemoresistance genes and proteins were decreased after inhibition of expression of cyclin D1 (36,37). In our study, after treatment with wnt-c59 in Panc-1/TSA cells, the transcriptional activity of Wnt/ β -catenin was inhibited and the apoptosis of Panc-1/TSA cells was significantly increased as detected by FACS. Expression of cyclin D1 and c-myc was also downregulated. We speculate that one of the mechanisms reversing drug resistance involved inhibition of Wnt/ β -catenin signaling, which re-sensitized the cells to apoptosis and downregulated cyclin D1. At the same time, c-myc may decrease the expression of multiple chemoresistance genes and proteins.

In addition to multiple chemoresistance genes and proteins, accumulating evidence suggests that chemoresistance is associated with the acquisition of the EMT phenotype in PC cells (38,39). Li *et al* found that PC cells that were sensitive to gemcitabine showed high expression of epithelial marker E-cadherin, whereas PC cells that were resistant to gemcitabine exhibited high expression of mesenchymal markers (including vimentin and ZeB1) at the gene and protein levels (40). These findings suggest that the EMT phenotype contributes to drug resistance in patients with PC. Moreover, drug-resistant cells from PC patients were more tumorigenic and had higher metastatic potential compared to cancer cells that were not drug resistant (41,42). This propensity is largely responsible for the poor prognosis of PC with chemoresistance. Upon Wnt ligand binding to its receptors, β -catenin enters the nucleus and transcriptionally upregulates VEGF and IL-8. IL-8 can induce overexpression of MMP-2 and MMP-9, which can dissolve the extracellular matrix (ECM) in direct and indirect ways and serve as the initiator of proliferation and movement of endothelial cells (33,43,44). In the present study, after inhibition of Wnt/ β -catenin signaling, the results of RTCA revealed that the migration ability of Panc-1/TSA cells in the TSA and wnt-c59 combination group was decreased significantly compared to the TSA alone group. We also found that the expression of E-cadherin protein was upregulated while the expression of vimentin and MMP-9 protein was downregulated. These results showed that the EMT phenotype was partly reversed after inhibition of Wnt/ β -catenin signaling.

In conclusion, inhibition of the Wnt/ β -catenin pathway can reverse drug resistance by downregulating the expression of multiple chemoresistance genes, proteins, and the EMT phenotype and by increasing cell apoptosis. Targeting Wnt/ β -catenin signaling could be useful for devising targeted therapeutic approaches for use in combination with conventional therapeutics to treat human PC. Compared to conventional approaches, these new treatments are expected to lead to better outcomes.

Acknowledgements

The authors are grateful for the funding from the Health Ministry of Zhejiang Province (Grant no. 11-ZC24), Zhejiang Provincial Top Key Discipline in Surgery, Administration of Traditional Chinese Medicine of Zhejiang Province (Grant

no. 2012-XK-A28) and the Wenzhou Government (Grant no. Y20090028), China.

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