

# Suppression of Akt1 expression by small interference RNA inhibits SGC7901 cell growth *in vitro* and *in vivo*

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**Abstract.** The Akt/PKB kinase family, including Akt1, 2 and 3, plays critical roles in regulating cell growth, proliferation, survival, metabolic and many other cellular activities. Recent evidence indicates that PKB/Akt is frequently constitutively active in many types of human cancer including gastric cancer. In the present study, we applied immunohistochemistry to tissue microarray to detect the expression of Akt1, followed by Akt1 small interference RNA (siRNA) to examine knock down of the Akt1 gene on the growth inhibition of human gastric cancer SGC7901 cells. Our results indicate that the expression of Akt1 was significantly increased in gastric cancer compared to normal gastric tissue and adjacent non-cancer tissue. The *in vitro* study shows that cell growth was significantly inhibited and G0/G1 arrest was observed in siRNA-Akt1-treated group. *In vivo*, the size of tumors was significantly smaller in SGC7901 subcutaneous mice model treated with siRNA-Akt1 than those treated with siRNA-nonsense and PBS. Our studies demonstrated siRNA-Akt1 can inhibit Akt1 expression, exerted growth inhibition effect on SGC7901 cells *in vitro* and *in vivo*. Suppression of Akt1 expression by siRNA could be a new strategy in gastric cancer treatment.

## Introduction

Over the past five decades, the mortality associated with gastric cancer has decreased markedly in the world (1,2). However,

in some individual cases, five-year survival rate after diagnosis of gastric cancer is still low (3,4). Gastric cancer remains the second leading cause of cancer death worldwide (1). In the USA alone, gastric cancer accounts for 2% of all cancer deaths (1).

It is believed that various human cancers, including gastric carcinoma, develop through the accumulation of several genetic alterations such as the inactivation of tumor suppressor genes or the activation of oncogenes. Protein kinase B or Akt (PKB/Akt) is a serine/threonine kinase, it appears that Akt/PKB lies in the crossroads of multiple cellular signaling pathways and acts as a transducer of many functions initiated by growth factor receptors, and regulates various cell processes such as cell survival, cell cycle, metabolism, protein synthesis, apoptosis as well as transcriptional regulation (5,6).

Recent evidence indicates that PKB/Akt is frequently constitutively active in many types of human cancer, including gastric cancer (7,8). Constitutive PKB/Akt activation can occur due to amplification of PKB/Akt genes or as a result of mutations in components of the signaling pathway that activates PKB/Akt.

Although Akt overexpression has been detected in certain tumor types such as colon, ovarian and pancreatic carcinoma (9,10), the role of Akt in tumor biology has not been well studied in other tumor systems. Most of the studies are centered on the functional importance of overall Akt activity in various cell types without sufficient consideration of the roles of individual Akt isoforms, despite the obvious significance for specific therapeutic targeting. In the current study, we constructed the siRNA-Akt1 and transfected them into the gastric cancer cell line SGC7901 to investigate the influence of down-regulation of Akt1 on tumor biology.

## Materials and methods

**Tissue microarray/immunohistochemistry.** We constructed a tissue array consisting of 100 representative 1-mm cores from formalin-fixed, paraffin-embedded tissue blocks from each of 45 gastric adenocarcinoma (25 samples of poorly differentiated adenocarcinoma, 10 samples of moderately differentiated adenocarcinoma, 10 samples of well-differentiated adenocarcinoma), and the gastric mucosa surrounding carcinomas

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**Key words:** Akt1, small interference RNA, gastric cancer, cell proliferation

Table I. siRNA sequences targeting Akt1.

Name	Sequence
Nonsense siRNA	5'-AATTCTCCGAACGTGTCACGTTT-3'
Akt1 siRNA	
Sense	5'-GGAGAUCAUGCAGCAUCGC-3'
Anti-sense	5'-GCGAUGCUGCAUGAUCUCC-3'

as positive control, and 10 samples of normal gastric tissues as negative control. Immunohistochemistry steps were the same as described in the *in vivo* study.

*siRNA constructs.* Small interfering RNA specifically targeting Akt1 were constructed as described by Elbashir and colleagues (11), and chemically synthesized (Shanghai GenePharma Co., Ltd., China). The short RNA oligonucleotids used are listed in Table I.

*Cell culture and transfection.* Human gastric cancer SGC7901 cells were kindly provided by Dr Daiming Fan. Cells were cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% heat-inactivated fetal calf serum, incubated in 5% CO<sub>2</sub> at 37°C. For gene transfection, 2x10<sup>5</sup> cells per well were plated into six-well plates and grown overnight until they were 50-80% confluent. siRNA-Akt1 was transfected into SGC7901 cells by oligofectamine as instructed by the manufacturer.

*Western blot analysis.* After 24-48 h of transfection, cells were washed 3 times with ice-cold phosphate-buffered saline (PBS). Then the cells were lysed by RIPA (Solarbio Company, China) and PMSF protease inhibitor mixture. Homogenates were clarified by centrifugation at 12000 rpm for 15 min at 4°C and protein concentrations were determined by Nanodrop ND1000 spectrophotometer (Gene Company, USA). The same amount of protein samples were applied onto SDS-PAGE at 60 V for 1 h; separate proteins were transferred to PVDF membranes (Millipore, USA) at 80 V followed by blocking with 10% non-fat dry milk. The membranes were incubated with primary antibody against Akt1 (1:1000 dilution, Santa Cruz Biotechnology), followed by incubation with HRP-conjugated secondary antibody (1:1000 dilution, Zhongshan-jinqiao, China). The blots were then reacted with enhanced chemiluminescence Western blot detection system (Pierce, USA). After washing with stripping buffer, the membrane was re-probed with antibody against β-actin (1:500 dilution, Santa Cruz Biotechnology) using the same procedures described above.

*Real-time PCR.* The total RNA was extracted using Trizol reagents (Promega, USA), according to the manufacturer's instructions and quantified by Nano-drop ND-1000 (Nanodrop, USA). Isolated RNA was electrophoresed through 1% agarose-formaldehyde gels to verify the quality of the RNA. The first strand cDNA was generated by reverse transcription. The reaction system was: 5X RNA PCR buffer 4 μl; dNTP

Table II. The sequences of PCR primers (Takara Bio).

Name	Sequence
Akt1	
Forward	5'-GGCCCAGATCACCATCAC-3'
Reverse	5'-CTATCGTCCAGCGCAGTCCA-3'
β-actin	
Forward	5'-GCCGGGACCTGACTGACTA-3'
Reverse	5'-TGCGGATGTCCACGTCACT-3'

mixture (10 mM) 2 μl; RNase inhibitor 0.5 μl; MV reverse transcriptase 2 μl; Oligo-dT, adaptor primer 0.5 μl; total RNA from sample (≤1 μg) 1 μl; RNase Free dH<sub>2</sub>O up to 20 μl. After sufficient cDNA was obtained, we performed PCR amplification using a real-time PCR cyclor. The sequences of PCR primers (Takara Bio) are listed in Table II.

To determine the expression level of Akt1, we mixed 10 μl 2X PCR premix (Mg<sup>2+</sup>), 0.5 μl Primer-1, 0.5 μl Primer-2, 1 μl cDNA and 8 μl dH<sub>2</sub>O up to 20 μl reaction system (Takara Bio). The reactions were incubated in a 96-well optical plate at 94°C for 12 min, followed by 40 cycles of 94°C for 30 sec, 56°C for 30 sec, 72°C for 30 sec, 76°C for 2 sec and 72°C for 5 min. Melting curve was from 65°C to 95°C, stay for 2 sec and read the plate per 0.2°C. The expression of Akt1 was examined by normalization of the cycle threshold (Ct) of these genes to that of the control gene (β-actin),  $\Delta Ct = Ct_{Akt1} - Ct_{\beta-actin}$ .

*Immunofluorescence.* After 24-48 h of transfection, cells were washed 3 times with ice-cold phosphate-buffered saline (PBS), then fixed with 4% paraformaldehyde for 40 min, washed with PBS 3 times, then permeabilized with 1% Triton X-100 for 5 min and blocked in 1% BSA for 20 min at room temperature. Then incubated with primary antibody (1:100 dilution) overnight at 4°C, then fluorescent secondary antibody (1:100 dilution) was added at 37°C for 1 h, after washing with PBS, cells were incubated with Hoechst for 20 min and mounted. The results were observed with a confocal microscopy.

*Flow cytometry analysis.* Transfected and control cells in the log phase of growth were harvested by trypsinization. Cells were centrifuged for 5 min at 1000 rpm and washed with PBS, fixed with ethanol. Nuclei of cells were stained with propidium iodide (JingMei Company, Shanghai) for an additional 30 min. Total of 10,000 nuclei were examined in a FACS Calibur flow cytometer (Becton-Dickinson, Heidelberg, Germany) and DNA histograms were analyzed by Modifit software. Cells were stained with Annexin V and propidium iodide was performed as recommended by the manufacturer, and apoptosis was determined by flow cytometry.

*Subcutaneous tumor model and gene therapy.* Four to six-week-old female immune-deficient nude mice (BALB/c-nu) were purchased from the animal center of the Cancer Institute of Chinese Academy of Medical Sciences, bred at the facility of laboratory animals, Tianjin University, and housed in

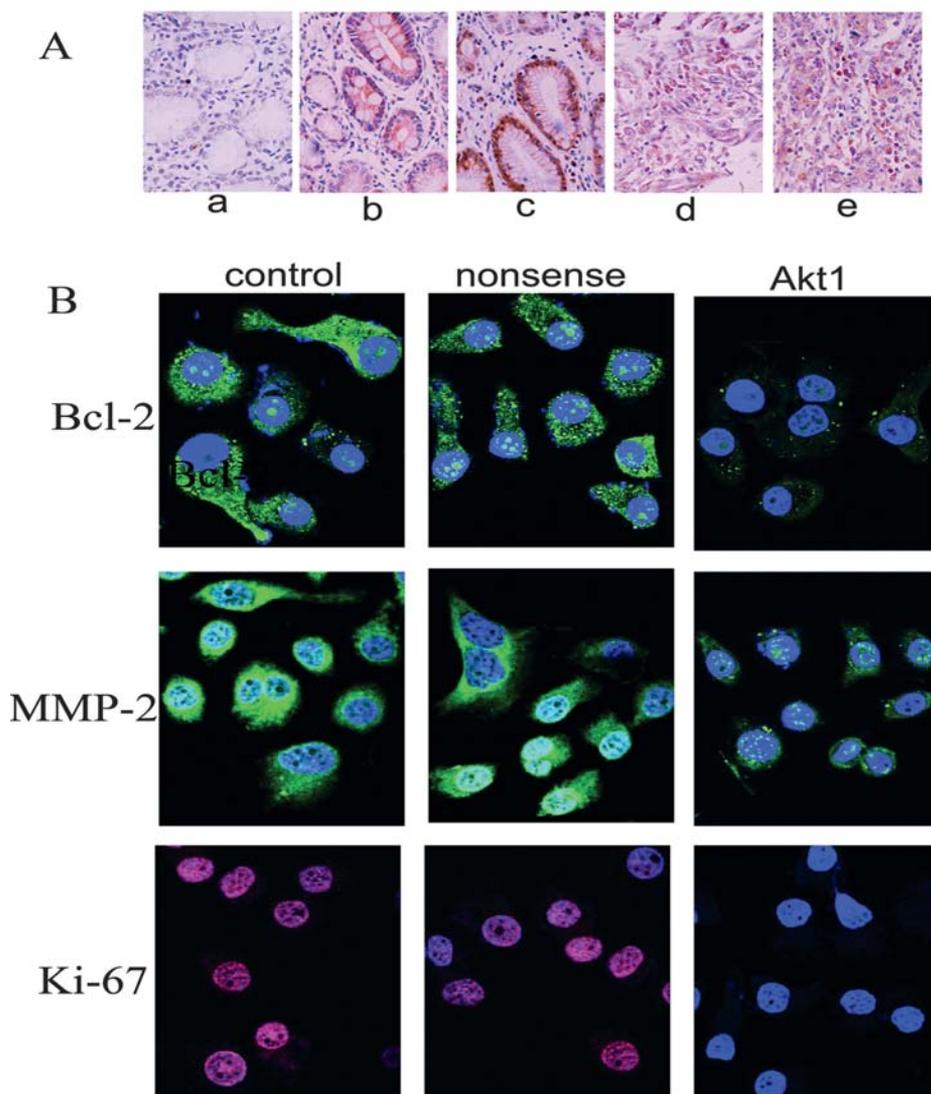


Figure 1. A tissue microarray consisting of biopsies from 45 gastric adenocarcinomas to analyse Akt1 protein expression was constructed (A). The results indicated that Akt1 expression in gastric cancer tissue significantly increased compare to normal tissue and adjacent non-cancer tissue ( $P < 0.05$ ), the difference was not significant ( $P > 0.05$ ) in well-differentiated, moderately differentiated or poorly differentiated gastric adenocarcinoma, but between well-differentiated and poorly differentiated gastric adenocarcinoma, the difference was significant ( $P < 0.05$ ). (a) Normal gastric mucosa, (b) adjacent non-cancer tissue, (c) well-differentiated gastric adenocarcinoma, (d) moderately-differentiated gastric adenocarcinoma, (e) poor-differentiated gastric adenocarcinoma (x200). (B) The results of immunofluorescence indicate that suppression of Akt1 expression by siRNA-Akt1 in SGC7901 cells induces down-regulation of Bcl-2, Ki-67 and MMP-2, however, in control and siRNA-nonsense groups, the expression of Bcl-2, Ki67 and MMP-2 present strong, positive immunoreactivity.

microisolator individually ventilated cages with water and food. All experimental procedures were carried out according to the regulations and internal biosafety and bioethics guidelines of Tianjin Medical University and the Tianjin Municipal Science and Technology Commission.

Five mice were injected subcutaneously with  $5 \times 10^7$ /ml of SGC7901 cells, in a volume of  $50 \mu\text{l}$  of DMEM (serum-free) pre-mixed. Mice were monitored daily. When the tumor size reached approximately 10 mm in length, the tumors were surgically removed, cut into pieces of 1-2 mm<sup>3</sup> and re-seeded in other 24 mice individually. When the tumor size reached approximately 5 mm in length, the mice were randomly grouped as control, siRNA-nonsense, and siRNA-Akt1;  $150 \mu\text{l}$  siRNA and  $60 \mu\text{l}$  oligofectamine mixed. In siRNA-nonsense and siRNA-Akt1 groups,  $25 \mu\text{l}$  of mixture were injected into subcutaneous tumors of each nude mouse in a multi-site

injection manner. Mice in control group received  $25 \mu\text{l}$  of PBS only. Another administration was conducted every four days in the same manner until the fourth week, the tumor volume was measured with slider caliper, using the formula: volume = length x width<sup>2</sup>/2.

**Immunohistochemistry.** After 28 days, mice were sacrificed and samples were fixed with 10% formalin, paraffin-embedded tissue sections were used for examination of Akt1, proliferating cell nuclear antigen (PCNA), matrix metalloproteinase-2 (MMP2) and Bcl-2 expression. Sections were dewaxed, treated with 3% H<sub>2</sub>O<sub>2</sub> for 7 min and incubated with primary antibody (1:100) overnight at 4°C. Biotinylated secondary antibody (1:100) was added at 37°C for 1 h, followed by the incubation with ABC-peroxidase for an additional 1 h. After washing with PBS, the sections were

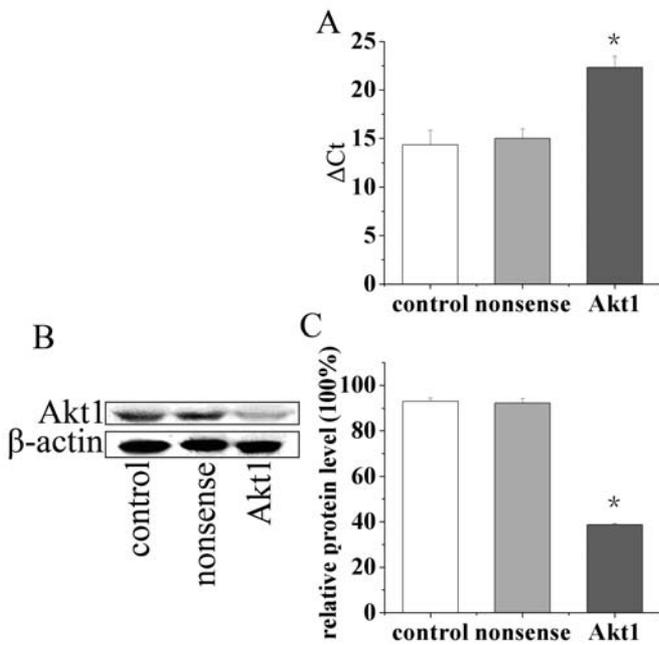


Figure 2. Suppression of Akt1 expression by siRNA-Akt1 in SGC7901 cells. (A) Real-time PCR of SGC7901 cells treated with siRNA-Akt1 and siRNA-nonsense. (B and C) Western blot analysis of Akt1 expression in SGC7901 cells transfected with siRNA-nonsense, siRNA-Akt1 and in the control group.

incubated with DAB (3,3' diaminobenzidine, 30 mg dissolved in 100 ml Tris-buffer containing 0.03%  $H_2O_2$ ) for 5 min, rinsed in water and counterstained with hematoxylin.

**Detection of apoptosis.** Apoptosis was detected by TUNEL staining. Briefly, sections were dewaxed, incubated with blocking solution (0.3%  $H_2O_2$  in double diluted water) for 30 min, and permeabilized with 0.1% Triton X-100 in PBS for another 2 min on ice. Apoptosis was detected using an *in situ* cell death kit (Boehringer Mannheim, Germany). Positive cells were visualized by confocal microscopy. The reaction mixture was incubated without enzyme in a control section to detect nonspecific staining.

**Statistical analysis.** A statistical package SPSS11.5 was used for statistical analysis. One-way analysis of variance (ANOVA) and  $\chi^2$  test was used to analyze the significance between groups. Statistical significance was determined at  $P < 0.05$  level.

## Results

**Tissue microarray.** We constructed a tissue microarray consisting of biopsies from 45 gastric adenocarcinoma and the gastric mucosa surrounding carcinomas, and 10 samples of normal gastric tissues. We analyzed Akt1 protein expression. The results indicated that in normal gastric mucosa, adjacent non-cancer tissue and gastric adenocarcinoma, the positive rate of Akt1 was 30.00, 66.72 and 71.11%, respectively, and the difference was significant ( $P < 0.05$ ); in well-differentiated, moderately differentiated and poorly differentiated gastric adenocarcinoma, the positive rate of Akt1 was 50, 60 and 84.4%, respectively, and the difference was not significant ( $P > 0.05$ ), but between well-differentiated and poorly differentiated gastric adenocarcinoma, the difference was significant ( $P < 0.05$ ) (Fig. 1A).

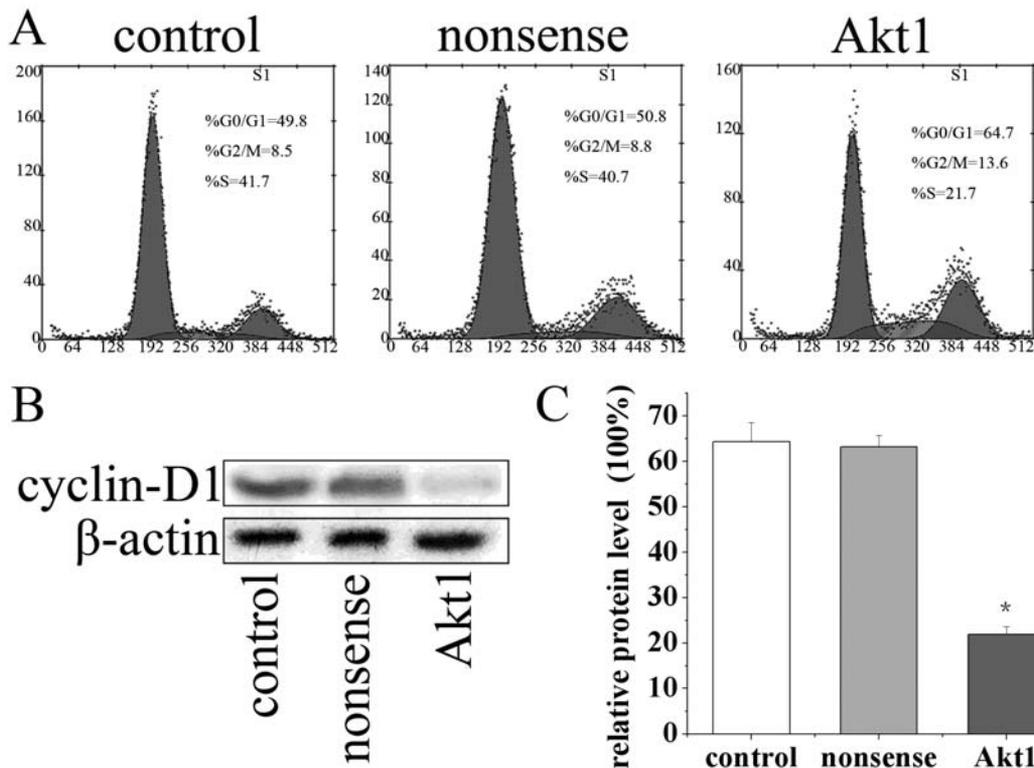


Figure 3. Suppression of Akt1 expression by siRNA-Akt1 in SGC7901 cells induces G0/G1 arrest and inhibits cell growth (A). Cell cycle distribution of SGC7901 cells after transfection detected with FCM. Transfected and control cells were harvested by trypsinization, washed and fixed overnight, nuclei of cells were stained with propidium iodide, a total of 10,000 nuclei was examined. The expression of cyclin-D1 after transfection with siRNA (B).

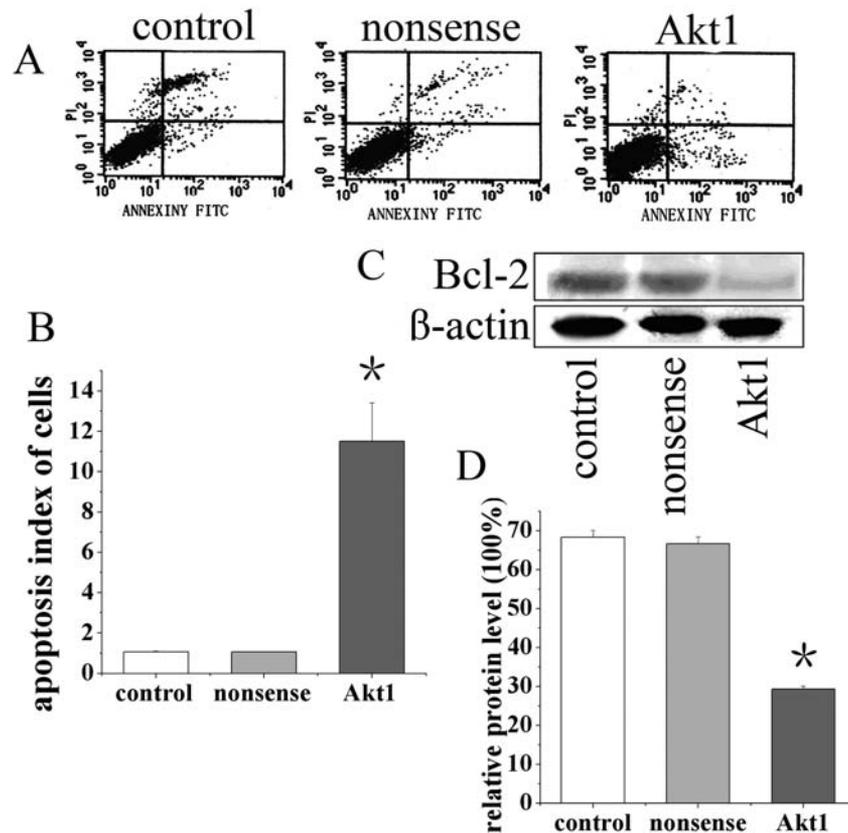


Figure 4. Apoptosis of SGC7901 cells transfected with siRNA-nonsense and siRNA-Akt1 evaluated by Annexin V staining detected with FCM (A and B); the expression of Bcl-2 after transfection with siRNA (C and D).

**siRNA-induced gene silencing in SGC7901 cells.** Real-time PCR showed that transfection of siRNA-nonsense had little effect on Akt1 expression. However, Akt1 expression was dramatically down-regulated in SGC7901 cells transfected with siRNA-Akt1 ( $P < 0.05$ ) (Fig. 2A). Meanwhile, Western blot analysis demonstrated similar down-regulation of Akt1 expression consistent with results from real-time PCR (Fig. 2B and C). These results suggest that siRNA-Akt1 can potently and specifically inhibit the Akt1 expression in SGC7901 cells.

**Immunofluorescence.** The expression of Bcl-2, Ki-67 and MMP2 was examined to determine the role of Akt1 in apoptosis, proliferation and invasion in SGC7901 cells. The results indicated the strong, positive immunoreactivity of Bcl-2, Ki-67 and MMP2 in control groups and siRNA-nonsense groups. However, the expression of Bcl-2, Ki-67 and MMP2 was down-regulated in the Akt1-siRNA groups (Fig. 1B).

**Effect of the Akt1 knock-down on SGC7901 cell proliferation.** The cell cycle distribution of control and transfected cells was analyzed by flow cytometry. As shown in Fig. 3A, the G2/M phase fraction of parental SGC7901 cells and cells transfected with siRNA-nonsense and siRNA-Akt1 was 8.5, 8.8 and 13.6%, respectively. The S phase fraction in parental SGC7901 cells and cells transfected with siRNA-nonsense and siRNA-Akt1 were 41.7, 40.4 and 21.7%, respectively, and decreased significantly in siRNA-Akt1 transfected cells. The

G0/G1 phase fraction in parental SGC7901 cells and in cells transfected with siRNA-nonsense and siRNA-Akt1 were 49.8, 50.8 and 64.7%, respectively. These results suggest that siRNA-Akt1 approach can induce the arrest of cells at G0/G1 phases, delay the progression of cell cycle and inhibit the cell proliferation, and the expression of cyclin-D1 decreased significantly in cells transfected with siRNA-Akt1 (Fig. 3B and C). As shown in Fig. 4A and B, apoptosis of SGC7901 cells treated with siRNA-Akt1 increased significantly compared with siRNA-nonsense and control groups ( $P < 0.05$ ), and the expression of Bcl-2 decreased in cells treated with siRNA-Akt1 (Fig. 4C and D), suggesting that down-regulation of Akt1 may be involved in the regulation of cell apoptosis.

**Cell invasion assessed by MMP-2 and MMP-9.** To further evaluate the anti-invasive activity of SGC7901 cells by siRNA-Akt1 approach, we assessed the inhibitory effect on cell invasion through detecting the expression of MMP-2 and MMP-9. Western blot analysis showed that the expression of MMP-2 (Fig. 5A and B) and MMP-9 (Fig. 5C and D) decreased in cells treated with siRNA-Akt1 ( $P < 0.05$ ). The result suggests that the knock-down of human Akt1 expression by siRNA approach can significantly reduce gastric cancer cell invasion.

**Anti-tumor effect of the Akt1 knock-down in SGC7901 gastric cancer xenograft model.** Our *in vitro* experiments demonstrated that siRNA-Akt1 approach can efficiently inhibit cell

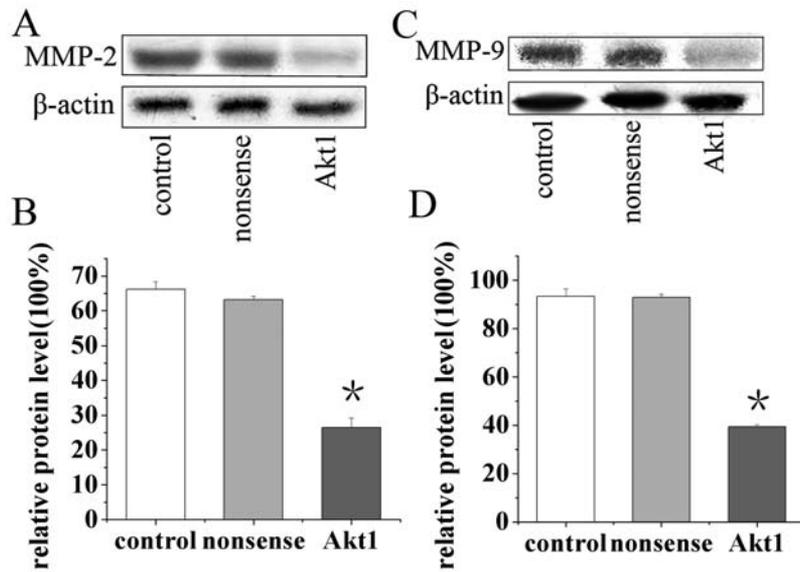


Figure 5. Cell invasive capability of SGC7901 cells. The expression of MMP-2 (A and B) and MMP-9 (C and D) after transfection with siRNA.

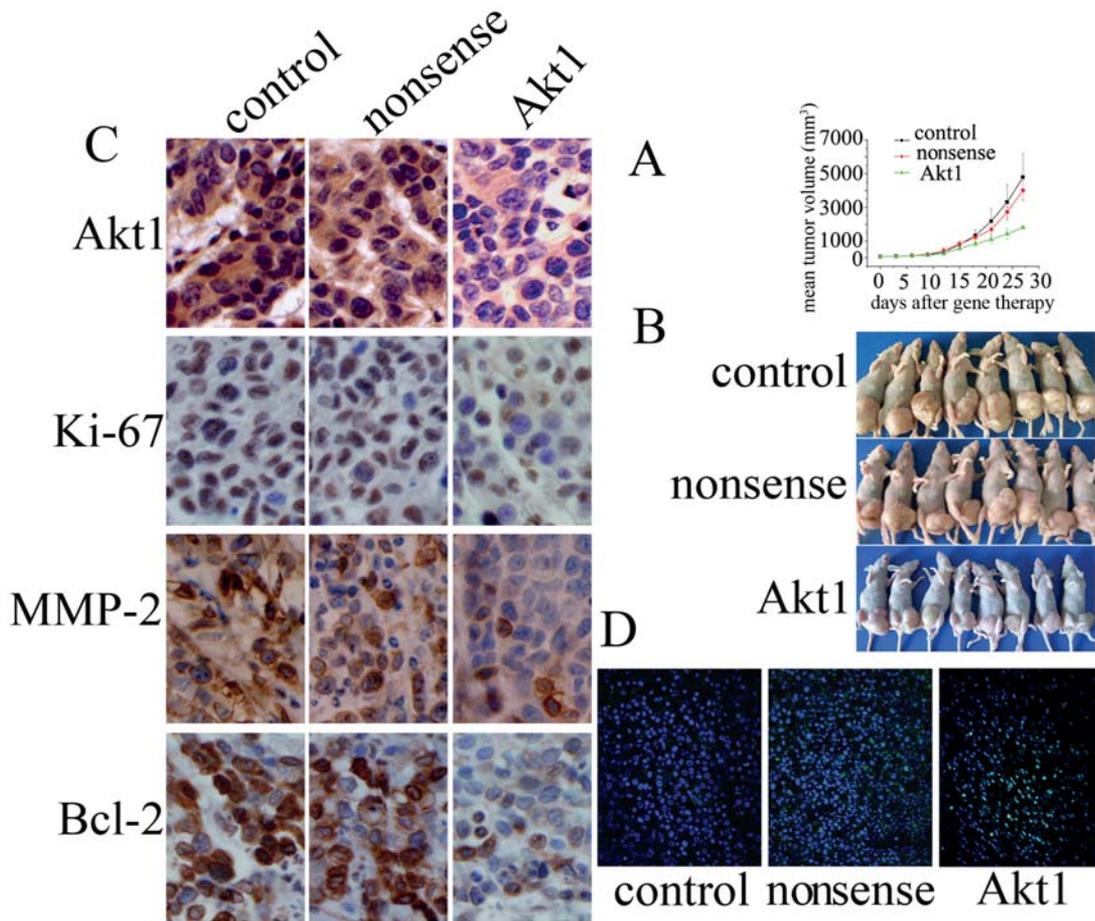


Figure 6. Compared to the control and siRNA-nonsense groups, tumor growth in nude mice in the group of siRNA-Akt1 was inhibited (A and B); compared to the control and siRNA-nonsense groups, Akt1, MMP2, Bcl-2 and Ki-67 were down-regulated (C); apoptotic cells were obvious by TUNEL staining (D).

proliferation, induce the G0/G1 cell cycle arrest, cell apoptosis and prevent the invasion of SGC7901 cells. Therefore, we further investigated the anti-tumor effect of such approaches in *in vivo* using SGC7901 gastric cancer xenograft model and oligofectamine-mediated gene therapy as indicated in Materials and methods. Tumors were established

in the right leg of 24 mice by SGC7901 xenografts, after reaching approximately 5 mm in length, and each mouse was challenged by *in situ* injection of PBS as control (n=8) or oligofectamine-mediated gene therapy for siRNA-nonsense (n=8), siRNA-Akt1 (n=8), respectively. The mean tumor volume of each group at the time of gene therapy was

64.80±15.00 mm<sup>3</sup>, 64.65±15.08 mm<sup>3</sup>, 63.65±13.22 mm<sup>3</sup>, respectively. No statistically significant difference was noted among the groups (P=0.985). The mice were monitored every three days for 4 weeks, and the tumor volumes were measured and compared. The mean volumes of tumors were 4789.563±1367.747 mm<sup>3</sup>, 4003.975±544.457 mm<sup>3</sup> and 1797.350±342.161 mm<sup>3</sup> after treatment with PBS oligofectamine-mediated gene therapy for control, siRNA-nonsense and siRNA-Akt1, respectively. The Akt1 knock-down resulted in statistically significant reduction in size of the tumor compared to control and siRNA-nonsense group (Fig. 6A and B).

After mice were observed for 28 days, the tumors were removed and paraffin-embedded section were prepared for immunohistochemical examination using anti-Akt1, MMP2, Bcl-2 and Ki-67 antibodies. Representative sections from mice treated with PBS or oligofectamine-mediated gene therapy for siRNA-Akt1, siRNA-nonsense are shown in Fig. 6A. In control and siRNA-nonsense treated groups, strongly positive immunoreactivity of Akt1, MMP2, Bcl-2 and PCNA were observed (Fig. 6C). TUNEL staining showed that there were nearly no apoptotic cells found in PBS and siRNA-nonsense treated groups. However, apoptosis was prominently increased in siRNA-Akt1-treated group (Fig. 6D).

## Discussion

*Gene therapy targeting Akt1 in gastric cancer therapy.* Akt was discovered as a cellular homologue (c-Akt) of the viral oncogene (v-Akt) from the acutely transforming retrovirus Akt8, isolated from a murine T-cell lymphoma (12). At the same time, it was identified as a novel kinase similar in many respects to protein kinase A (PKA) and protein kinase C (PKC) and hence is also referred to as protein kinase B (PKB) (13). Akt gene encodes a 56 kDa of serine/threonine protein kinase. In mammals, there are three isoforms of Akt (Akt1, 2 and 3, or PKB $\alpha$ ;  $\beta$  and  $\gamma$ ), which have a broad tissue distribution. The expression patterns of the three Akt isoforms differ, as do their apparent biological activities. Akt2 appears to be expressed at greater levels in insulin responsive tissues. Consistent with these data, Akt2 knockout mice develop mild type 2 diabetes, while Akt1 knockout mice are non-diabetic, but small in size (14). Akt3 expression is more limited, occurring in brain, heart, kidney, and in certain poorly differentiated cancers. All three isoforms are composed of a central catalytic domain, a C-terminal hydrophobic domain, and an N-terminal pleckstrin homology domain (PH domain), making it a target for phospholipids produced by PI3 kinase activity. Akt activity is induced by ligand stimulation of growth factor receptors, such as the insulin-like growth factor-I receptor (IGF-IR) and the EGF family of receptors. It appears that Akt/PKB lies in the crossroads of multiple cellular signaling pathways and acts as a transducer of many functions such as regulating cell growth, proliferation, survival, cell motility, anti-apoptosis, metabolism and other activities. Therefore, dysregulation of Akt-mediated signaling pathways may play an important role in tumor development and progression.

Although *in vitro* or *in vivo* studies on several types of human cancer have identified the role of activated Akt, its

role in the development and malignancy of gastric carcinoma *in vivo* and *in vitro* remains unclear. A previous study reported that Akt was expressed in 74% of the gastric cancers, and this was not associated with any other clinicopathological factors, and their study also indicated that activated Akt expression was highly detected in early-stage pTNM. This observation is compatible with a study that the overexpression and phosphorylation of Akt is an early event during sporadic colon carcinogenesis (15). Akt activation appears to be required for cell growth and proliferation in early-stage human gastric carcinoma. Oki *et al* reported that the activated Akt was associated with increased resistance to multiple chemotherapeutic agents (5-fluorouracil, adriamycin, mitomycin C and cisplatin) in gastric cancer patients (8). Therefore, down-regulation of the Akt activity seems to be a promising target for gastric cancer therapy. A previous study indicated that small interfering RNA (siRNA) directed against the Akt1 can efficiently suppress its expression in human gastric cancer cell lines leading to the induction of arrest at G1 by inhibiting p27 nuclear import (16). Collectively, the increased expression of Akt1 and the key role that it may play in gastric cancer development and progression led to the investigation of targeting Akt1 for anticancer therapies.

*Biological changes by silenced Akt1 expression.* Akt/PKB is elevated in most gastric cancers (17). One of the earliest studies that identified the two human orthologues, Akt1 and Akt2, based on the homology to the viral oncogene v-Akt, showed a 20-fold amplification of the Akt1 locus in gastric adenocarcinoma (18). It has been reported that in gastric cancer AGS cells transfected with Akt1 siRNA, the expression of Akt1 was dramatically decreased, and the down-regulation of Akt1 significantly down-regulated the expression of Bcl-2, and up-regulated the expression of Bax, but did not alter the expression of PTEN in gastric cancer cells. Our results also indicated down-regulation of Akt1 down-regulated the expression of Bcl-2 and increased the apoptosis of tumor cells. Fan *et al* showed that overexpression of constitutively active Akt protects cells from COX-2 inhibitor SC236-induced cell death and up-regulates the expression of several survival proteins. These results indicate that inhibition of the Akt activity contributes to SC236-induced apoptosis in gastric cancer (19). Akt1 siRNA could significantly enhance the sensitivity of AGS cells to vincristine, adriamycin, 5-fluorouracil and cisplatin (20). Another study reported that inhibition of Akt1 is sufficient to affect cell migration, invasion, and proliferation. Expression of Akt1 siRNA had a similar effect as p110 $\alpha$  siRNA in the cells (21). Previous studies indicated that deficiency of Akt1 results in impaired adhesion and migration of endothelial cells to various matrix proteins including fibronectin, vitronectin and fibrinogen, and showed that Akt1<sup>-/-</sup> endothelial cells adhere to fibronectin, vitronectin and fibrinogen with lower affinity compared to WT (22). It was also suggested that Akt1 activation can accelerate the early stages of ErbB-2 mediated tumorigenesis through increased cyclin D1 expression and cellular proliferation (23). A recent study discovered that phospho-Akt1 was overexpressed in human gastric cancers and its levels correlated with tumor differentiation and pTNM. Akt1 activation promoted cell survival because the phosphatidyl-

inositol 3-kinase (PI3K) inhibitor LY294002 inhibited Akt1 phosphorylation and inhibited cell growth, especially in cells with active Akt1. Dominant negative Akt inhibited proliferation of gastric cancer cells and induced G1 cell-cycle arrest whereas constitutively active Akt increased cell proliferation (24). In the present study, we also found that the expression of Akt1 was decreased via siRNA-Akt1, and biological ability of proliferation, apoptosis and invasion of SGC7901 cells were suppressed. Thus, signal transduction pathways downstream of Akt were found to be critical in mediating important cellular functions, including survival, proliferation, migration, and invasion. Therefore, inhibition of overexpressed and/or activated PI3-K/Akt mediated signaling pathways possess great significance in gastric cancer gene therapy.

*Use of siRNA gene therapy for gastric cancer.* Gene silencing of target mRNA by RNA interference (RNAi) was first discovered in *Caenorhabditis elegans* as a response to double-stranded RNA (dsRNA), which resulted in sequence specific gene silencing and has dramatically expanded the arsenal of genetic tools that can be used to study signaling pathways in mammalian systems (25). The triggers for RNAi are small double-stranded RNAs of 21-23 nucleotides in length called small interfering RNAs or siRNA; one of the two strands forms a complex with a set of proteins (the RNA induced silencing complex, or RISC), with the RNA serving as a sequence-specific guide element for identification of the target RNA (26). The mechanism of RNAi was reported in two steps. The first step, referred to as the RNAi initiating step, involves binding of the RNA nucleases to a large dsRNA and the activation of a family of RNase III ribonucleases termed Dicer enzymes, which initiates the cleavage of the dsRNA into ~22 nt double-stranded duplexes with 2 nt 3'-overhangs and 5' phosphate termini, termed the small interfering RNA (siRNA). In the second step, the siRNA is utilized by RISC, which uses the siRNA as a template to recognize and cleave RNA targets with similar nucleotide sequences (27-32). The specific mediator of RNAi is the short dsRNA.

Application of RNAi in mammalian cells demonstrated a particular challenge since introduction of exogenous long dsRNA activates an innate immune (IFN) response, which leads to the inhibition of protein translation by the PKR pathway and activation of RNase L (33). However, transfection of target-specific synthetic short dsRNA (21-23 nt in length) into mammalian cells yielded gene silencing capabilities, which allowed for the routine application of siRNA in mammalian cells (34). Current siRNA selection criteria are based on guidelines first published by Elbashir *et al.* (34). These criteria and subsequent revisions were the result of trial-and-error observations of randomly selected siRNA. Briefly, the requirements include: 21 nt sense and antisense strands paired as to have a 2 nt 3'-overhang, target regions starting 50-100 nt downstream of a start codon, 50% G/C content. The guidelines provide a starting point to design siRNA, but provide little specificity to ensure siRNA knock-down efficacy.

The therapeutic application of siRNA, however, is largely dependent on the development of a delivery vehicle that can efficiently deliver the siRNA to target cells. DNA vector-

mediated RNAi technology has made it possible to develop therapeutic applications of this technology in mammalian cells (35). Several examples using retroviral (36-39) or adenoviral vector systems (40) to deliver siRNA for stable or transient expression, respectively, have been reported. In the present study, the expression of Akt1 was decreased using siRNA-Akt1, sequentially, the biology of SGC7901 cells including proliferation, invasion and apoptosis changed. More studies are needed to test silenced efficacy of various target genes of Akt/PKB pathways, which may become a therapeutic target of gastric cancer.

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