

Blockade of sonic hedgehog signaling decreases viability and induces apoptosis in retinoblastoma cells: The key role of the PI3K/Akt pathway

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Abstract. Retinoblastoma (RB) is the most common type of malignant intraocular cancer in teenagers. One of the proteins abnormally expressed during oncogenesis of RB is sonic hedgehog (SHH), which possesses the capability to selectively activate transcription factors of different genes. However, the detailed function of SHH in RB remains unknown. Thus, the present study sought to investigate the role of SHH in the development of RB. The human RB WERI-Rb-1 cell line was used as an *in vitro* model for the knockdown of *SHH* by a specific short hairpin RNA (shRNA). To assess the effect of SHH inhibition on cell growth and apoptosis, cell viability, colony formation and flow cytometry assays were conducted. WERI-Rb-1 cells transfected with an shRNA targeting *SHH* were treated with the phosphoinositide-3 kinase (PI3K)/Akt agonist insulin-like growth factor 1 (IGF-1) to investigate the possible mechanism by which SHH promotes RB. The present results revealed that the silencing of *SHH* induced G1 cell-cycle arrest and apoptosis in WERI-Rb-1 cells and led to a decrease in cell viability, indicating that SHH has a critical role in the determination of RB cell survival. Moreover, according to the results of the IGF-1 assays, suppression of PI3K/Akt was a prerequisite for SHH inhibition, illuminating its potential role in the treatment of RB. The findings outlined in the present study elucidate a clear link between SHH and the PI3K/Akt pathway in RB cell survival, which could provide valuable inspiration for the advancement of therapies against RB.

Introduction

Retinoblastoma (RB) is the most common type of malignant intraocular cancer in children <5 years old (1). In the majority of

cases, particularly in developing countries, RB is only detected at the late stages of the cancer, which severely threatens the lives of the patients (2). Prior to the 1990s, surgical removal of the afflicted eyes, followed by external-beam radiotherapy, was the major treatment method for RB; however, the approach was associated with high morbidity and craniofacial deformity (3). At present, the most effective therapy against RB is chemotherapy combined with immunotherapy, which is cytotoxic to RB cells, increasing cell apoptosis (2). Despite its high efficacy, the current therapy remains somewhat unsatisfactory, owing to the adverse effects of chemotherapy. Thus, the development of mild therapeutic modalities to improve the outcome and survival of RB is required.

Numerous signaling molecules are relevant to the oncogenesis and development of cancer. One of these signaling transduction pathways is the hedgehog (HH) pathway, which governs processes such as embryonic development, cell differentiation, cell proliferation and tissue patterning (4). The most studied member of the vertebrate HH family is sonic hedgehog (SHH). SHH exerts its function by selectively activating the transcription factors of different genes. Altered SHH pathway activation has been observed in different types of solid and non-solid cancer (5). The selective regulation of SHH production can influence cell apoptosis and the metastasis of pancreatic and ovarian cancer cells through multiple mechanisms (6-8). These findings reveal the close connection between the SHH pathway and other critical pathways, which include phosphoinositide-3 kinase (PI3K)/Akt and nuclear factor κ -light-chain-enhancer of activated B cells pathways. Considering that tumorigenesis initiates aberrant apoptotic processes, the aforementioned data suggest that the SHH pathway may exert control over apoptosis in cancer cells. Moreover, a recent study regarding the SHH pathway has validated the critical role of the SHH pathway in the progression of RB (9). From a therapeutic point of view, it is therefore desirable to further evaluate how regulation of the SHH pathway affects RB cell survival.

In the current study, the expression of *SHH* in the human RB WERI-Rb-1 cell line was regulated using specific short hairpin RNAs (shRNAs). Cell viability and apoptosis were measured to investigate the function of SHH in the survival of RB cells. As PI3K/Akt serves such a central role in cell

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apoptosis, further experiments subjecting WERI-Rb-1 cells to insulin-like growth factor 1 (IGF-1) were conducted to assess the possible mechanism by which SHH drives the promotion of RB. IGF-1 is a pleiotropic polypeptide with a wide range of actions in the central and peripheral nervous systems, which protects neurons against cell death induced by amyloidogenic derivatives, glucose or serum deprivation via the activation of intracellular pathways implicating PI3K/Akt (10). Inhibition of SHH significantly increased the apoptotic rate in the WERI-Rb-1 cells and resulted in a decrease in cell viability. Downregulation of PI3K/Akt is therefore required for SHH inhibition-based anti-RB therapies. The findings in the present study demonstrate a clear interaction between SHH and the PI3K/Akt pathway in the survival of RB cells and could aid the development of therapies for the prevention of this malignancy.

Materials and methods

Chemicals and cell culture. Antibodies against SHH and phosphorylated (p)-PI3K (cat. nos. bs-1544R and bs-5538R, respectively) were purchased from Beijing Biosynthesis Biotechnology Co., Ltd. (Beijing, China). Antibodies against cleaved caspase-3 and cleaved poly (ADP-ribose) polymerase (PARP) (cat. nos. ab2302 and ab32561, respectively) were purchased from Abcam (Cambridge, MA, USA). Antibodies against B-cell CLL/lymphoma 2 (Bcl-2), Bcl-2 associated X (Bax) and PI3K (cat. nos. BA0412, BA0315 and BA1352, respectively) were purchased from Boster Systems, Inc. (Pleasanton, CA, USA). Antibodies against p-Akt (Ser 473), Akt and β -actin (cat. nos. sc-8312, sc-135651 and sc-47778, respectively) were purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). The secondary IgG antibodies conjugated to horseradish peroxidase (cat. nos. WLA023 and WLA024) were purchased from Wanleibio Co., Ltd. (Shanghai, China). The PI3K/Akt agonist IGF-1 (cat. no. 10598-R023) was obtained from Sino Biological Inc. (Beijing, China). Transfection agents (cat. no. c1507) were purchased from Applygen Technologies Inc. (Beijing, China). A specific *SHH*-targeted shRNA (5'-CCCGACATCATATTTAAGGAT-3') and control shRNA (5'-GCTGTTGGACAGCGAGACCAT-3') were constructed by Wanlei Life Sciences (Shenyang, China) in accordance with a previous study (11). The human RB WERI-Rb-1 cell line was obtained from Wanlei Life Sciences and cultured in RPMI-1640 medium (cat. no. 31800-014; Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) supplemented with 20% fetal bovine serum (FBS; cat. no. SH30084.03; HyClone; GE Healthcare Life Sciences, Logan, UT, USA) and a 1% (v/v) antibiotics mixture (penicillin/streptomycin; cat. no. E485-20ML; Amresco, LLC, Solon, OH, USA) in an atmosphere of 95% air and 5% CO₂ at 37°C.

Knockdown of *SHH* gene in WERI-Rb-1 cells. Sequences of different shRNAs (1.2 μ g/100 μ l) were ligated into pRNAH1.1 plasmid (Sangon Biotech, Co., Ltd., Shanghai, China) to form transfection vectors, and transfection was conducted using HiGene transfection reagent (cat. no. C1507) from Applygen Technologies Inc., according to the manufacturer's protocol. Cells were divided into three groups: A control group of untransfected WERI-Rb-1 cells; an shNC group of WERI-Rb-1

cells transfected with a pRNAH1.1-NC (non-targeted shRNA) plasmid; and an shSHH group of WERI-Rb-1 cells transfected with a pRNAH1.1-SHH (*SHH*-targeted shRNA) plasmid. Each treatment was performed in triplicate. Stable transfected cells were screened in RPMI-1640 medium in the presence of G418 (0.5 μ g/ μ l) for subsequent experiments. The knockdown efficiency of *SHH* by shRNA was determined using reverse transcription-quantitative PCR (RT-qPCR) and western blot analysis.

RT-qPCR. For the detection of mRNA, whole RNAs in cells under different conditions were extracted using an RNA extraction kit according to the manufacturer's instructions (cat. no. RP1201; BioTeke Corporation, Beijing, China). β -actin was selected as the internal reference gene. cDNA templates of *SHH* were attained by reverse transcribing the RNA using an RT-PCR kit (cat. no. PR6502; BioTeke Corporation), and the final RT-qPCR reaction mixture of volume 20 μ l contained 10 μ l SYBR Green master mix, 0.5 μ l of each primer (10 μ M) (*SHH*, forward, 5'-AGCAACAGCAACGCAAAG-3' and reverse, 5'-AATAGCCAGGAGAGGAGGA-3'; and β -actin forward, 5'-TTAGTTGCGTTACACCCTTTCTTG-3' and reverse, 5'-CTGTACCTTCACCGTTCCAGTTT-3'), 1 μ l of the cDNA template and 8 μ l RNase-free H₂O. Thermal cycling parameters for the amplification were as follows: A denaturation step at 95°C for 10 min, followed by 40 cycles of 95°C for 10 sec, 60°C for 20 sec and 72°C for 30 sec. The relative expression level of the targeted gene was calculated with Data Assist Software version 3.0 (Applied Biosystems; Thermo Fisher Scientific, Inc.) according to the 2^{- $\Delta\Delta$ C_q} method (12). Each assay was repeated \geq 3 times.

Western blotting assay. The whole protein product in each group was extracted using Whole Protein Extraction kit according to the manufacturer's instructions (Wanleibio Co., Ltd.). β -actin was used as an internal reference protein. Protein concentration was determined using a bicinchoninic acid assay. Next, 40 μ g protein was subject to SDS-PAGE with a 10% gel. Following the transfer of targeted proteins onto polyvinylidene difluoride sheets, the membranes were washed with Tris-buffered saline plus Tween 20 (TTBS; cat. no. WLA025; Wanleibio Co., Ltd.) for 5 min and then blocked with skimmed milk powder solution for 1 h. Primary antibodies against SHH (1:500), cleaved caspase-3 (1:1,000), cleaved-PARP (1:1,000), Bcl-2 (1:400), Bax (1:400), p-PI3K (1:500), PI3K (1:400), p-Akt (1:200), Akt (1:200) or β -actin (1:1,000) were added and the membranes were incubated at 4°C overnight. After an additional four washes with TTBS, secondary IgG antibodies conjugated to horseradish peroxidase (1:5,000) were incubated with the membranes for 45 min at 37°C. Following a further six washes with TTBS, the blots were developed using Beyo ECL Plus reagent (Beyotime Institute of Biotechnology, Haimen, China) and the results were recorded in the Gel Imaging System (cat. no. WD-9413B; Beijing Liuyi Instrument Factory, Beijing, China). The relative expression levels of SHH in different groups were calculated with the Gel-Pro-Analyzer (Media Cybernetics, Inc., Rockville, MD, USA).

3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) test. The viability of different groups of WERI-Rb-1

cells was measured using an MTT assay. A total of 50 μ l exponentially growing cells (4×10^3 cells/ml) were seeded into a 96-well plate and cultured at 37°C for 120 h. For each group, the cell viabilities at 24, 48, 72, 96 and 120 h were all determined and each time point was represented by five replicates. In brief, 5 mg/ml MTT was added to each well. After incubating for an additional 4 h at 37°C, 200 μ l dimethyl sulfoxide was added and the cell viabilities of different groups were measured with a microplate reader at 490 nm.

Flow cytometry. WERI-Rb-1 cells in different groups were cultured in RPMI-1640 medium supplemented with 10% FBS at 37°C and collected by centrifugation at 626 x g for 5 min at 37°C. A flow cytometry assay was conducted to determine the effect of *SHH*-knockdown on the cell-cycle distribution of WERI-Rb-1 cells. In brief, cells were fixed with 70% ethanol at 4°C for 2 h. A total of 500 μ l propidium iodide (PI)-fluorescein isothiocyanate (FITC) (2 μ g/ml) was then added to different samples to stain DNA in the dark at 4°C for 30 min. After a 20-min incubation at room temperature, the DNA contents of the cells were analyzed using a flow cytometer (Accuri C6; BD Biosciences, Franklin Lakes, NJ, USA).

Colony formation in soft agar. The capability of cancer cells for anchorage-independent growth was measured by colony formation assay. Cells that had been transfected with different plasmids were seeded into RPMI-1640 medium containing 3% methylcellulose at a concentration of 500 cells per plate. After 3 weeks in culture, the numbers of colonies were counted using an inverted phase contrast microscope (cat. no. AE31; Motic Incorporation, Ltd., Causeway Bay, Hong Kong). Colony formation rate = colony number/inoculation cell number x 100.

Determination of the effect of *SHH*-knockdown on the apoptotic process in WERI-Rb-1 cells. To determine the effect of *SHH*-knockdown on the apoptotic process of WERI-Rb-1 cells, cells were first probed with an Annexin V-FITC Apoptosis Detection kit (Wanleibio Co., Ltd.) to determine the apoptotic rates according to the manufacturer's protocol. In brief, 5 μ l annexin V was added to different wells. Following incubation with annexin V for 10 min at room temperature, the cells were resuspended with 1X binding buffer (Annexin V-FITC Apoptosis Detection kit) and 5 μ l PI was added. Next, the apoptotic rates were analyzed using FACScan flow cytometry (Accuri C6; BD Biosciences, Franklin Lakes, NJ, USA). The apoptotic cell rate [upper right quadrant advanced-stage apoptosis cell percentage (UR) + lower right quadrant prophase apoptosis cell percentage (LR)-whole apoptosis cell percentage] was equal to the sum of the late apoptotic rate (UR) and the early apoptotic rate (LR). Additionally, the morphological changes of the nuclei in different groups were detected using a Hoechst staining kit, according to the manufacturer's instructions (Beyotime Institute of Biotechnology) and the results were recorded using a fluorescence microscope (IX53; Olympus Corporation, Tokyo, Japan) under x400 magnification. Cells with nuclei stained bright blue were judged to be Hoechst-positive and undergoing apoptosis. The expression of apoptosis-associated molecules, including cleaved caspase-3, cleaved PARP, Bcl-2, Bax, p-PI3K, PI3K, p-Akt and Akt, was determined by western blot analysis, as aforementioned.

Determination of effect of *SHH*-knockdown on the PI3K/Akt pathway in WERI-Rb-1 cells. To explain the mechanism driving the function of *SHH* in RB, WERI-Rb-1 cells in shNC and shSHH groups were administered with 1 μ g/10 μ l of the PI3K/Akt agonist IGF-1 for 48 h. In total, four groups were set up: The NC group, with shNC-transfected WERI-Rb-1 cells; the shSHH group, with shSHH-transfected WERI-Rb-1 cells; the NC+IGF-1 group, with shNC-transfected cells treated with IGF-1; and the shSHH+IGF-1 group, with shSHH-transfected cells treated with IGF-1. Following incubation, the cell viability, apoptotic rates and expression of p-PI3K, PI3K, p-Akt, Akt, cleaved caspase-3, cleaved PARP, Bcl-2 and Bax were determined as aforementioned.

Statistical analysis. All statistical analysis and graph manipulation was conducted using R version 3.2.1 (<https://cran.r-project.org/doc/FAQ/R-FAQ.html#Citing-R>). Data are expressed as the mean \pm standard deviation. Differences between multiple groups were determined based on the least-significant difference method, with $P < 0.05$ considered to indicate a significant difference.

Results

Transfection of specific shRNA reduces the production of *SHH* at mRNA and protein levels. Transfection of WERI-Rb-1 cells with *SHH*-specific shRNA markedly attenuated the expression of *SHH* at the mRNA and protein levels, and the difference between the shSHH group and the control or shNC groups was statistically different (Fig. 1; both $P < 0.05$).

Knockdown of *SHH* gene decreases cell viability and anchorage independent growth ability of WERI-Rb-1 cells. The cell viability of the different groups were measured using an MTT assay. The transfection of *SHH*-specific shRNA reduced the viability of the RB cells (Fig. 2A). The negative effect of shRNA transfection on cell viability could be observed from 48 h after incubation, and the difference between the shSHH group and the control or shNC groups was statistically different at the last four sampling time points (all $P < 0.05$). Additionally, the anchorage-independent growth ability of WERI-Rb-1 cells was markedly decreased by knockdown of *SHH* (Fig. 2B). The colony formation rate of cells in the shSHH group (20.7 \pm 2.2%) was almost half of those in the control (44.2 \pm 4.7%) and shNC (41.1 \pm 4.4%) groups, and the difference was statistically significant (both $P < 0.05$).

Knockdown of *SHH* gene induces G1 cell cycle arrest and apoptosis in WERI-Rb-1 cells. The cell-cycle distribution and apoptosis of cells under different conditions were analyzed with flow cytometry. The cell-cycle distribution analysis demonstrated a G1 cell-cycle arrest in WERI-Rb-1 cells administered with *SHH*-specific shRNA (Fig. 3A). The percentage of cells arrested in the G1 phase in the shSHH group (75.4 \pm 5.0%) was markedly higher than those in control (58.9 \pm 5.8%) and shNC (58.3 \pm 5.6%) groups. The downregulation of *SHH* also concomitantly decreased the fraction of cells in the S phase, representing a halt in cell proliferation through G1-phase cell-cycle arrest. Consistent with the results of cell-cycle distribution, the cell apoptotic rate in the shSHH

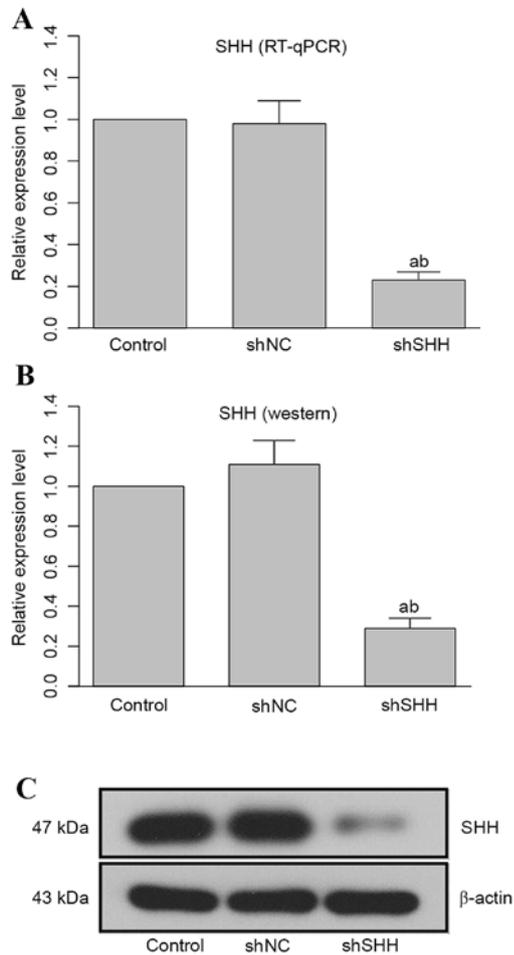


Figure 1. Validation of *SHH*-knockdown efficiency by RT-qPCR and western blotting assay. (A) Quantitative analysis result of RT-qPCR validation of *SHH*-knockdown efficiency. (B) Quantitative analysis result showing *SHH*-knockdown efficiency. (C) Representative image of western blot analysis, confirming the knockdown of *SHH*; β -actin was used as a control. ^a $P < 0.05$ vs. control group; ^b $P < 0.05$ vs. shNC group. SHH, sonic hedgehog; RT-qPCR, reverse transcription-quantitative polymerase chain reaction; shNC, non-targeting short hairpin RNA.

group was also significantly higher than those in the other two groups ($P < 0.05$; Fig. 3B). The morphological alteration of cell nuclei in different groups was observed with Hoechst staining. More Hoechst-positive nuclei were recorded in the shSHH group than in the control or shNC groups (Fig. 3C). To further reveal the function of SHH in the induction of apoptosis in WERI-Rb-1 cells, the expression of indicators associated with cell apoptosis was analyzed: It was found that the expression of pro-apoptotic molecules (cleaved caspase-3, cleaved PARP and Bax) was upregulated, whereas the expression of anti-apoptotic molecules (Bcl-2, p-PI3K and p-Akt) was downregulated (Fig. 3D). On the basis of quantitative analyses, the difference in relative expression level between the shSHH group and the control or shNC groups was statistically different for these 6 groups (Fig. 3D; $P < 0.05$).

Knockdown of SHH gene inhibits the activation of the PI3K/Akt pathway. The effect of *SHH*-knockdown on the PI3K/Akt pathway was further analyzed by administering cells in the shNC and shSHH groups with IGF-1. IGF-1 treatment

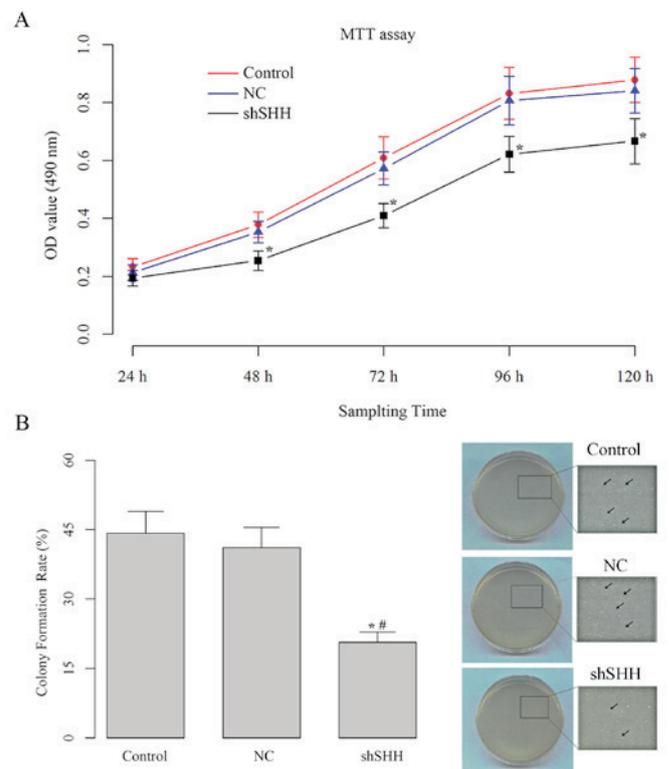


Figure 2. Knockdown of *SHH* inhibits the viability and anchorage-dependent growth ability of WERI-Rb-1 cells. (A) Quantitative analysis results of MTT assay. (B) Quantitative analysis result and representative images of colony formation assay. Arrows indicate colonies. ^a $P < 0.05$ vs. control group; ^b $P < 0.05$ vs. shNC group. SHH, sonic hedgehog; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; shNC, non-targeting short hairpin RNA; OD, optical density.

increased cell viability, which was attenuated by transfection with shSHH (Fig. 4A). However, the difference between the untreated shSHH and shSHH+IGF-1 groups was not statistically significant ($P > 0.05$). Contrary to the results of the MTT assay, increased apoptosis induced by knockdown of *SHH* was markedly inhibited by IGF-1; the difference between the untreated shSHH and shSHH+IGF-1 groups was statistically significant ($P < 0.05$; Fig. 4B). Moreover, following administration of IGF-1, the expression patterns of apoptosis-related molecules in the shSHH+IGF-1 group were reversed with regards to the control when compared with the untreated shSHH group (Fig. 4C).

Discussion

Recent studies have demonstrated that levels of SHH are elevated in multiple types of human cancer (13-17), which subsequently results in the aberrant activation of the HH signaling cascade. Thus, the SHH-dependent initiation of the HH pathway has been hypothesized to be a critical mechanism for the promotion of tumor cell viability. Emerging research based on different types of cancer has validated this hypothesis by revealing multiple important pro-cancer pathways associated with the dysregulated expression of SHH (7,8,18). Consequently, the role of SHH in RB was investigated in the present study, the major finding of which was that knockdown of *SHH* decreased the proliferation of RB cells and induced

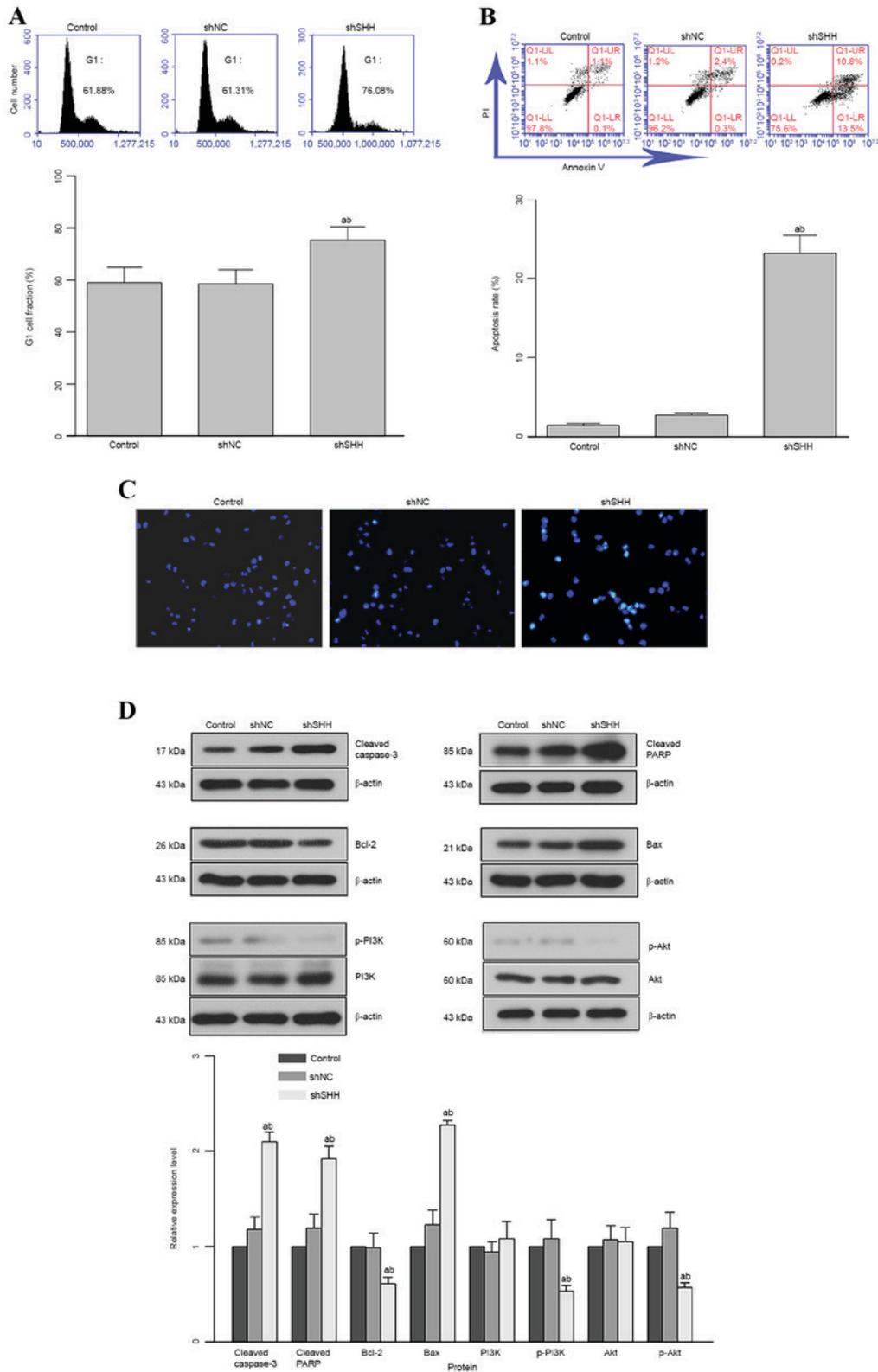


Figure 3. Knockdown of *SHH* induces cell-cycle arrest and apoptosis in WERI-Rb-1 cells. (A) Representative image and quantitative analysis of cell-cycle distribution. (B) Representative image and quantitative analysis result of apoptosis. (C) Representative images of Hoechst staining; nuclei of cells in shSHH group were stained bright blue and abnormally shaped (x400 magnification). (D) Representative image and quantitative analysis result of western blotting assay. ^aP<0.05 vs. control group; ^bP<0.05 vs. shNC group. SHH, sonic hedgehog; shSHH, short hairpin RNA targeting SHH; shNC, non-targeting short hairpin RNA; Bcl-2, B-cell chronic lymphocytic lymphoma 2; PARP, poly (ADP-ribose) polymerase; Bax, Bcl-2-associated X; p-PI3K, phosphorylated phosphoinositide-3 kinase; PI, phosphoinositide.

apoptosis. Moreover, the effect of SHH in RB was mediated through a PI3K/Akt-dependent manner.

Choe *et al* (9) verified the aberrant expression of SHH in patients with RB and identified SHH as potential therapeutic

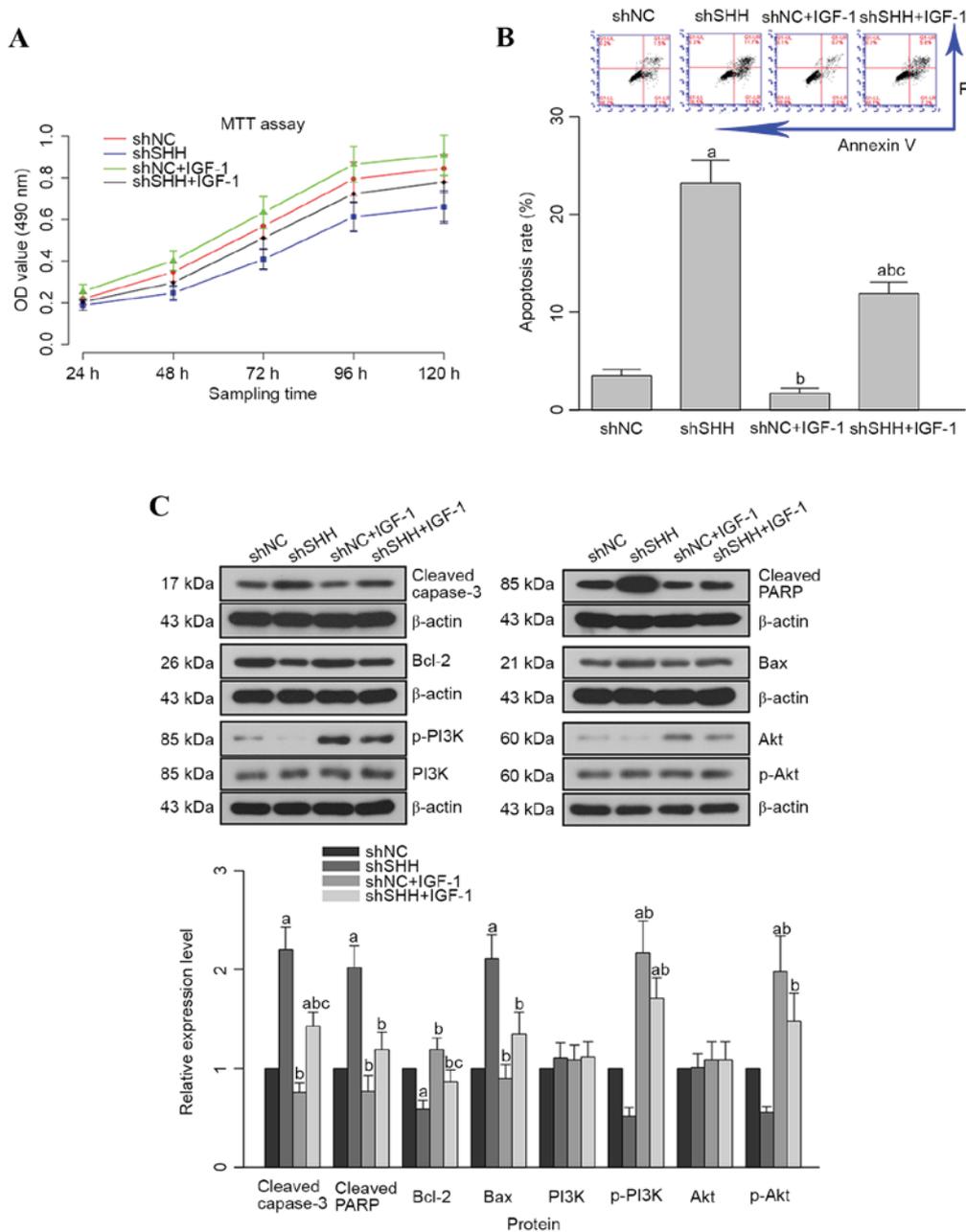


Figure 4. Administration of PI3K/Akt agonist IGF-1 increases cell viability and inhibits apoptosis in WERI-Rb-1 cells. (A) Quantitative analyses results of MTT assay. (B) Representative image and quantitative analysis result of western blotting assay; the expression of anti-apoptosis molecules (Bcl-2, p-PI3K and p-Akt) was upregulated, while the expression of pro-apoptosis molecules (cleaved caspase 3, cleaved PARP and Bax) was inhibited. (C) Representative image and quantitative analysis result of apoptosis. ^aP<0.05 vs. shNC group; ^bP<0.05 vs. shSHH group; ^cP<0.05 vs. shNC+IGF-1 group. p-PI3K, phosphorylated phosphoinositide-3 kinase; IGF-1, insulin-like growth factor 1; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; Bcl-2, B-cell chronic lymphocytic lymphoma 2; PARP, poly (ADP-ribose) polymerase; Bax, Bcl-2-associated X; shSHH, short hairpin RNA targeting SHH; shNC, non-targeting short hairpin RNA; OD, optical density.

target for RB treatment. However, the study failed to provide a comprehensive mechanism to underpin these findings. Indicating that SHH is involved in oncogenesis and the development of RB, the expression of *SHH* was downregulated by a *SHH*-specific shRNA in the human RB WERI-Rb-1 cell line in the current study. Following transfection, the cell viability and anchorage-independent growth ability were markedly decreased. Indeed, the anchorage-independent growth ability was representative of the oncogenicity of cells. Thus, it can be inferred that the knockdown of *SHH* evidently

reduces the ability of RB to transform into a malignancy. SHH serves a central role in embryonic development and controls key genes that modulate cell proliferation (19) directly and indirectly (4,20). During carcinogenesis, SHH exerts its function in a similar pattern to normal development and organ growth, but this aberrant signaling activity promotes the recapitulation of the development ability in cancer cells.

In addition, in the present study, expression of *SHH* not only promoted the growth of RB cells, but also inhibited apoptosis in RB cells. As illustrated by flow cytometry,

knockdown of *SHH* induced G1 cell-cycle arrest and apoptosis in the WERI-Rb-1 cell line. At the protein level, the expression of certain apoptosis-related indicators was also detected. Pro-apoptotic factors, including cleaved caspase-3, cleaved PARP and Bax, were all upregulated following *SHH* shRNA transfection. By contrast, the activity of anti-apoptotic factors, including Bcl-2, PI3K and Akt, were inhibited when *SHH* expression was knocked down. As aforementioned, SHH-related pathways do not always proceed directly, but rather induce a complex network of cascades that cross-talk with other pathways to successfully exert a role in different biological processes (21,22). The PI3K/Akt signal transduction pathway is closely associated with SHH. Previous studies have shown that SHH can promote the epithelial-mesenchymal transition (EMT) in ovarian cancer, and promote metastasis and lymphangiogenesis in gastric cancer in a PI3K/Akt dependent manner (8,18). Riobó *et al* (23) suggested that PI3K/Akt has a synergistic function in the SHH signaling in embryonic development and cancer. To investigate further the interaction between PI3K/Akt and SHH in RB, WERI-Rb-1 cells that had been transfected with *SHH*-specific shRNA were treated with IGF-1 in the present study. This treatment significantly increased the expression of anti-apoptotic molecules, while decreasing the apoptotic rate of *SHH*-knockdown RB cells. These results clearly demonstrated that the activation of the PI3K/Akt pathway blocked the antitumor effect of *SHH*-knockdown. Prior investigation suggests that it is possible that the effect of PI3K/Akt pathway on SHH signaling is modulated through the enhancement of Gli-dependent transcriptional activity (18), which requires further study. However, the role of PI3K/Akt in regulating the SHH pathway varies between cell types, as PI3K/Akt is critical for SHH-mediated EMT and metastasis in multiple cancer types (18). Therefore, the suppression of the PI3K/Akt pathway is key for the development of SHH-based anti-RB therapies. It is thus recommended that treatment modalities targeting SHH signaling should take into account the influence on the PI3K/Akt pathway.

In conclusion, the results of the present study suggest that SHH signaling plays an important role in the development of RB, and that knockdown of *SHH* could decrease cell growth and induce apoptosis in RB cells. At the molecular level, the function of SHH was found to closely associate with the activation of the PI3K/Akt pathway. However, the present study did not detect a change in *SHH* level following administration of IGF-1 and could not provide a detailed explanation for the response of SHH expression to the regulation of the PI3K/Akt pathway. Thus, further studies should be conducted to reveal further the SHH-associated signaling transduction during RB.

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