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ORIGINAL RESEARCH

Bag-I Silence Sensitizes Non-Small Cell Lung Cancer Cells To Cisplatin Through Multiple Gene Pathways

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Purpose: B-cell lymphoma-2 (Bcl-2) associated athanogene 1 (Bag-1) is a multifunctional protein, and Bag -1 overexpression is associated with progression, metastasis, and drug resistance in lung cancer. This study assessed the effects of Bag-1 siRNA on sensitization of cisplatin on non-small cell lung cancer (NSCLC) cells.

Material and methods: NSCLC A549 cell line was transfected with Bag-1 or negative control siRNA and then treated with cisplatin for cell viability, CCK-8, LDH, and flow cytometry assays. The Ca^{2+} levels were analyzed using Fluo-3/AM fluorescence staining, and the protein levels were assessed using Western blot analysis.

Results: Bag-1 siRNA significantly knocked down Bag-1 expression and inhibited cell invasion versus the negative control siRNA, while Bag-1 silence sensitized cisplatin to induce A549 cells to apoptosis by induction of cell cycle G1 arrest. At protein level, Bag-1 silence reduced the expression ratio of Bcl-2 to Bcl-2 associated X protein (Bax), downregulated activity of the PI3K/AKT and mitogen-activated protein kinase (MAPK) pathways, and potently upregulated the calcium signaling-mediated pathway. **Conclusion:** This study demonstrated that Bag-1 silencing sensitized A549 to cisplatin to enhance A549 cell apoptosis by modified multiple gene pathways. Further study will evaluate the usefulness of Bag-1 siRNA as a potential targeting therapy for NSCLC.

Keywords: bag-1, apoptosis, non-small cell lung cancer, chemotherapy

Introduction

Lung cancer is the leading cause of cancer-related deaths,¹ while lung cancer has become the first ranked cause of cancer death in People's Republic of China since the beginning of the twenty-first century.^{2,3} 70–80% of lung cancer patients are diagnosed at an advanced stage of disease and lose the opportunity for curative surgery.^{2,3} Thus, platinum-based chemotherapy is the standard regimen to control non-small-cell lung cancer (NSCLC), prolonging survival of these patients. However, drug resistance (e.g., cisplatin-induced therapeutic resistance in lung cancer patients) causes a significant obstacle to satisfactory overall survival of NSCLC patients. Thus, it is urgent to understand lung carcinogenesis and drug resistance mechanisms in order to develop novel strategies to effectively control lung cancer.

Apoptosis is essential for maintaining the cell and tissue physiological state and B-cell lymphoma-2 (Bcl-2) associated athanogene 1 (Bag-1) is a

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multi-functional protein⁴ and can bind to Bcl-2 to inhibit cell apoptosis. For the different initiation sites during bag-1 protein translation, Bag-1 has four isoforms, including Bag-1L (52 kDa), Bag-1 (34 kDa), Bag-1M (46-48 kDa), and Bag-1S (29 kDa),⁵ expressed indifferent nuclear and cytoplasmic localization patterns.⁶ Bag-1 was overexpressed in various human cancers, including lung cancer.⁷ The association between expression of Bag-1 and a prognosis in lung cancer remains controversial. For example, Wang YD et al.⁷ showed that Bag-1 expression was associated with prolonged survival of patients with NSCLC. Differently, previous studies showed that high expression of BAG1 associated with poor outcome of NSCLC.^{8,9} Additional studies reported that Bag-1 expression was associated with a favorable prognosis of NSCLC after chemotherapy.^{10,11} Thus, in this study, we assessed the effects of Bag-1 silencing on regulation of cisplatin sensitivity to NSCLC cells in vitro and explored the underlying molecular events. We expected to provide useful information regarding the role of Bag-1 in mediation of chemotherapy efficacy for future control of NSCLC.

Materials And Methods Cell Culture

A human NSCLC A549 cell line was provided by the Teaching and Research Section of Microbiology of University Medical Qingdao College (Qingdao, People's Republic of China) and the use of this cell line was approved by the institutional review board of Qingdao University. A549 cells were cultured in Dulbecco's modified Eagle's medium (DMEM) (Hyclone, Logan, UT, USA) supplemented with 10% fetal bovine serum (Hyclone), penicillin (100 U/mL), and streptomycin (100 µg/mL) in a humidified atmosphere with 5% CO₂ at 37°C, and the culture media was replaced every 2-3 days.

Bag-1 siRNA And Cell Infection

Lentivirus carrying Bag-1 and negative control siRNA was purchased from Shanghai Genechem Company (Shanghai, People's Republic of China). A549 cells were grown and divided into three groups: 1) a group of non-treatment cells without any gene transfection, 2) a control siRNA group where the cells were infected with the negative control siRNA, and 3) a Bag-1 siRNA group where the cells were infected with Bag1 siRNA. The cells were then infected with these lentiviruses using the manufacturer's instructions. The infected cells were then selected with 1 μ g/mL puromycin to reach 80% confluency, and the cell culture medium was changed to DMEM containing various doses of cisplatin (Kocak Pharma, Turkey) and cultured for 24 hrs for the following experiments.

Transwell Cell Invasion Assay

A Transwell invasion assay was performed to assess tumor cell invasion with Transwell filters that were precoated with Matrigel (Corning, Corning, NY, USA). Specifically, 200 µl transfected cells $(7 \times 10^4/\text{mL})$ were seeded into the upper chamber of 24-well plate inserts with serum-free medium, while 600 µl of medium supplemented with 20% FBS was added to the lower chamber as a chemoattractant. The cells were then grown for 48 hrs at 37°C with 5% CO₂. At the end of the experiment, the cells that invaded the reverse side of the filters were fixed in methanol for 30 mins and stained with crystal violet (0.1%)for 20 mins at 25°C. The number of cells was counted in five randomly selected fields under a 400-fold magnification and summarized.

Cell Viability Assay

The changed cell viability was measured using the Cell Counting Kit-8 (CCK-8; Boster, Wuhan, People's Republic of China). Briefly, 5×10^3 cells/well were seeded into 96-well plates and incubated for 24 hrs and then treated with various doses (0, 0.625, 1.25, 2.5, 5, 10, 20, 30, or 40 µg/mL) of cisplatin for 24 hrs. Subsequently, 10 µl CCK-8 reagent was added to the cell culture medium, and the cells were further incubated for 1 hr. At the end of the experiment, the absorbance rate of each well was measured at 450 nm using a microplate reader (Bio-Tek, Winooski, VT, USA), while 100 µl DMEM was used as a blank control. The experiments were in triplicate and repeated at least three times. To calculate the IC_{50} , the optical density (OD) values were measured after the cells were treated with different doses (0, 0.625, 1.25, 2.5, 5, 10, 20, 30, or 40 µg/mL) of cisplatin for 24 hrs.

Flow Cytometry Cell Cycle Assay

A549 cells were seeded into 6-well plates at a density of 2×10^5 cells/well and incubated for 24 hrs and infected with Bag-1 or negative control siRNA for 48

hrs and then treated with 5 μ g/mL cisplatin for an additional 24 hrs. After that, the cells were measured for cell cycle distribution using the Propidium Iodide (PI) flow cytometry kit (Abcam, Cambridge, UK) using the manufacturer's recommendations. In brief, the cells were washed and incubated with trypsin for 3 mins at 37° C and then fixed with 66% ethanol on ice. 1×10^{6} cells per group were stained with 5% PI in phosphatebuffered saline (PBS) containing 0.5% RNase A and subjected to DNA content analysis using FACStar Plus flow cytometer (BD Biosciences, Franklin Lakes, NJ, USA). The mean value of three independent experiments was analyzed by using FlowJo (BD Biosciences).

Flow Cytometry Apoptosis Assay

A549 cells were seeded into 6-well plates at a density of 2×10^5 cells/well and incubated for 24 hrs and infected with Bag-1 or negative control siRNA for 48 hrs and then treated with 5 µg/mL cisplatin for an additional 24 hrs. After that, the cells were harvested using trypsin without EDTA and resuspended in 1× binding buffer to a concentration of $1-5 \times 10^6$ cells/mL from the Annexin V-APC and propidium iodide (PI) apoptosis detection kit (eBioscience, San Diego, CA, USA) using the manufacturer's protocol. Next, the apoptotic cells were quantified with a FACStar Plus flow cytometer (Becton Dickinson analyzed using BD Accuri C6 software). The mean value of triplicate experiments was used to analyze data.

Lactate Dehydrogenase (LDH) Activity And The Intracellular Ca²⁺ Concentration Assays

To measure the LDH activity in cells, five parallel wells were set for each experimental group and infected with Bag-1 or negative control siRNA for 48 hrs and then treated with 5 μ g/mL cisplatin for an additional 24 hrs. Next, the culture medium was collected for detection of the LDH activity using an LDH activity assay kit (Nanjing Jian Cheng Biotechnology, Nanjing, People's Republic of China) using the manufacturer's protocol. Each sample was run in triplicate.

To assay the intracellular Ca²⁺ concentration, the duplicate cells were subjected to the Fluo-3/AM kit (Jiangsu Biyuntian Biological Technology Research Institute, Jiangsu, People's Republic of China) analysis using the manufacturer's protocol. Each sample was run in triplicate.

Quantitative Reverse Transcriptase-Polymerase Chain Reaction (qRT-PCR)

Total cellular RNA was isolated from the cells by using the Trizol reagent (Invitrogen, Carlsbad, CA, USA) using the manufacturer's instructions and quantified by UV spectrophotometry (Implen, Germany) and then reversely transcribed into cDNA using a M-MLV Reverse Transcriptase Kit (Roche, Swiss) using the manufacturer's protocol. These cDNA samples were then mixed with FastStart Essential DNA Green Master Mix (Roche, Basel, Switzerland) and amplified in a Light Cycler 96 (Roche, Indianapolis, IN, USA) using the manufacturer's protocol. All gene primers were purchased from Genetech Co., Ltd, People's Republic of China, and the Bag-1 primer sequences were 5'-CTTCATGTTACCTC CCAGCA-3' and 5'-ACGGTGTTTCCATTTCCTTC-3', and the GAPDH primer sequences were 5'-ATCTCT GCCCCCTCTGCTGA-3' and 5'-GATGACCTTGCCCA CAGCCT-3'. Each experiment was repeated in triplicate. The relative level of Bag-1 mRNA was calculated using the 2- $\Delta\Delta$ Ct method.

Western Blot

The experiment used the methods described previously.¹² Briefly, A549 cells were collected and washed three times with ice-cold PBS, and the total cellular protein was extracted with a lysis buffer containing 200 µl ice-cold RIPA lysis buffer, 2 µl ice-cold PMSF lysis buffer, and 2 µl ice-cold phosphatase inhibitors for 30 mins. The supernatant fraction was collected after the mixture was centrifuged for 15 mins at 12,000 rpm. The protein concentration was determined using the bicinchoninic acid (BCA) protein assay kit (TransGen Biotech, Beijing, People's Republic of China), and the protein samples were then denaturated at 100°C for 5 mins in the loading buffer and separated in 10% sodium dodecyl sulfatepolyacrylamide gel electrophoresis (SDS-PAGE) gels and blotted onto a PVDF membrane (Bio-Rad, Hercules, CA, USA) and Western blot procedures.¹¹ The primary antibodies used were an anti-Bag-1 (1:1000; Cell Signaling Technology, Danvers, MA, USA), anti-Hsp70 (1:1000; Abcam), anti-Bcl-2 (1:1000; Cell Signaling Technology), anti-Bax (1:1000; Technology), anti-Cleaved-Caspase-3 Cell Signaling (1:1000; Cell Signaling Technology), anti-IP3R (1:1000; Cell Signaling Technology), anti-P-IP3R (1:1000; Cell Signaling Technology), anti-AKT (1:10,000; Abcam), anti-P-AKT (1:5000; Abcam), anti-ERK (1:1000; Abcam), anti-P-ERK (1:1000; Abcam), and anti- β -Actin antibodies (1:1000; Boster, Wuhan, People's Republic of China). The secondary antibodies (the goat anti-mouse for Bag-1, while the goat anti-rabbit antibody for other primary antibodies were from Boster (Wuhan, People's Republic of China), and the positive bands were visualized using the Fusion FX Imaging System (Vilber Lourmat, Paris, France) and quantified by using the Bio-Rad quantity one software (Hercules, CA, USA).

Statistical Analysis

All values were expressed as a mean plus or minus the standard deviation (SD) and analyzed with the SPSS 17.0 statistical software (SPSS, Chicago, IL, USA). A One-way analysis of variance (ANOVA) followed by a Tukey's test was performed to compare differences in multiple comparisons. A *p*-value less than 0.05 was considered to be statistically significant.

Results

Effect Of Bag-1 Silencing On Regulation Of Tumor Cell Invasion

After 48 hrs of infection with lentivirus carrying Bag-1 siRNA, the fluorescence microscopic data showed that

A549 cells were successfully infected by the lentivirus (Figure 1). Western blot results showed that expression of Bag-1 protein isoforms was significantly downregulated in Bag-1 siRNA group vs negative control siRNA group (Figure 2A), while qRT-PCR confirmed a dramatic decrease in Bag-1 mRNA level in Bag-1 siRNA-infected A549 cells vs the negative control siRNA group and the non-treatment cells (Figure 2B). As shown in Figure 3A and B, Bag-1 knockdown inhibited A549 cell invasion vs the cells infected with negative control siRNA (p = 0.011).

Bag-I Silence Increased A549 Cell Cytotoxicity After Cisplatin Treatment

After that, we first assessed the effect of Bag-1 silencing on regulation of cell viability. With increase in cisplatin concentrations, the cell viability of each group was decreased, but viability of Bag-1 siRNA-infected cells was even lower than that of the negative control siRNA group and non-treatment group. There was no statistical difference between the non-treatment and the negative control siRNA groups, whereas a lower IC50 was observed in Bag-1 siRNA-infected A549 cells (Figure 4A). After cisplatin concentration reached 5 μ g/ mL, the cell viability of Bag-1 siRNA group was significantly lower than that of non-treatment group (*p*=0.005) and the negative control siRNA group (*p*=0.003;



Figure I Infection of A549 cells with lentivirus carrying Bag-I siRNA. A549 cells were grown and infected by Bag-I or negative control siRNA. (A) Green fluorescence microscopy 48 hrs after infection. (B) Light field of the fluorescence microscopy 48 hrs after infection.



Figure 2 Silencing of Bag-I expression using Bag-I siRNA.A549 cells were grown and infected by Bag-I or negative control siRNA for 48 hrs. (**A**) Western blot results. (**B**) This graph is data of the Bag-I mRNA levels. **p*<0.001 vs the non-treatment group and **p* <0 0.001 vs the negative control siRNA group.

Figure 4B). We, therefore, used this 5 μ g/mL of cisplatin as a choice for our further experiments.

Next, we performed the flow cytometric assay to assess the changed cell apoptosis in Bag-1 silencing cells after 5 µg/mL cisplatin treatment. Our data showed that Bag-1 silencing enhanced the levels of both early and late apoptotic cells compared to that of non-treatment group (p=0.007) and the negative control siRNA group (p=0.01), while there was no difference occurred between the non-treatment and negative control siRNA groups (p=0.74; Figure 5). Cell cycle distribution assay showed that Bag-1 silencing decreased the percentage of cells at the S phase of the cell cycle but significantly increased the percentage of cells in the G1 phase of the cell cycle compared to those of the non-treatment and negative control siRNA cells (Figure 6).

Furthermore, the LDH, a glycolytic enzyme, leaks from the cytoplasm to extracellular during apoptosisinduced cell membrane permeability and damage. Thus, the LDH is widely used for cell cytotoxicity detection.^{13,14} We found that bag-1 silencing of A549 cells increased LDH (p=0.000 vs that of the non-treatment group and p=0.001 vs.that of the negative control siRNA group; Figure 7). Therefore, these data suggested that Bag-1 silencing promoted cisplatin-induced A549 cell apoptosis in vitro.

Downregulation Of BcI-2, PI3K/AKT, And The Mitogen-Activated Protein Kinase (MAPK) Pathway And Upregulation Of Caspases-3 Activity After Bag-1 Silencing

We then investigated the molecular events underlying Bag-1 silencing in A549 cells and cisplatin treatment. Our data showed that Bag-1 silencing lowered Bcl-2 levels and downregulated the phosphorylation of AKT and extracellular signal-regulated kinase (ERK1/2). However, Bag-1-silenced cells did not alter the expression of 70 kDa heat shock proteins (HSP70) and Bcl-2-associated x protein (Bax) compared to those of the non-treatment cells and negative control siRNA cells (Figure 8). To examine the effect of Bag-1 silencing on the regulation of caspase activity, we assayed the levels of caspase 3 and observed an increase in cleaved caspase-3 (17 kDa) in Bag-1 siRNA cells compared to the non-treatment and negative control siRNA cells (Figure 8).

Effect Of Bag-I Silencing On Regulation Of The Intracellular Concentration Of Ca²⁺ And Inositol 1,4,5-Trisphosphate Receptor (IP3R) Phosphorylation In A549 Cells

As a second messenger in cells, Ca^{2+} plays an extremely important role during cell injury.^{15,16} To investigate whether Bag-1 silencing increased the sensitivity of cisplatin in the induction of A549 cell apoptosis through calcium overload, we performed Fluo-3/AM staining. Our data showed that Bag-1 silencing significantly elevated intracellular Ca²⁺ levels versus the non-treatment and negative control siRNA cells (both *p*-values were less than 0.001; Figure 9A), while phosphorylated IP3R levels were also enhanced in the Bag-1 siRNA group versus the non-treatment and negative control siRNA cells (Figure 9B). To further confirm whether calcium



Figure 3 Effect of Bag-1 silencing on the inhibition of tumor cell invasion. A549 cells were grown and infected with lentivirus carrying Bag-1 or negative control siRNA for 48 hrs and then subjected to Transwell tumor cell invasion assay. (A) Invasion cells under a microscope. (B) the relative invasion rate. *p=0.0011 vs the negative control siRNA group.

signaling is involved, we overexpressed Bag-1 in A549 cells using lentivirus carrying Bag-1 cDNA (e-Bag-1) or vector-only (control e-Bag-1). Our data showed the opposite results and changes the intracellular Ca^{2+} and phosphorylated IP3R levels (Figure 9C and D).

Discussion

Activity of the apoptosis signaling pathways is frequently disturbed in lung cancer cells, resulting in infinite tumor cell growth and proliferation but resistance to chemotherapy. Thus, better understanding of the molecular mechanism by which agents promote tumor cells to undergo apoptosis becomes an important direction of research in lung cancer treatment. In the current study, we investigated the pro-apoptotic effects of Bag-1 silencing in combination with cisplatin treatment in NSCLC A549 cells. Cisplatin is a well established and effective chemotherapeutic agent against



Figure 4 Effects of Bag-I silence and cisplatin on the regulation of A549 cell viability. **(A)** Calculation of the IC₅₀, and the data is presented as the mean plus or minus the standard deviation of three independent experiments. **(B)** Cell viability assay. A549 cells were grown and infected by Bag-I or negative control siRNA for 48 hrs and then treated with cisplatin for 24 hrs and subjected to a cell viability assay. $^{*}p$ <0.05 vs the non-treatment group; $^{*}p$ <0.05 vs the negative control siRNA group; $^{*}p$ <0.01 vs the non-treatment group $^{\#}p$ <0.01 vs the negative control siRNA group.

various human cancers and induces DNA damage and apoptosis in highly proliferating cancer cells.¹⁷ Thus, "the background apoptotic effect" of cisplatin was quite substantial (>20%) in the "non-treatment" and "control siRNA treated" cells. We first performed lentivirus infection carrying Bag-1 or negative control siRNA into A549 cells and found that Bag-1 siRNA effectively silenced Bag-1 expression in A549 cells and inhibited cell invasion. After that, these infected A549 cells were subjected to various doses of cisplatin treatment. Our data showed that Bag-1 silencing reduced A549 cell viability and triggered them to undergo apoptosis. The caspase family is a group of proteolytic enzymes that execute cell apoptosis and Caspase-3 is the final step of the apoptosis cascades of proteolytic reactions responding to the apoptotic signals; thus, Caspase-3 is the key executor of apoptotic events.¹⁸ In our current study, Bag-1 silencing led to cleavage of caspases-3, indicative of activation of caspase 3 in cells. This phenomenon was in line with the cytotoxic effects induced by Bag-1 silencing and cisplatin.

To further identify the potential mechanisms of Bag-1 silencing, we assessed the therapeutic potential of Bag-1 silencing and explored the mechanism by which Bag-1 silencing sensitized A549 cells to cisplatin-induced apoptosis. Oltvai et al revealed that Bcl-2 can bind to Bax to form Bcl-1/Bax heterodimer¹⁹ and the ratio of Bcl/Bax determines the survival of cells after receiving the apoptotic signals. In other words, when Bax is dominant, cell apoptosis is accelerated. In the current study, we showed the decrease in ratio of Bcl to Bax, although the Bax levels did not show significant change in A549 cells after Bag-1 silencing and cisplatin treatment. Furthermore, the Bag family is one of those known cochaperones to Hsp70.²⁰⁻²² Bag-1 binds to Hsp70 to modulate its chaperone activity in vitro,²³ which is closely related to the anti-apoptotic function, tumorigenesis, and cell cycle redistribution in different physiological and diseased processes.²⁴ Bag-1 can positively or negatively regulate the function of Hsp70. For example, Bag-1L overexpression could elevate HSP70 level in neuroblastoma cells,²⁵ while Liu et al¹¹ reported that Bag-1 siRNA increased the levels of Hsp70 protein in A549 cells. In our current study, we were unable to show change in Hsp70 expression in A549 cells after Bag-1 silencing, thus, further study of Bag-1 and its interaction with Hsp70 is warranted to explore the underlying mechanisms for NSCLC resistance to cisplatin.

In physiological conditions, the PI3K/AKT pathway plays a role in the regulation of cell growth proliferation and angiogenesis, and dysregulation of the PI3K/AKT pathway promotes cancer development.²⁶ The P-AKT can activate the transcription factors FOXO1, FOXO 3, and FOXO4 to promote NSCLC cell progression to the S phase of the cell cycle.²⁷ Ali et al observed that the PI3K inhibitors were able to delay tumor growth in an animal model.²⁸ The PI3K/AKT pathway was highly activated in approximately 90% of NSCLC cell lines and involved in drug resistance.^{29,30} In our current study, AKT phosphorylation was potently suppressed after Bag-1 silencing, accompanied by G1 phase cell



Figure 5 Effects of Bag-1 siRNA and cisplatin on the regulation of A549 cell apoptosis. A549 cells were grown and infected by Bag-1 or negative control siRNA for 48 hrs and then treated with 5 μ g/mL cisplatin for 24 hrs and subjected to a flow cytometric apoptosis assay. (A) Representative results of the Annexin V-APC/PI staining of A549 cells. Q3-UL necrosis, Q3-UR late apoptosis, Q3-LR early apoptosis, and a Q3-LL viable cell. (B) The graph is quantified data of the assay. *p=0.007 vs the non-treatment group and #p=0.010 vs the negative control siRNA group.

cycle arrest. As we know, cells in the G0/G1 phase are insensitive to chemotherapy, suggesting that Bag-1 silencing may downregulate the PI3K/AKT pathway to promote tumor cell apoptosis after cisplatin. In addition, the MAPK signal transduction pathway is the most important cell division regulatory system and plays an



Figure 6 Effects of Bag-I siRNA and cisplatin on the regulation of cell cycle redistribution. (**A**) A549 cells were grown and infected by Bag-I or negative control siRNA for 48 hrs and then treated with 5 μ g/mL cisplatin for 24 hrs and subjected to a flow cytometric cell cycle assay. (**B**) The columns indicate the percentages of A549 cells at different phases. G0/GI phase, *p=0.02 vs the non-treatment group and #p=0.04 vs the negative control siRNA group. The S phase, *p=0.009 vs the non-treatment group and #p=0.02 vs the negative control siRNA group.



Figure 7 Effects of Bag-1 siRNA and cisplatin on the regulation of LDH activity in A549 cell culture medium. A549 cells were grown and infected by Bag-1 or negative control siRNA for 48 hrs and then treated with 5 μ g/mL cisplatin for 24 hrs and subjected to the LDH assay. **p*=0.000 vs.the non-treatment group and **p*=0.001 vs the negative control siRNA group.



Figure 8 Effects of Bag-I siRNA and cisplatin on the regulation of various protein expression and activity. A549 cells were grown and infected by Bag-I or negative control siRNA for 48 hrs and then treated with 5 μ g/mL cisplatin for 24 hrs and subjected to Western blot analysis of Hsp70, Bax, Bcl-2, P-AKT, AKT, P-ERK, ERK, and cleaved-Caspase-3.

important role in the process of cell division and proliferation.³¹ The ERK1/2 are the main and classical pathway of the MAPK families and involved in

regulation of cell cycle progression and cell proliferation.³² Thus, targeting of the MAPK pathway could enhance the efficiency of chemotherapy. In our current study, we found that the level of phosphorylated ERK1/2 was affected by Bag-1 silencing.

Normally, intracellular calcium is mainly stored in the endoplasmic reticulum (ER)/sarcoplasmic reticulum. To maintain an intracellular Ca²⁺ dynamic homeostasis, Ca²⁺ in the ER is mainly released by IP3R and RyR into the cell cytoplasm. At the same time, Ca²⁺ in the cytoplasm is ingested into the ER through SERCA.³³ A previous study showed that calcium can be released from the ER to induce calcium overload, which, in turn, initiates mitochondrial-dependent apoptosis.³⁴ As a universal intracellular calcium release channel, the IP3R is considered to be activated by its phosphorylation.³⁵ In our current study, we found that the Ca²⁺ concentration was significantly elevated in the Bag-1 siRNA group, and, meanwhile, IP3R was activated by phosphorylation, while Bag-1 overexpression had the opposite change. Thus, we speculate that P-IP3R may be involved in Bag-1 silence-induced A549 cell apoptosis. Furthermore, Bcl-2 protein can block the release of calcium from the ER and decrease the activity of calcium-dependent endonuclease, thereby blocking cell apoptosis.^{36,37} A previous study revealed that Bcl-2 was able to dephosphorylate IP3R and suppress the channel activity.^{38,39} In our current study, we found that Bag-1 silencing enhanced IP3R phosphorylation and downregulated Bcl-2 expression in the cisplatin-treated A549 cells, suggesting that Bcl-2 downregulation could upregulate IP3R phosphorylation to improve the effectiveness of cisplatin treatment of A549 cells. In the present study, Bag-1 silencing inhibited A549 cell invasion and sensitized them to cisplatin-induced apoptosis via multiple pathways. Silencing of Bag-1 expression using Bag-1 siRNA like some tumor suppressors⁴⁰ might represent a potential targeting therapy for NSCLC.

Conclusions

Bag-1 silencing sensitized NSCLC A549 cells to cisplatininduced apoptosis through multiple pathways, i.e., decrease in the Bcl to Bax ratio, downregulation of the PI3K/AKT and MAPK pathway activity, and activation of IP3R.



Figure 9 Effects of Bag-1 siRNA and cisplatin on the regulation of the intracellular calcium overload. (**A** and **C**) The Fluo-3/AM and flow cytometry assay. A549 cells were grown and infected by Bag-1 siRNA, negative control siRNA, lentivirus carrying vector-only (control e-Bag-1), or carrying Bag-1 cDNA (e-Bag-1) for 48 hrs and then treated with 5 μ g/mL cisplatin for 24 hrs and subjected to an intracellular calcium assay. The percentage of total cells stained by Fluo-3/AM for intracellular Ca²⁺ levels was determined by flow cytometry. The values are the mean plus or minus the standard deviation. (**B** and **D**) Western blot. A549 cells were grown and infected with Bag-1 siRNA, negative control siRNA, lentivirus carrying vector-only (control e-Bag-1), or carrying Bag-1 cDNA (e-Bag-1) for 48 hrs and then treated with Bag-1 siRNA, negative control siRNA, lentivirus carrying vector-only (control e-Bag-1), or carrying Bag-1 cDNA (e-Bag-1) for 48 hrs and then treated with 5 μ g/mL cisplatin for 24 hrs and subjected to Western blot of phosphorylated IP3R. *p<0.001 vs the non-treatment group and [#]p<0.001 vs the negative control siRNA group. *p=0.026 vs the non-treatment group and [#]p=0.011 vs the negative control siRNA group.

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Author Contributions

Jiling Lv designed the study. Jiling Lv, Fang Zhang, Congying Zhai, Gejin Wang and Yan Qu carried out the study. Jiling Lv and Congying Zhai participated in the experiments and statistical analysis. Jiling Lv wrote the manuscript. Jiling Lv, Fang Zhang and Yan Qu revised the manuscript. All authors contributed to data analysis, drafting or revising the article, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

Disclosure

The authors report no conflicts of interest in this work.

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