S100A9 regulates cisplatin chemosensitivity of squamous cervical cancer cells and related mechanism

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Objective: Our previous research has shown that the expression of S100 calcium-binding protein A9 (S100A9) in tumor cells was associated with neoadjuvant chemotherapy sensitivity in cervical squamous cell carcinoma. In the present study, we altered the expression of S100A9 through infecting lentivirus, investigated its effect on the chemosensitivity to cisplatin of cervical cancer cells and then made a primary exploration of the involved mechanism.

Materials and methods: Lentivirus was employed to upregulate and downregulate S100A9 expression in SiHa cells. The protein expression level of apoptotic-related proteins Bcl-2 and Bax, drug resistance-related proteins multiple drug resistance protein 1 (MRP1), P glycoprotein (P-gp), glutathione-S-transferase- π (GST- π), lung resistance-related protein (LRP), and FOXO1 signaling pathway related proteins was detected by Western blot. The CCK-8 assay was used to examine chemosensitivity to cisplatin, and the proportion of apoptosis cells was analyzed by the flow cytometry.

Results: S100A9 overexpression could obviously increase the IC₅₀ value of SiHa cells to cisplatin and decrease the apoptosis rate induced by cisplatin. Downregulation of S100A9 led to the opposite results. In S100A9 overexpression SiHa cells, the expression level of Bcl-2, LRP, GST- π , p-AKT, p-ERK, p-FOXO1, and Nanog was significantly increased, while FOXO1 expression was decreased. The opposite results were observed in S100A9 knockdown SiHa cells.

Conclusion: Downregulation of S100A9 could significantly increase apoptosis rate, resulting in enhancing sensitivity of SiHa cells to cisplatin, which may be related to Bcl-2, GST- π , and LRP protein and by altering the AKT/ERK-FOXO1-Nanog signaling pathway.

Keywords: S100A9, cervical cancer, Bcl-2, GST- π , LRP, FOXO1, cisplatin, chemosensitivity

Introduction

In spite of the widespread use of screening tests and the advancement in treatment methods, cervical cancer is still the fourth most frequently diagnosed cancer in women worldwide, leading to an estimated 527,600 new cases and 265,700 deaths in 2012.¹ In America, there were 61,380 estimated new cases of cervical cancer and 10,920 estimated deaths in 2017.² Most early-stage cervical cancer can be cured by surgery and/or radiation with or without chemotherapy.³ For individuals with locally advanced cervical cancer, primary treatment options contain cisplatin-based combination chemotherapies and concurrent radiation.⁴ However, systemic chemotherapies, especially cisplatin-based regimens, often have severe side effects, such as nephrotoxicity, myelosuppression and digestive tract reactions.⁵ Moreover, patients often develop resistance to chemotherapy, resulting in compromising the efficacy of cisplatin.⁶ The molecular mechanisms underlying cisplatin resistance are complex and are not well understood

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yet.⁶ Therefore, it is essential to screen novel strategies to overcome cisplatin resistance in cervical cancer.

The S100 family contains more than 20 members, which exhibit a high degree of sequence and structural similarity. The term S100 is derived from the solubility of these proteins in 100% saturated ammonium sulphate.⁷ S100A9 belongs to the S100 family of calcium-binding proteins, and is also called MRP-14 (myeloid related protein of molecular weight 14 kD).⁸ S100A9 may exist as a homodimer, heterodimer or heterotetramer.⁹ S100A9 protein has been shown to regulate cell proliferation, differentiation, apoptosis, survival, and migration/invasion through interacting with some enzymes, receptors, transcription factors.¹⁰ Arachidonic acid, receptor for advanced glycation end products, Toll-like receptor 4, the major fatty acid transporter CD36, matrix metalloproteinases, fibronectin, and heparin sulfate glycosaminoglycans are ligands of calcium bound S100A9.¹¹ Increased levels of S100A9 were found in many tumors, including ovarian, gastric, esophageal, colon, pancreatic, bladder, thyroid and breast carcinomas.¹²

In our previous study, we detected the expression of S100A proteins in the paired tumor samples (pre- and post-chemotherapy) obtained from 68 squamous cervical cancer patients who were treated with cisplatin-based neoadjuvant chemotherapy and radical hysterectomy, and revealed that S100A9 protein expression in tumor cells correlated with tumor response to chemotherapy.¹³ Similar to our research, Zhu et al¹⁴ found that cervical squamous carcinoma patients, who received the concurrent chemoradiotherapy, with negative staining of S100A9 showed significantly lower 5-year overall survival rate than those with positive staining. The above results suggested that S100A9 may be associated with the chemosensitivity of squamous cervical cancer. However, we also found the expression intensity of S100A9 was statistically higher in post-chemotherapy cervical cancer tissues in comparison with that in pre-chemotherapy.¹³ In our further study,¹⁵ four squamous cervical cancer cell lines were selected with diverse human papillomavirus expression, SiHa, C-33A, CaSki, and MS751, and found that the S100A9 mRNA and protein were moderately expressed in SiHa cells. In addition, S100A9 could promote squamous cervical cancer cell proliferation, migration and invasion. Therefore, whether S100A9 promotes or inhibits cisplatin sensitivity remains controversial.

In the present study, we used lentivirus to upregulate or downregulate the expression of S100A9 in SiHa cells, followed by exploring the role of S100A9 expression on the cisplatin sensitivity in squamous cervical cancer and elucidating the related molecular mechanisms, in order to overcome

the chemoresistance in cervical cancer and provide a thread to explore new treatment strategies.

Materials and methods

Cell lines and culture

The human cervical cancer cell line SiHa was obtained from Shanghai Cell Biology Medical Research Institute, Chinese Academy of Sciences, and maintained as a monolayer in Dulbecco's modified Eagle's medium (DMEM) (Gibco, Thermo Fisher Scientific, Waltham, MA, USA) supplemented with 10% fetal bovine serum (Gibco, Thermo Fisher Scientific) and 100 µg/mL streptomycin. The cells were incubated at 37°C in a humidified atmosphere of 5% CO₂.

Lentivirus production and transduction of target cells

The S100A9 sequences were amplified by polymerase chain reaction, confirmed by sequencing, and then inserted into the lentiviral expression vector pLVX-IRES-ZsGreen 1 (donated by the School of Medical Lab Science, Wenzhou Medical University). The four pGFP-C-shLenti vectors inserted with four different S100A9 small hairpin RNAs (shRNAs) and the negative control plasmid were purchased from Origene (Rockville, MD, USA). To produce virus particles, together with two packaging plasmids psPAX2 and G protein of the vesicular stomatitis virus (VSV-G) envelope plasmid pMD2.G (donated by Dr Luzhe Sun, The University of Texas Health Science Center at San Antonio), S100A9-pLVX-IRES-ZsGreen1, S100A9-pGFP-C-shLenti and the control vector (5.0 µg) were transfected into 293 T cells at the ratio of 5:2:3 for 48 hours, using lipofectamine 3,000 (Thermo Fisher Scientific). SiHa cells were seeded in 60-mm dish and cultured in DMEM medium without antibiotics for 24 hours. Then, SiHa cells were transduced with recombinant lentiviral particles with the help of polybrene (final concentration 8 µg/mL). To produce polyclonal cells with stable expression of S100A9, the infected GFP-positive cells were sorted by flow cytometry (FACSaria II, BD Biosciences, San Jose, CA, USA); the S100A9 down-regulation cells were selected by 2 µg/mL puromycin (Sigma-Aldrich Co., St Louis, MO, USA). The primers and sh-RNA used for gene overexpression or downexpression are listed in Table 1. All cells were confirmed by Western blot.

Western blot analysis

The cells were washed with PBS and lysed with RIPA buffer (150 mM NaCl, 1.0% Nonidet P-40, 0.5% sodium deoxycholate, 0.1% sodium dodecyl sulfate, 50 mM Tris, pH 8.0) containing protease inhibitor cocktail (Beyotime, Shanghai,

Table I Description of the selected genes and specific primers used in this study

Gene	Primer name	Primer sequence (5'–3')
S100A9	S100A9-F	CCG CTC GAG ATG ACT TGC AAA ATG TCG CAG
	S100A9-R	ATA TAT GGA TCC TTA GGG GGT GCC CTC CCG AG
S100A9shA	S100A9shRNA-A	TGT CGC AGC TGG AAC GCA ACA TAG AGA CC
S100A9shB	S100A9shRNA-B	CAT CAA CAC CTT CCA CCA ATA CTC TGT GA
S100A9shC	S100A9shRNA-C	CAT CAA CAC CTT CCA CCA ATA CTC TGT GA
S100A9shD	S100A9shRNA-D	TGA GCT TCG AGG AGT TCA TCA TGC TGA TG

China). The lysates were incubated on ice for 30 minutes. After centrifugation at 12,000 g at 4°C for 20 minutes, the supernatant was collected and bicinchoninic acid assay (Beyotime) was used for protein qualification. Equal amounts of protein were loaded onto a 12% sodium dodecyl sulfate-polyacrylamide gel and then transferred onto a 0.45 µm or 0.22 µm polyvinylidene difluoride membrane (Millipore, Boston, MA, USA). The membranes were blocked with 5% non-fat milk in Tris-buffered saline containing 0.1% Tween-20 at room temperature for 2 hours, and incubated at 4°C overnight with each primary antibodies: Bax, Bcl-2, AKT, p-AKT, ERK, p-ERK, FOXO1, p-FOXO1, Nanog (1:1,000, Cell Signaling Technology, Beverly, MA, USA), MRP1, P-gp, LRP, GST-π (1:1,000, Abcam, San Francisco, CA, USA), and α-tubulin antibody (1:2,000, Beyotime). Then the membranes were incubated with the secondary antibody for 2 hours at room temperature. Enhanced chemiluminescence reagent (Beyotime) was used to detect the protein signals. All values were normalized to those of α-tubulin. All experiments were performed in triplicate.

Cell sensitivity to cisplatin

The viability of the SiHa cells after treatment with cisplatin (Sigma-Aldrich Co.) was determined using the CCK-8 assay (Cell Counting Kit-8; Dojindo Laboratories, Tokyo, Japan). The cells were digested and cultured in 96-well plates for 24 hours. Subsequently, the cells were treated with various concentrations of cisplatin (0, 1, 2.5, 5, 10, 20, 30, 40, 50, and 60 µM) for 24 or 48 hours. Then the drug solution was replaced with fresh medium, and 10 µL/well CCK-8 solution was added to the medium. The cells were incubated at 37°C for 2 hours, and absorbance measured at 450 nm absorption spectra in a microplate reader (Bio-Rad Laboratories Inc., Hercules, CA, USA). Cell viability was calculated as follows: % cell viability = (OD₄₅₀ of test well – OD₄₅₀ of blank well) / (OD₄₅₀ of control well – OD₄₅₀ of blank well) × 100%. The experiments were repeated three times. The half-maximal inhibitory concentration (IC₅₀) was defined as the concentration of cisplatin that inhibited cell viability by 50% which was calculated by GraphPad Prism software.

Apoptosis assay

Apoptosis was assessed using the annexin V-phycoerythrin (PE) and 7-amino-actinomycin D (7-AAD) apoptosis detection kit (BD, Franklin Lakes, NJ, USA) according to the manufacturer's instructions. After treatment with 10 µM cisplatin for 24 hours, SiHa cells were collected, washed with phosphate buffer saline twice and digested by 0.25% trypsin and dissociated into single cell. Then the cells were double-stained with 5 µL annexin V-PE and 7-AAD. Stained cells were analyzed by flow cytometry (BD). The experiment was repeated three times.

Plate clone formation assay

Four different kinds of cells were seeded into each well of 6-well plates at the density of 400/well and cultured in DMEM containing 10% fetal bovine serum for 14 days. After washing with PBS, each well was fixed with methyl alcohol for 15 minutes and stained with crystal violet for 10 minutes.

Statistical analysis

Statistical analysis was performed with Statistical Product and Service Solutions (SPSS) 19.0 statistical software (IBM Corporation, Armonk, NY, USA). All data values in the text were continuous variables and normal distribution, expressed as mean ± SD. The differences were analyzed using one-way ANOVA with least significance difference method. Statistically significant differences between two groups were determined by two-tailed unpaired Student's *t*-test. Differences were considered statistically significant at a 2-sided *P*-value of less than 0.05.

Results

Lentivirus-mediated upregulation and downregulation of S100A9 in SiHa cells

To elucidate the functional importance of S100A9, SiHa cells were transfected with S100A9-pLVX-IRES-ZsGreen1 lentivirus or S100A9-pGFP-B-shLenti to stably enhance or knockdown the expression of S100A9. The efficiency

of upregulation and downregulation of S100A9 is shown in Figures 1 and 2. The levels of S100A9 expression in the empty-vector group showed no significant difference compared to the control group ($P>0.05$)

As shown by Western blot analysis (Figure 2), transfection of SiHa cells with S100A9-pLVX-IRES-ZsGreen1 lentivirus dramatically enhanced the expression of S100A9 by ~10-fold. SiHa cells transfected with S100A9-pGFP-B-shLenti

showed the maximum suppression of S100A9 expression by approximately 3-fold.

Effect of S100A9 on the sensitivity of SiHa cells to cisplatin

The cell proliferation treated with cisplatin was further evaluated with CCK-8 assay. The IC₅₀ values were calculated from the cell viability plots. Results revealed that upregulation of

Figure 1 The transfection efficiency of lentivirus after transfecting SiHa cells for 48 hours.

Notes: (A) Image of SiHa cells at normal light ($\times 100$). (B) GFP expression in SiHa cells following transfection with S100A9-pLVX-IRES-ZsGreen1 lentivirus at fluorescent ($\times 100$). The transduction efficiency of lentivirus was 93%.

Figure 2 S100A9 upregulation and downregulation by S100A9-pLVX-IRES-ZsGreen one lentivirus and S100A9-pGFP-C-shLenti lentivirus.

Notes: (A and B) S100A9 protein expression in SiHa cells transfected with S100A9-pLVX-IRES-ZsGreen one lentivirus was detected by Western blot. (C and D) S100A9 protein expression in SiHa cells transfected with four various S100A9 shRNA. Data are mean $\pm$ SD from triplicate experiments. * $P<0.05$ compared with the control and negative control group (control-Lt or control shRNA group). ** $P<0.01$ compared with the control and negative control group.

Abbreviations: control-Lt, control lentivirus; S100A9-Lt, S100A9 lentivirus.

S100A9 decreased the sensitivity of SiHa cells to cisplatin compared with the control-Lt group, whether SiHa cells were treated for 24 hours or 48 hours (Figure 3A and C) and the IC₅₀ value was dramatically increased (Table 2). As illuminated by Figure 3B and D, compared to the negative control group, down-regulation of S100A9 led to decrease survival rate of SiHa cells with a significantly decreased IC₅₀ value to cisplatin (Table 3).

Influences of S100A9 expression on cell apoptosis caused by cisplatin

The underlying mechanisms of S100A9 effect on chemosensitivity were further studied. SiHa cells were treated with cis-

platin at 10 μ M concentration for 24 hours and flow cytometry was used to detect the apoptotic cells. S100A9 overexpression SiHa cells had significantly lower apoptotic rate and S100A9 knockdown SiHa cells had obviously higher apoptotic rate, compared with their negative control group respectively (Figure 4), indicating that S100A9 could significantly inhibit the cell apoptosis of SiHa cells caused by cisplatin.

S100A9 affected apoptosis related protein Bcl-2 expression in SiHa cells

As S100A9 protein could affect the apoptosis rate of SiHa cells induced by cisplatin, the expression level of apoptosis

Figure 3 Effect of S100A9 on the sensitivity of SiHa cells to cisplatin.

Notes: (A and B) S100A9 overexpression and knockdown SiHa cells treated with cisplatin for 24 hours. (C and D) S100A9 overexpression and knockdown SiHa cells treated with cisplatin for 48 hours. Data are mean $\pm$ SD from triplicate experiments. *Compared with control group, $P < 0.05$. #Compared with negative control group (control-Lt or control shRNA group) $P < 0.05$.

Abbreviations: control-Lt, control lentivirus; S100A9-Lt, S100A9 lentivirus.

Table 2 IC₅₀ values of cisplatin for S100A9-overexpressed SiHa cells

Cell line	Cisplatin IC ₅₀	
	24 hours	48 hours
Control	17.24	6.54
Control-Lt	16.78	6.97
S100A9-Lt	26.30	12.21

Abbreviations: control-Lt, control lentivirus; S100A9-Lt, S100A9 lentivirus.

Table 3 IC₅₀ values of cisplatin for S100A9 knockdown SiHa cells

Cell line	Cisplatin IC ₅₀	
	24 hours	48 hours
Control	17.89	6.85
Control-shRNA	17.21	7.49
S100A9-shRNA	11.22	3.91

related-proteins Bcl-2 and Bax in SiHa cells after the change of S100A9 protein was detected by Western blot. As shown in Figure 5, the Bcl-2 protein level was significantly enhanced after S100A9 upregulated by lentivirus in SiHa cells, and decreased after S100A9 downregulated. Therefore, S100A9 could influence Bcl-2 protein expression. However, the expression level of Bax protein was not affected by S100A9 expression.

S100A9 upregulated drug resistance-associated proteins expression in SiHa cells

Multiple drug resistance protein 1 (MRP1), P glycoprotein (P-gp), lung-related resistance protein (LRP) and glutathione-S-transferase- π (GST- π) are well-known drug resistance-related proteins in cancer cells. The expression levels of

Figure 4 The effect of S100A9 on the apoptosis of SiHa cells induced by cisplatin.

Notes: (A and B) The apoptosis percentage of blank vector group and S100A9 overexpression group. (C and D) The apoptosis percentage of control group and S100A9 knockdown group. Apoptosis percentage was analyzed by annexin V-PE/7-AAD staining. Data are mean $\pm$ SD from triplicate experiments. ** $P < 0.01$.

Abbreviations: control-Lt, control lentivirus; S100A9-Lt, S100A9 lentivirus; PE, phycoerythrin.

Figure 5 Levels of apoptosis-related proteins Bcl-2 and Bax in S100A9 overexpression and knockdown SiHa cells.

Notes: (A and C) Bcl-2 and Bax proteins expression was analyzed by Western blot. (B and D) The data are presented as relative expression level of Bcl-2 and Bax proteins normalized to tubulin. Data are mean $\pm$ SD from triplicate experiments. * $P < 0.05$.

Abbreviation: Bcl-2, B-cell lymphoma-2; control-Lt, control lentivirus; S100A9-Lt, S100A9 lentivirus.

MRP1, P-gp, GST- π and LRP in SiHa cells were detected by Western blot. As showed in Figure 6, P-gp protein level tended to increase compared to the control groups when S100A9 expression was upregulated. While S100A9 expression was downregulated, P-gp protein had a tendency to decrease. However, there was no significant difference. The expression of MRP1 protein was not affected by S100A9 expression.

Western blot was performed to evaluate the expression of LRP and GST- π protein, another two MDR-related proteins, in S100A9 upregulated and downregulated SiHa cells. Results revealed a remarkable increased LRP and GST- π protein expression in S100A9 overexpression SiHa cells. Meanwhile, LRP and GST- π protein expression significantly decreased in S100A9 knockdown SiHa cells (Figure 6).

S100A9 levels correlate with p-AKT, p-ERK, p-FOXO1, and Nanog protein expression in SiHa cells

To further identify the potential mechanism for regulation of tumor cell cisplatin chemosensitivity by S100A9, Western blot analysis was performed. We detected a striking increase of p-AKT, p-ERK, p-FOXO1, and Nanog in S100A9-overexpressed SiHa cells, and the opposite results were observed in the S100A9 knockdown SiHa cells (Figure 7). Moreover, FOXO1 was downregulated in S100A9-overexpressed group

and upregulated in S100A9 knockdown group (Figure 7). Furthermore, S100A9 overexpression remarkably increased the number of clones formed, whereas the knockdown of S100A9 inhibited their ability to form colonies, as assessed by the colony formation assay (Figure 8). These data suggest that upregulation of S100A9 could augment the tumorigenicity of SiHa cells and promote SiHa cells acquiring stemness.

Discussion

S100A9 is reported to be overexpressed in a variety of cancers¹⁶ and is implicated in chemotherapy resistance.^{17–22} The results of studies concerning the effect of S100A9 protein on chemotherapy are contradictory. Needle-biopsied tissues of breast cancer patients prior to neoadjuvant chemotherapy (doxorubicin and docetaxel) were collected, and mass spectrometry and immunohistochemistry found that S100A9 was robustly upregulated in the sensitive group.¹⁷ In an in vitro study, knocking down S100A9 could enhance human cervical squamous carcinoma cell SiHa and C-33A invasion ability and tumor cell viability against paclitaxel treatment.¹⁸ Conversely, other studies concluded that S100A9 could enhance the chemotherapy resistance of cancers. In NK/T cell lymphoma patients treated with pegaspargase/gemcitabine, higher levels of S100A9 protein were found in non-responders compared with responders through serum

Figure 6 Levels of drug resistance-related proteins MRP1, P-gp, LRP and GST-π in S100A9 overexpression and knockdown SiHa cells.

Notes: (A and C) MRP1, P-gp, LRP, and GST-π proteins expression was analyzed by Western blot. (B and D) The data are presented as relative expression level of MRP1, P-gp, LRP, and GST-π proteins normalized to tubulin. Data are mean $\pm$ SD from triplicate experiments. * $P < 0.05$ compared with the control and negative control group. ** $P < 0.01$ compared with the control and negative control group.

Abbreviations: MRP1, multiple drug resistance protein 1; P-gp, P glycoprotein; LRP, lung-related resistance protein; GST-π, glutathione-S-transferase-π; control-Lt, control lentivirus; S100A9-Lt, S100A9 lentivirus.

proteomic analysis.¹⁹ Benzyl butyl phthalate could increase the secretion of S100A9 in tumor myeloid-derived suppressor cells, and therefore exacerbate the breast cancer resistance to doxorubicin with cyclophosphamide.²⁰ In muscle invasive bladder cancer, S100A9 positivity predicted disease progression after chemotherapy.²¹ In epithelial ovarian cancer, S100A9 was also found upregulated in chemoresistant tumors using Affymetrix HGU133A microarray.²² Nevertheless, the relationship between S100A9 and cervical cancer cell sensitivity to cisplatin was still unclear. In the present study, the results revealed that upregulated S100A9 protein significantly increased the IC₅₀ values of cisplatin in SiHa cells, while inhibition of the S100A9 protein expression obviously lowered the IC₅₀ values of cisplatin. Therefore, we considered that S100A9 might relate to the cisplatin resistance of SiHa cells.

Cisplatin, also named cis-diamminedichloroplatinum (II), is a platinum coordination compound and a major chemotherapeutic drug for the treatment of cervical carcinomas. DNA is the cytotoxic target of cisplatin. Cisplatin damages DNA through forming covalent adducts with purine DNA bases and the drug–DNA adducts induce cell cycle arrest and apoptosis.²³ However, two problems in cisplatin treatment are that the sensitivity needs to be improved and the cellular toxicity also increases significantly along with the increase of dosage.²⁴ The molecular mechanism underlying cisplatin resistance is multifactorial, and mainly associated with the following factors: 1) less accumulation of intracellular platinum compounds; 2) enhanced ability of tumor cells to repair damaged DNA; 3) upregulation of antiapoptotic proteins; and 4) enhanced epithelial–mesenchymal transition.⁶ The precise mechanism of how S100A9 protein regulates cervical cancer

Figure 7 Levels of p-AKT, AKT, p-ERK, ERK, p-FOXO1, FOXO1, and Nanog in S100A9 overexpression and knockdown SiHa cells.

Notes: (A and C) p-AKT, AKT, p-ERK, ERK, p-FOXO1, FOXO1, and Nanog proteins expression was analyzed by Western blot. (B and D) The data were presented as relative expression level of p-AKT, AKT, p-ERK, ERK, p-FOXO1, FOXO1, and Nanog proteins normalized to tubulin. Data are mean $\pm$ SD from triplicate experiments. * $P < 0.05$ compared with the control and negative control group. ** $P < 0.01$ compared with the control and negative control group.

Abbreviations: ERK, extracellular regulated protein kinases; p-ERK, phosphorylated extracellular regulated protein kinases; FOXO, forkhead box protein O; p-FOXO, phosphorylated forkhead box protein O; control-Lt, control lentivirus; S100A9-Lt, S100A9 lentivirus.

Figure 8 The effects of S100A9 expression on colony formation.

Notes: (A) S100A9 overexpression, (B) S100A9 knockdown.

Abbreviations: Control-Lt, control lentivirus; S100A9-Lt, S100A9 lentivirus.

cell sensitivity to cisplatin is poorly understood up to now. In the current study, we provided evidence that S100A9 was related to the cervical cancer cells cisplatin resistance, and may be through the inhibition of apoptosis.

Apoptosis pathways involve multiple proteins, and the Bcl-2 family proteins, especially Bcl-2 and Bax, are the critical ones.⁶ In various human cancers, the decrease of Bcl-2 expression conferred cisplatin sensitivity.^{25–28} In cervical cancer, down-regulation of NDRG2 could enhance the sensitivity of HeLa cells to cisplatin treatment via inhibiting Bcl-2 expression.²⁵ microRNA 873 decreased the level of the Bcl-2 protein and enhanced sensitivity to cisplatin in the glioma cells.²⁶ In cervical cancer, suppression of N-Myc downstream regulated gene two could enhance sensitivity of HeLa cells to cisplatin through significantly inhibited Bcl-2 expression.²⁷ Knocking down phosphoglycerate dehydrogenase significantly increased cisplatin chemotherapy sensitivity of HeLa cells and was connected with the reduction of Bcl-2 protein.²⁸ In this study, the results indicated that overexpression of S100A9 raised the level of Bcl-2 protein, while downexpression of S100A9 inhibited the level of Bcl-2 protein, but Bax protein expression was not varied. Therefore, S100A9 influenced the sensitivity of SiHa cells to cisplatin maybe by means of regulating apoptosis via Bcl-2 protein but not Bax protein.

The ATP-binding cassette transporter proteins, including MRP1 and P-gp, are known to play a role in the phenomenon of multidrug resistance in cancer.²⁹ MRP1 and P-gp are transport proteins and commonly protect cells by effluxing chemotherapeutic agents.³⁰ The overexpression of MRP1 and P-gp in cancer cells results in reduction of intracellular

platinum compound concentration.³¹ Whether the effect of S100A9 on SiHa cells cisplatin sensitivity involved MRP1 and P-gp has not been reported yet. In the present study, results showed that the expression of MRP1 and P-gp protein was not altered by S100A9 expression. So, MRP1 and P-gp might not be involved in the S100A9 regulation of cisplatin sensitivity in SiHa cells.

LRP is another protein related to chemoresistance which does not belong to the ATP-binding cassette transporter family. LRP is encoded by the MVP gene which is on the short arm of chromosome 16 (16p11.2).³² Clinical trials showed that the expression of LRP was significantly correlated with the MDR of various tumors, and the expression level of LRP was positively correlated with multidrug resistance.³³ Several groups have demonstrated that ectopic expression of LRP in hematopoietic tumors such as multiple myeloma or acute leukemia may result in poor therapeutic effect.^{34,35} Our results found that LRP protein expression was downregulated in S100A9 knockdown SiHa cells, while LRP protein expression was upregulated in S100A9-overexpressed SiHa cells. This may represent one of the mechanisms to explain the decreased sensitivity of S100A9-overexpressed SiHa cells to cisplatin.

GST- π , a member of the GST family, is also a non-transporter-based MDR, which functions by altering the activity of enzyme systems, resulting in drug sequestration in intracellular vesicles.³⁶ As a pivotal multifunctional enzyme, GST- π plays an important role in cellular detoxification. A growing body of evidence showed that GST- π was overexpressed in many tumors compared to the surrounding normal tissues and in various drug-resistant cancer cell line.³⁷ More importantly, GST- π has been demonstrated as a drug resistance protein in gynecological cancers.³⁸ Recent studies have demonstrated that GST- π plays a key role in regulating the MAP kinase pathway which participates in fundamental cellular processes such as growth, proliferation, differentiation, migration and apoptosis.³⁹ Given our findings that S100A9 could regulate GST- π expression in SiHa cells, we suspect that S100A9 is likely implicated in the regulation of drug resistance in cervical cancer.

FOXO1 is a member of the Forkhead box (Fox) family. The evolutionarily conserved transcription factor is a subclass of the forkhead family, which consists of four members.⁴⁰ FOXO1 regulates diverse gene expression and controls a wide spectrum of cellular processes. Deregulation of FOXO1 has a crucial role in the development and progression of many types of cancer.⁴¹ FOXO1 exerts its antitumor effect by promoting apoptosis, inducing differentiation, and inhibiting tumorigenicity and self-renewal.⁴² FOXO1 is mainly regulated by a variety of post-translational modifications, including

phosphorylation, acetylation and ubiquitylation.⁴³ Under stress conditions, FOXO1 enters the nucleus to promote the expression of pro-apoptotic genes.⁴² Aberrant overactivation of PI3K/AKT and MEK/ERK signaling can phosphorylate and inhibit FOXO1.⁴⁴ After being phosphorylated by related proteins, FOXO1 is blocked in the cytoplasm and is degraded, inhibiting its transcriptional activity.⁴⁴ FOXO1 also plays an important role in maintaining cellular pluripotency, and can directly regulate the expression of Nanog, OCT4, and SOX2 in human embryonic stem cells.^{45,46} But, the role of FOXO1 in cervical cancer cells is unclear.

In an in vitro study reported by Zhu et al,¹⁸ the effects of S100A9 on the biological behavior of SiHa and C-33A human cervical squamous carcinoma cell were found to be related to the phosphatidylinositol 3-kinase/Akt signaling pathway. In the present study, the results showed that overexpression of S100A9 in SiHa cells could promote p-AKT, p-ERK, p-FOXO1 and Nanog expression. Our results suggest that ectopic S100A9 expression induced a significant upregulation of p-AKT and p-ERK protein expression in SiHa cells. Subsequently, p-AKT and p-ERK interacts with FOXO1 and promotes its phosphorylation. AKT and ERK induced FOXO1 phosphorylation promotes its nuclear exclusion, and subsequent degradation of FOXO1. Reduced FOXO1 expression loses the inhibitory effect on tumor stem cells and further promotes SiHa cells acquiring stem cell-like properties, which is also confirmed by a plate clone formation assay in the present study. Therefore, the expression of pluripotent markers, such as Nanog, increased. This makes SiHa cells resistant to cisplatin chemotherapy.

Taken together, we hypothesize that S100A9 reduces the sensitivity of cervical cancer SiHa cells to cisplatin by the following effects. First, S100A9 may change the expression of multidrug-resistant portion GST- π , LRP and apoptosis related protein Bcl-2 and affect the apoptosis process. Secondly, S100A9 can promote the phosphorylation of FOXO1 through AKT and ERK signaling pathway, which further inhibits FOXO1-mediated transcription of pro-apoptotic gene and promotes tumor cell survival. Finally, the phosphorylation of FOXO1 induced by S100A9 overexpression can promote the expression of pluripotent markers and promote SiHa cells to maintain a stem cell-like phenotype, consequently resistant to cisplatin chemotherapy. Downregulation of S100A9 could significantly increase apoptosis rate, resulting in enhanced sensitivity of SiHa cells to cisplatin, which may be related to Bcl-2, GST- π and LRP protein and the altered AKT/ERK-FOXO1-Nanog signaling pathway. The detailed mechanisms and precise signaling pathway need further study.

Acknowledgments

This work was sponsored by grants from the National Natural Science Foundation of China (no. 81372381) and Science and Technology project of Wenzhou City (no. H20150003). The study sponsors had no involvement in the collection, analysis and interpretation of data, or in the writing of the manuscript.

Disclosure

The authors report no conflicts of interest in this work.

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