Downregulation of DNA-PKcs suppresses P-gp expression via inhibition of the Akt/NF-κB pathway in CD133-positive osteosarcoma MG-63 cells

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Abstract. The development of chemoresistance is closely linked to the plateau of the survival rate in osteosarcoma (OS) patients. CD133-positive (CD133+) OS cells are known as cancer stem cells (CSCs) in OS and exhibit the characteristic of chemoresistance. In this study, CD133+ and CD133-negative (CD133−) MG-63 cells were isolated by magnetic activated cell sorting (MACS). We verified that CD133+ MG-63 cells were more resistant to cisplatin (CDDP) than CD133− MG-63 cells. DNA-dependent protein kinase catalytic subunit (DNA-PKcs) and P-glycoprotein (P-gp) were expressed at higher levels in the CD133+ MG-63 cells compared with those levels in the CD133− MG-63 cells, whereas downregulation of DNA-PKcs by small interfering RNA (siRNA) decreased chemoresistance to CDDP and P-gp expression at the mRNA and protein levels in these cells. This indicated that DNA-PKcs was correlated with P-gp expression in the CD133+ MG-63 cells. The Akt/NF-κB pathway was hyperactivated in the CD133+ MG-63 cells, whereas inhibition of the Akt/NF-κB pathway downregulated P-gp expression. In addition, downregulation of DNA-PKcs suppressed the activity of the Akt/NF-κB pathway. These results revealed that downregulation of DNA-PKcs could decrease P-gp expression via suppression of the Akt/NF-κB pathway in CD133+ MG-63 cells. Therefore, inhibition of DNA-PKcs decreases P-gp expression and sensitizes OS CSCs to chemotherapeutic agents in vitro, which needs to be further validated in vivo.

Introduction

Osteosarcoma (OS) is the most common primary solid tumor of bone in children and adolescence among various types of bone tumors (1). With the introduction of chemotherapy in the 1970’s, the 5-year survival rate after surgery has increased to 50-70% in patients without metastasis (2-4). However, there has been no further improvement during the last three decades in terms of the survival rate and it remains at 20-30% for patients with detectable metastasis (4,5). The development of chemoresistance in OS contributes to the plateau of the survival rate to a certain extent. It is necessary to investigate the mechanisms of OS drug resistance.

The cancer stem cell (CSC) model is one emerging model for the development of drug resistance in malignancies. CSCs markedly promote drug resistance in various cancers (6). It has been demonstrated that CD133-positive (CD133+) cells in OS exhibit CSC characteristics (7-9). However, the mechanisms of drug resistance in CD133+ OS cells need to be further elucidated.

DNA-dependent protein kinase catalytic subunit (DNA-PKcs) is a member of the large phosphatidylinositol 3-kinase (PI3K)-related kinase (PIKK) family. DNA-PKcs, along with accessory heterodimeric complexes, Ku70 and Ku80, are involved in DNA damage repair via non-homologous end joining (NHEJ). Our previous studies revealed that inhibition of DNA-PKcs sensitized OS cells to chemotherapeutic agents (10), indicating that DNA-PKcs plays a significant role in chemoresistance. Moreover, DNA-PKcs was found to be overexpressed in OS CSCs (11), which might be one of the causes of chemoresistance in OS.

P-glycoprotein (P-gp), a member of the ATP-binding cassette (ABC) transporters, is encoded by the ABCB1 gene and plays an important role in chemoresistance in tumors. Hence, it is necessary to understand the mechanisms of the regulation of P-gp. It has been demonstrated that P-gp expression is at a higher level in OS CSCs compared with that in non-CSCs (9). Although DNA-PKcs and P-gp are involved in chemoresistance and are overexpressed in OS CSCs, there has been no study concerning the relationship between DNA-PKcs and P-gp in OS CSCs to date. Previous studies have revealed that the expression of P-gp is regulated by the PI3K/Akt/NF-κB pathway.
pathway in other cancers (12,13). This prompted us to investigate the relationship between DNA-PKcs and P-gp in OS CSCs, as well as the role of the Akt/NF-κB pathway in this relationship.

We hypothesize that DNA-PKcs regulates P-gp via the Akt/NF-κB axis in CD133+ OS cells. The purpose of this study was to investigate the role of DNA-PKcs in P-gp expression and the underlying molecular mechanism in drug-resistant CD133+ MG-63 cells. Compared with CD133-negative (CD133-) MG-63 cells, CD133+ MG-63 cells showed increased expressions of DNA-PKcs and P-gp, as well as higher activity of the Akt/NF-κB pathway. Downregulation of DNA-PKcs significantly decreased the P-gp expression and activity of the Akt/NF-κB pathway, and inhibition of the Akt/NF-κB pathway downregulated the P-gp expression. All of these results revealed that DNA-PKcs regulates P-gp via the Akt/NF-κB pathway in CD133+ OS cells.

Materials and methods

Cell culture. The human MG-63 cell line was purchased from the American Type Culture Collection (ATCC; Manassas, VA, USA). Cells were cultured in Dulbecco’s minimal essential medium (DMEM) supplemented with 10% fetal bovine serum (both from Gibco, Grand Island, NY, USA) at 37°C, 5% CO₂ in a 95% humidified atmosphere.

Magnetic activated cell sorting (MACS). MACS was performed using CD133 MicroBead kit (Miltenyi Biotec, Auburn, CA, USA) following the manufacturer’s instructions. Briefly, a single-cell suspension was prepared in the MACS separation buffer. Cells were incubated with FcR Blocking Reagent and CD133 MicroBeads at 4°C for 15 min. After washing steps, magnetic separation was performed using an LS Column and MACS Separator (Miltenyi Biotec). The magnetically labeled CD133+ cells and unlabeled CD133- cells were collected, respectively.

Cell viability assay. Cells were seeded in 96-well plates at a density of 5,000 cells/well. Then cisplatin (CDDP; Qilu Pharmaceutical Co., Ltd., Shandong, China) was added at a density of 5,000 cells/well. Then cisplatin (CDDP; Qilu Pharmaceutical Co., Ltd., Shandong, China) was added at 24 h post-treatment using the Cell Counting Kit-8 (CCK-8; BestBio, Shanghai, China) according to the manufacturer’s instructions. The cell survival rate was presented as the percentage of viable cells compared with the corresponding viable control cells. The half maximal inhibitory concentration (IC₅₀) was calculated from the relative survival curve.

Transfection of small interfering RNA (siRNA) and inhibitor treatment. The CD133+ MG-63 cells were seeded into ultra-low attachment 6-well plates in serum-free medium. The serum-free medium consisted of Dulbecco’s modified Eagle’s medium (DMEM), 20 ng/ml epidermal growth factor (EGF), 20 ng/ml basic fibroblast growth factor (bFGF) (both from PeproTech, Rocky Hill, NJ, USA) and N-2 Supplement (Gibco). For siRNA transfection, the siDNA-PKcs, siNF-κB/p65 or control siRNAs (GenePharma Co., Ltd., Shanghai, China) were transfected into the cells using Lipofectamine 2000 (Invitrogen, Carlsbad, CA, USA) according to the manufacturer’s instructions. After 24 or 48 h of transfection, the cells were harvested for further experiments at the gene or protein level, respectively. For the inhibition experiment, the CD133+ MG-63 cells were treated with Akt inhibitor MK-2206 2HCl (10 µM) (Selleck, Houston, TX, USA) for 24 h, and subjected to gene and protein expression experiments.

Immunofluorescence. The cells were seeded on 24-well chamber slides. After adherence, the cells were fixed with 4% paraformaldehyde for 15 min. The fixed cells were incubated in 0.3% Triton X-100 for 10 min to permeabilize and 10% normal goat serum for 1 h to block non-specific protein-protein interactions. Then the cells were incubated with the rabbit polyclonal anti-human P-gp (ab129450, 1:200) and mouse monoclonal anti-human DNA-PKcs primary antibodies (ab1832, 1:100) (both from Abcam, Cambridge, MA, USA) overnight at 4°C. The Alexa Fluor 488 goat anti-rabbit (ZF-0511, 1:200) and Alexa Fluor 594 goat anti-mouse (ZF-0513, 1:200) (both from Beijing Zhongshan Golden Bridge Biotechnology Co., Ltd., Beijing, China) secondary antibodies were used for detection. 4’,6-Diamidino-2-phenylindole was used to stain the cell nuclei for 5 min at room temperature. Slides were observed on an inverted fluorescence microscope (BX52; Olympus Corp., Tokyo, Japan).

Quantitative real-time polymerase chain reaction (qPCR). Total RNA was extracted with TRIzol lysis buffer (Toyobo, Osaka, Japan) according to the manufacturer’s instructions. First-strand complementary DNA (cDNA) synthesis was carried out using ReverTra Ace qPCR RT kit (Toyobo). Briefly, 1 µg of total RNA was used in a total volume of 10 µl containing 2 µl 5X RT buffer, 0.5 µl RT Enzyme Mix, 0.5 µl Primer Mix and nuclelease-free water. The reverse transcription was performed in a thermal cycler (TGradient 96; Biometra GmbH, Göttingen, Germany) with a temperature cycling program of 15 min at 37°C, 5 min at 98°C. The cDNAs were used as templates for PCR amplification using SYBR® Green Realtime PCR Master Mix kit (Toyobo). In brief, reaction mixtures (20 µl) for PCR were assembled using 2 µl cDNA template, 6.4 µl distilled water, 10 µl 2X SYBR® Green Realtime PCR Master Mix, 0.8 µl forward primers (10 µM) and 0.8 µl reverse primers (10 µM). The cycle parameters were 95°C for 30 sec followed by 40 cycles at 95°C for 5 sec, 55°C for 10 sec and 72°C for 15 sec. The human GAPDH PCR product was used as an internal control. The results were standardized with the formula: ΔCT = CTtarget - CTcontrol and further converted to the fold of the target gene over the control gene (2^-ΔΔCT). The primer sequences of the genes used in this study are presented in Table I.

Western blot analysis. The cells were harvested and total proteins were extracted with RIPA lysis buffer, and the protein concentrations were quantified with the Enhanced BCA Protein Assay kit (both from Beyotime, Shanghai, China). Equal amounts of protein were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and electroblotted onto a polyvinylidene fluoride (PVDF) membrane. Non-specific sites were blocked with 5% non-fat milk in Tris-buffered saline and Tween-20 (TBST)
at room temperature. The PVDF membranes were incubated at 4°C overnight with primary antibodies. Primary antibodies included: rabbit polyclonal anti-human γH2AX (S139) (ab2893, 1:1,000), rabbit polyclonal anti-human DNA-PKcs (ab230, 1:2,000), rabbit polyclonal anti-human P-gp (ab129450, 1:1,000) antibodies from Abcam; rabbit polyclonal anti-human Akt (9272S, 1:1,000), rabbit polyclonal anti-human phospho-Akt (S473) (9271S, 1:1,000), mouse monoclonal anti-human phospho-IκB-α (9246S, 1:1,000), and rabbit monoclonal anti-human phospho-NF-κB/p65 (3033S, 1:1,000) antibodies from Cell Signaling Technology (Danvers, MA, USA); rabbit polyclonal anti-human NF-κB/p65 (10745-1-AP, 1:500) antibody from Proteintech (Chicago, IL, USA); mouse monoclonal anti-human β-actin (TA-90, 1:500) antibody from Beijing Zhongshan Golden Bridge Biotechnology Co., Ltd. After washing with TBST, membranes were incubated with goat anti-rabbit secondary antibody (ZB-2301, 1:5,000) or goat anti-mouse secondary antibody (ZB-2305, 1:5,000) (both from Beijing Zhongshan Golden Bridge Biotechnology Co., Ltd.) conjugated with horseradish peroxidase for 1.5 h at room temperature. Immunoreactive bands were detected by enhanced chemiluminescence substrate (EMD Millipore, Billerica, MA, USA).

Table I. Primer sequences for qPCR.

<table>
<thead>
<tr>
<th>Primers (5’-3’)</th>
<th>F</th>
<th>R</th>
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<tbody>
<tr>
<td>PRKDC</td>
<td>ACAGAGATCCAGAAAGTGAGACA</td>
<td>AGCAACCGGTCCAAGGTATT</td>
</tr>
<tr>
<td>ABCB1</td>
<td>ACAGAGGGGATGGTCAGTGT</td>
<td>TCACGGCCATAGCGAATGGT</td>
</tr>
<tr>
<td>GAPDH</td>
<td>CAGGAGCATTGCGTGATGAT</td>
<td>GAAGGCTGGGGGCTATT</td>
</tr>
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qPCR, quantitative real-time polymerase chain reaction; F, forward; R, reverse.

Statistical analysis. Each experiment was performed three times independently. Data are expressed as means ± standard deviation (SD). Student's t-test was used for comparisons. Differences were considered statistically significant if P<0.05. Statistical analysis was carried out using GraphPad Prism 5.0 (GraphPad Software, Inc., La Jolla, CA, USA).

Results

CD133+ MG-63 cells are more resistant to CDDP compared with CD133 MG-63 cells. After MACS, the CD133+ and CD133- cells were treated with CDDP at different concentrations for 24 h, respectively. The cell viability was measured and the result showed that the CD133+ cells were more resistant to CDDP (Fig. 1A). The IC50 value of the CD133+ cells was significantly higher than that of the CD133- cells (23.55 vs. 14.57 µM; P<0.05) (Fig. 1B). In addition, the expression of DNA double-strand break (DSB) marker γH2AX

Figure 1. Downregulation of DNA-PKcs sensitizes CD133+ MG-63 cells to CDDP. (A and B) CD133+ MG-63 cells were more resistant to CDDP and had a higher IC50 value than the CD133 MG-63 cells. (C) After CDDP (10 µM) treatment for 24 h, the expression of γH2AX (S139) in the CD133+ cells was much lower than that in the CD133- cells. (D and E) Downregulation of DNA-PKcs by siRNA decreases the resistance to CDDP and the IC50 value in the CD133+ MG-63 cells, and (F) increased the expression of γH2AX (S139) after CDDP (10 µM) treatment for 24 h. The results represent the mean ± SD of three independent experiments; *P<0.05. DNA-PKcs, DNA-dependent protein kinase catalytic subunit; CD133+, CD133-positive; CDDP, cisplatin; CD133-, CD133-negative; siRNA, small interfering RNA; SD, standard deviation.
Figure 2. DNA-PKcs and P-gp are markedly elevated in the CD133+ MG-63 cells compared with levels in the CD133- MG-63 cells. (A) The expression levels of PRKDC and ABCB1 genes were significantly higher in the CD133+ MG-63 cells than levels in the CD133- MG-63 cells. (B) Immunofluorescence and (C and D) western blot analysis showed higher levels of DNA-PKcs and P-gp in the CD133+ MG-63 cells compared with levels in the CD133- MG-63 cells. The results represent the mean ± SD of three independent experiments; *P<0.05, **P<0.01. DNA-PKcs, DNA-dependent protein kinase catalytic subunit (or PRKDC); P-gp, P-glycoprotein; CD133+, CD133-positive; CD133-, CD133-negative; SD, standard deviation.

Figure 3. Downregulation of DNA-PKcs decreases P-gp expression at the gene and protein levels. (A) The expression level of the ABCB1 gene assessed by qPCR was significantly reduced in the CD133+ MG-63 cells transfected with siDNA-PKcs. (B) Immunofluorescence and (C and D) western blot analysis showed that P-gp expression was decreased in the CD133+ MG-63 cells after transfection with siDNA-PKcs. The results represent the mean ± SD of three independent experiments; *P<0.05, **P<0.01. DNA-PKcs, DNA-dependent protein kinase catalytic subunit (or PRKDC); P-gp, P-glycoprotein; qPCR, quantitative real-time polymerase chain reaction; CD133+, CD133-positive; SD, standard deviation.
The expression of γH2AX (S139) in the CD133+ cells was lower than that in the CD133− cells after CDDP (10 µM) treatment for 24 h (Fig. 1C). Downregulation of DNA-PKcs sensitizes CD133+ MG-63 cells to CDDP. After transfection with siDNA-PKcs or siControl, the CD133+ MG-63 cells were treated with CDDP at different concentrations for 24 h. It was shown that the CD133+ MG-63 cells with transfection of siDNA-PKcs were more sensitive to CDDP compared with those levels in the CD133− MG-63 cells (Fig. 1D). The IC50 value of the siDNA-PKcs group was lower than that of the siControl group (12.83 vs. 22.67 µM; P<0.05) (Fig. 1E). The expression of γH2AX (S139) was markedly elevated in the CD133+ MG-63 cells with siDNA-PKcs transfection after CDDP (10 µM) treatment (Fig. 1F). The results revealed that downregulation of DNA-PKcs reduced the DNA damage repair and increased the sensitivity to CDDP in the CD133+ MG-63 cells.

DNA-PKcs is involved in the expression of P-gp. The expression level of DNA-PKcs and P-gp were first investigated in the CD133+ and CD133− MG-63 cells, respectively. The results of qPCR revealed that the PRKDC and ABCB1 genes were at higher levels in the CD133+ MG-63 cells (Fig. 2A). In addition, immunofluorescence and western blot analysis showed that the expression levels of DNA-PKcs and P-gp were significantly decreased in the CD133+ cells compared with levels in the CD133− cells (Fig. 2B-D).

Then the CD133+ MG-63 cells were transfected with siDNA-PKcs, and P-gp was examined at the gene and protein levels. It was shown that the ABCB1 gene (Fig. 3A) and P-gp (Fig. 3B-D) expression were significantly decreased following the downregulation of DNA-PKcs. Taken together, the results indicate that DNA-PKcs is involved in P-gp expression, and DNA-PKcs and P-gp are positively correlated with chemoresistance to CDDP in CD133+ MG-63 cells.

The Akt/NF-κB pathway is implicated in P-gp expression in CD133+ MG-63 cells. The expression levels of p-Akt (both S473 and T308) and p-NF-κB/p65 were examined in both the CD133+ and CD133− MG-63 cells. The results showed that p-Akt (S473), p-Akt (T308) and p-NF-κB/p65 were expressed at higher levels in the CD133+ MG-63 cells compared with levels in the CD133− MG-63 cells (Fig. 4A). Immunofluorescence showed that NF-κB/p65 was mainly located in the nuclei of the CD133+ MG-63 cells and in the cytoplasm of the CD133− MG-63 cells. (C and D) The expression levels of the ABCB1 gene and P-gp were significantly decreased after downregulation of NF-κB/p65. (D and E) The expression levels of p-NF-κB/p65 and P-gp, as well as the ABCB1 gene were downregulated following the inhibition of the activity of Akt by MK-2206 2HCl (10 µM). The results represent the mean ± SD of three independent experiments; *P<0.05, **P<0.01. P-gp, P-glycoprotein; CD133+, CD133-positive; CD133−, CD133-negative; SD, standard deviation.
localized in the nuclei of CD133+ MG-63 cells (Fig. 4B). These results indicate that CD133+ MG-63 cells display hyperactivation of the Akt/NF-κB pathway.

Moreover, the results showed that P-gp and ABCB1 gene expression levels were decreased following the downregulation of NF-κB/p65 via siNF-κB/p65 transfection (Fig. 4C and D). This suggests that NF-κB/p65 is involved in P-gp expression. Following Akt inhibitor MK-2206 2HCl (10 µM) treatment, the activity of NF-κB/p65 and expression levels of P-gp and ABCB1 genes in the CD133+ MG-63 cells were examined. It was shown that the expression levels of p-NF-κB/p65 and P-gp, as well as the ABCB1 gene were downregulated by inhibition of the activity of Akt (Fig. 4D and E).

The results above demonstrated that the Akt/NF-κB pathway is implicated in P-gp expression at the gene and protein levels.

**Downregulation of DNA-PKcs decreases the activation of the Akt/NF-κB pathway in CD133+ MG-63 cells.** The Akt/NF-κB pathway proteins were observed in the CD133+ MG-63 cells after siDNA-PKcs transfection. Inhibition of DNA-PKcs via siDNA-PKcs decreased the expression of p-Akt (S473), p-IκB-α, p-NF-κB/p65, as well as P-gp (Fig. 5). It is worth noting that the expression of p-Akt (T308) was consistent between the siDNA-PKcs and siControl group. These results revealed that downregulation of DNA-PKcs suppressed Akt/NF-κB pathway activation and P-gp expression in the CD133+ MG-63 cells.

Taken together, all the results above revealed that downregulation of DNA-PKcs decreased P-gp expression via suppression of the Akt/NF-κB pathway in the CD133+ MG-63 cells.

**Discussion**

It is well known that a tumor is populated by heterogeneous cell populations and drug-resistant clones exist within the tumor (14,15). Targeting drug-resistant cells could have significance in the treatment of OS. The CSC theory believes that CSCs are relatively resistant to chemotherapeutic agents. Therefore, if it was possible to target drug-resistant CSCs, this would improve the therapeutic outcomes of OS. It has been well established that CD133+ cells in OS display features of CSCs (7-9), thus CD133+ MG-63 cells were taken as the object of this study.

Some chemotherapeutic reagents lead to DNA DSBs which are lethal for tumor cells. However, DSBs can be repaired by two main pathways, homologous recombination and NHEJ. DNA-PKcs, along with Ku70 and Ku80, are essential in DNA damage repair via NHEJ. Overexpression of DNA-PKcs is found in various malignancies, which is associated with poor prognosis (16,17). However, inhibition of DNA-PKcs sensitizes cells to chemotherapy in various tumor cells including OS (10,18,19). This indicates that DNA-PKcs is correlated with chemoresistance in tumors. Studies have revealed that DNA-PKcs is overexpressed in CSCs (11,20). In this study, we found that CD133+ MG-63 cells displayed overexpression of DNA-PKcs and chemoresistance to CDDP, along with lower expression of γH2AX (5139) after CDDP treatment, whereas downregulation of DNA-PKcs increased DSBs after CDDP treatment and sensitivity to CDDP. This demonstrates that DNA-PKcs overexpression leads to enhanced DNA damage repair and is involved in increased chemoresistance in OS CSCs.

The ABC family of drug transporters contributes to resistance to chemotherapeutic agents when overexpressed in tumors. P-gp is a well-characterized member of the ABC membrane transporters which functions as a drug efflux pump and reduces intracellular drug concentrations (21). Increased expression of P-gp is one of the key causes of drug resistance in tumors. Studies have reported that inhibition of P-gp reversed drug resistance in OS (22-24). Previous data revealed that high expression of P-gp is present in OS CSCs and is considered as one of the mechanisms of drug resistance in OS (25,26).

The results of this study showed that DNA-PKcs and P-gp were markedly elevated in CD133+ MG-63 cells, which may explain the chemoresistance of these cells with a higher IC50. In contrast, as the expression of DNA-PKcs was downregulated by siRNA, P-gp and ABCB1 gene expression levels were significantly decreased. This indicates that, besides DNA damage repair, DNA-PKcs is involved in chemoresistance via the regulation of P-gp expression. Therefore, the molecular mechanism through which DNA-PKcs mediates P-gp expression needs to be further investigated.

The PI3K/Akt signaling pathway is an important mediator of cell growth, survival and motility. Dysregulation of
PI3K/Akt pathway is implicated in resistance to chemotherapy in a wide variety of neoplasias (27-29). Activated Akt targets many proteins, including IκB kinase (IKK) which is responsible for the phosphorylation and degradation of IκB. Then NF-κB is released from the IκB-bound complex. With NF-κB nuclear translocation and binding to its recognition sites, ABCB1 gene promoter activation is enhanced and gene expression is induced (13,30). It has been demonstrated that downregulation of NF-κB inhibits P-gp expression by blocking ABCB1 gene transcription (31-33). These findings suggest that the Akt/NF-κB pathway may be able to mediate P-gp expression. Moreover, DNA-PKcs is a member of the PIKK family and is involved in Akt/NF-κB pathway activation (18). Therefore, we postulate that DNA-PKcs is involved in P-gp expression via the Akt/NF-κB pathway in CD133+ MG-63 cells.

To verify our hypothesis, the relationship between the Akt/NF-κB pathway and P-gp expression was first investigated. Our results revealed that p-Akt and p-NF-κB/p65 were highly expressed and NF-κB/p65 was mainly localized in the nuclei in the CD133+ cells compared with the CD133- cells, which indicated that the Akt/NF-κB pathway was activated in these cells. The results are consistent with previous reports (34,35). However, inhibition of the Akt/NF-κB pathway via inhibition of Akt activity or downregulation of NF-κB/p65 decreased P-gp and ABCB1 gene expression. These results demonstrated that the Akt/NF-κB pathway was involved in P-gp expression in the CD133+ MG-63 cells.

Since P-gp expression was decreased following downregulation of DNA-PKcs in the CD133+ MG-63 cells, we downregulated the DNA-PKcs expression via siRNA and examined the activation of the Akt/NF-κB pathway and P-gp expression. The results showed that p-Akt (S473), p-IκB-α, p-NF-κB/p65, P-gp and ABCB1 gene were decreased after transfection of siDNA-PKcs in the CD133+ MG-63 cells, which demonstrated that downregulation of DNA-PKcs decreased P-gp expression at the mRNA and protein levels via suppression of the Akt/NF-κB pathway in these cells. The results were supported by a previous report that DNA-PKcs mediated Akt/NF-κB pathway activation followed by the expression of P-gp in multidrug-resistant glioblastoma cells (36).

In addition, we found that disruption of DNA-PKcs decreased the expression of p-Akt (S473) rather than p-Akt (T308). This suggests that, as a member of the PIKK family, DNA-PKcs phosphorylates Akt at Ser473 specifically, which is consistent with previous data reported in other studies (37-39). This can be attributed to the fact that Ser473 is located in a hydrophobic motif Phe-Xaa-Xaa-Phe-Ser-Tyr (Xaa is any amino acid) in the C terminus, and DNA-PKcs has predisposition for phosphorylation sites at the extreme terminus of its substrate, which is critical for DNA-PK activity (38,40).

CD133+ OS cells which are well-known as CSCs play important role in drug resistance. Our study presented evidence that DNA-PKcs and P-gp were significantly elevated and the Akt/NF-κB pathway was activated in the CD133+ MG-63 cells. Moreover, downregulation of DNA-PKcs decreased P-gp expression and chemoresistance to CDDP via suppression of the Akt/NF-κB pathway in these cells. We therefore propose that combining DNA-PKcs inhibition targeting CSCs with conventional chemotherapeutic agents may be considered as a strategy to improve the treatment outcome of OS.

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References


