Pretreatment with the γ-secretase inhibitor DAPT sensitizes drug-resistant ovarian cancer cells to cisplatin by downregulation of Notch signaling

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Abstract. Notch signaling is implicated in ovarian cancer tumorigenesis and inhibition of Notch signaling with γ-secretase inhibitor DAPT resulted in reduction of tumor cell viability and induction of apoptosis in ovarian cancer cells. This study investigated whether DAPT has the same effect on ovarian cancer cells that are resistant to cisplatin and the underlying molecular events. Ovarian cancer cell lines resistant to cisplatin were treated with DAPT, cisplatin or combination for cell viability MTT, flow cytometric cell cycle, ELISA apoptosis and colony formation assays. qRT-PCR and western blotting were used to detect gene expressions. We found that pretreatment of ovarian cancer cisplatin-resistant cell lines with DAPT for 24 h and then with cisplatin for 72 h showed a synergistic antitumor activity in these cell lines, while cisplatin treatment and then addition of DAPT just showed an additive or antagonistic effects on these cisplatin-resistant ovarian cancer cells. Moreover, pretreatment of ovarian cancer cell lines with DAPT and then with cisplatin also inhibited tumor cell colony formation capacity, arrested tumor cells at G2 phase of the cell cycle and induced apoptosis. The cell cycle and apoptosis-related genes, such as cyclin B1, Bcl-2 and caspase-3, were also modulated by the treatment. Pretreatment of ovarian cancer cell lines with DAPT and then with cisplatin downregulated Notch1 and Hes1 expression dose- and time-dependently. The current data demonstrate that DAPT pretreatment was able to sensitize cisplatin-resistant human ovarian cancer cells to cisplatin by downregulation of Notch signaling.

Introduction

Ovarian cancer is the second most common gynecologic cancer in women and the leading cause of cancer death of gynecologic malignancy in the world (1). More than 90% ovarian cancers are believed to arise from the surface epithelium of the ovary and frequently absent of early signs and symptoms; thus most ovarian cancer patients are diagnosed at advanced stage of disease, which makes curable surgery infrequent and has a relatively poor prognosis. For past decades, chemotherapy has been a general standard of care for ovarian cancer with highly variable protocols and used alone or after surgery to treat any residual disease (2). Cisplatin-based chemotherapy is used as a primary treatment of ovarian cancer and its combination with other agents has become standard chemotherapy for treatment of advanced ovarian cancer, but prolonged drug treatment results in development of acquired drug resistance impeding successful treatment (3). The antineoplastic effect of cisplatin is mediated by formation of DNA adducts and inter- and intra-strand crosslinks (4). These adducts distort the DNA template with deceleration of cells in S phase followed with G2 phase arrest (5) and also result in diverse effects, including DNA synthesis inhibition, RNA transcription suppression, cell cycle arrest and apoptosis. Cisplatin resistance is multifactorial and rather complicated, including reduced platinum accumulation and enhanced platinum detoxification and metabolism in cells, altered DNA damage repair, activation of phospholipid kinase and phosphatidyl inositol 3-kinase and other signaling pathways ultimately causing dysregulation of apoptotic pathway (6). Thus, this disappointing outcome strongly suggests that a better understanding of the mechanisms of chemoresistance could lead to novel therapeutic strategies for effective control of ovarian cancer.

Our study is focusing on the Notch signaling. A previous study demonstrated that activation of the Notch signaling was linked to chemoresistance of pancreatic cancer to cisplatin (7) and inhibition of Notch signaling with γ-secretase inhibitors could sensitize colon cancer cells to chemotherapy and was synergistic with some antineoplastic agents (8). Indeed, the Notch signaling pathway plays a key role in the proliferation and differentiation of many tissues. This evolutionarily...
conserved pathway can regulate critical cell fate decision (9). In mammals, the Notch family consists of four receptors (i.e., Notch1 to Notch4) and five ligands (Jagged-1, Jagged-2, Delta-like-1, Delta-like-3 and Delta-like-4) (10). Notch ligands and receptors are type I membrane proteins that regulate cell fate during cell-cell contact (10-12). Receptor-ligand interaction between two neighboring cells leads to γ-secretase-mediated proteolytic release of the Notch intracellular domain (NICD) (13). NICD then translocates into the nucleus, and in turn interacts with the transcriptional cofactor CBF1 and transactivates target genes, such as Hes and Hey families to affect numerous pathways involving cell-fate determination (14,15). Abnormal Notch signaling has been documented in many cancers, including ovarian cancer (16,17). Overexpression of Notch proteins was associated with poor prognosis of different cancer patients (18) and with tumor de-differentiation in ovarian cancer (19). Molecularly, activation of Notch proteins are triggered by γ-secretase, which cleaves the Notch receptor to activate the pathway (13). Previous studies showed that γ-secretase inhibitors were able to inhibit tumor cell viability and induced apoptosis in different cancer cell lines (20,21). We found that γ-secretase inhibitor (DAPT)-blocked Notch signaling reduced viability of ovarian cancer A2780 cells but induced them to undergo apoptosis (22). In this study, we hypothesized that inhibition of Notch signaling by DAPT could sensitize drug-resistant ovarian cancer cells to cisplatin chemotherapy. We assessed the effects of DAPT on sensitizing cisplatin-resistant ovarian cancer A2780/CP70 and OV2008/C13 cells to cisplatin-induced cell death and the underlying molecular events.

Materials and methods

Cell culture and reagents. Two pairs of cisplatin-sensitive and cisplatin-resistant human ovarian cancer cell lines A2780, A2780/CP70 and OV2008, OV2008/C13 were kindly provided by Dr Jun Hu (The Third Military Medical University, Chongqing, China) and maintained in RPMI-1640 medium containing 10% heat-inactivated fetal bovine serum (FBS; all from Xin Xing Tang Biotechnology Company, Beijing, China) at 37˚C in a humidified 5% CO₂ atmosphere. For cisplatin treatment, cells were maintained in medium with the desired doses of cisplatin for 1 h and then washed with PBS and followed by incubation in fresh drug-free medium for varying times post-treatment.

Cell viability MTT assay. Cells (5x10⁴) were seeded in 96-well cell culture plates, treated with different concentrations of γ-secretase inhibitor N-[N-(3,5-difluorophenacetyl)-L-alanyl]-S-phenylglycine t-butyl ester (DAPT), cisplatin (all from Sigma-Aldrich, St. Louis, MO, USA), or combination for 72 h. For combination experiments, ovarian cancer A2780/CP70 and OV2008/C13 cells were treated for 72 h with DAPT (30 µmol/l) or cisplatin (3 or 6 µmol/l) treatment and then subjected to the MTT assay. Specifically, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) was added to the cultures and the cells were incubated for an additional 4 h. The resulting formazan crystals were solubilized by addition of 150 µl dimethyl sulfoxide (DMSO) to each well. The optical density was measured at 570 nm using a microplate reader and cell viability was determined by the formula: cell viability (%) = (absorbance of the treated wells - absorbance of the blank control wells) / (absorbance of the negative control wells - absorbance of the blank control wells) x 100%. All experiments were performed in triplicate and repeated at least three times. Drug interactions and isobologram were analyzed using CalcuSyn software (Biosoft, Beijing, China).

Colony formation assay. Ovarian cancer A2780/CP70 and OV2008/C13 cells (5x10⁴/ml) were seeded into 6-well plates according to the manufacturer's instructions. Cells were treated with DAPT (30 µmol/l) either 24 h before or after cisplatin (3 or 6 µmol/l) treatment. After 14 days of incubation at 37˚C in a humidified atmosphere containing 5% CO₂ in air, colonies were counted using an inverted microscope (Leica, Heidelberg, Germany).

Flow cytometric cell cycle assay. The cell cycle was analyzed using flow cytometry. Briefly, cells (1x10⁶) were collected and washed in phosphate-buffered saline (PBS), then fixed in 75% ice-cold alcohol for 30 min at 4˚C. After washing with ice-cold PBS three times, cells were resuspended in 1 ml of PBS containing 40 µg of propidium iodide (Sigma-Aldrich) and 100 µg of RNase A (Sigma-Aldrich) and incubated for 30 min at 37˚C. Samples were then analyzed by FACS (BD Immunocytometry Systems, San Jose, CA, USA). Each experiment was repeated for at least three times.

ELISA apoptosis assay. A Cell Death Detection ELISA kit (Roche, Shanghai, China) was used to detect apoptosis in treated cells according to the protocol provided by the manufacturer. Briefly, cell culture supernatants were washed away to remove fragmented DNA from necrotic cells, and then cells were lysed and loaded into microtiter plate modules coated with an anti-histone antibody for incubation for 45 min at room temperature. Next, samples were incubated with the anti-DNA peroxidase followed by color development with ABTS substrate. After that, the absorbance rates of these samples were measured using a microplate reader (SLT, Spectra LabInstruments Deutschland GmbH) at 405 and 490 nm (reference wavelength).

RNA isolation and quantitative RT-PCR. Total RNA was isolated from ovarian cancer cells using the TRIzol reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer's instructions and then reverse transcribed into cDNA using a PrimeScript™ RT reagent kit (Takara, Dalian, China). These cDNA samples were then amplified in the ABI 7500 system (Applied Biosystems) using SYBR Premix Ex Taq™ II (Takara). Primers used for Hes1 were 5'-TGGAAATCG-3' and 5'-CTAAGTCATAGGAC-3' and 5'-TCGTTCATGCATGACTGACAGTGAAGCACCTC-3' and 5'-TCGTTCATGCACTC-3'. The internal control β-actin primers were 5'-TGGCAACCCAGCACAATGGA-3' and 5'-CTAAGTCATGACTGACAGTGAAGCACCTC-3'. The internal control β-actin primers were 5'-TGGCAACCCAGCACAATGGA-3' and 5'-CTAAGTCATGACTGACAGTGAAGCACCTC-3'. Thermocycling was set as follows: 94˚C for 5 min, 40 cycles of 94˚C for 30 min, 55˚C for 30 min and 72˚C for 60 min, and a final extension at 72˚C for 10 min and then permanently stored at 4˚C. Relative quantitation of mRNA expression levels was determined using the relative standard curve method according to the manufacturer's instructions (Applied Biosystems).
Protein extraction and western blot analysis. Total proteins from ovarian cancer A2780/CP70 and OV2008/C13 cells were lysed in a lysis buffer containing NaCl, sodium desoxycholate, sodium dodecyl sulfate and Tris and incubated at 4˚C for 15 min. The protein concentrations were determined using the Bio-Rad assay system (Bio-Rad, Hercules, CA, USA). These protein samples were then fractionated using sodium dodecyl sulfate polyacrylamide (10%) gels for electrophoresis (SDS-PAGE) and then transferred onto a nitrocellulose membrane (Kenker, USA). For western blotting, the membranes were blocked in 5% non-fat milk in Tris-buffered saline containing 0.1% Tween-20 (TBS-T) and then incubated with appropriate primary antibodies overnight at 4˚C. Horseradish peroxidase-conjugated anti-goat IgG was used as the secondary antibody, and the protein bands were visualized using the enhanced chemiluminescence (ECL) method (GE Healthcare, USA) and quantified by using laser densitometry. The data were summarized as the mean of 3 independent experiments with the standard deviation. The membranes were then stripped by incubated for 30 min at 50°C in a buffer that contained 2% SDS, 62.5 mmol/l Tris (pH 6.7), and 100 mmol/l 2-mercaptoethanol and further washed and incubated with the desired primary antibody. Antibodies against Hes1, cyclin B1, caspase-3, Bcl-2, and β-actin were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA).

Statistical analysis. The data were expressed as means ± standard error. The Student's t-test was performed to analyze the data between groups. A P-value <0.05 was considered as statistically significant.

Results

DAPT potentiates cisplatin-reduced viability of ovarian cancer cells in a drug sequence-dependent manner. In this study, we first determined whether γ-secretase inhibitor DAPT was able to sensitize ovarian cancer cells to low-dose cisplatin-reduced cell viability. We treated A2780/CP70 and OV2008/C13 cells with various concentrations of DAPT up to 90 µmol/l and cisplatin up to 9 µmol/l under two different drug administration scenarios, i.e., an initial 24-h DAPT exposure followed by 72-h cisplatin treatment or vice versa. Cell viability was assessed and the values of inhibiting concentration (IC₅₀) were calculated and processed using CalcuSyn software. As shown in Fig. 1A and D, if DAPT was administered before cisplatin, DAPT can synergistically sensitize cisplatin anti-tumor activity in cisplatin-resistant ovarian cancer cell lines, i.e., most of IC₅₀ were below the combination-isobol line. On the other hand, if DAPT was administered after cisplatin, DAPT was additive or antagonistic rather than synergistic effects with cisplatin, i.e., most of IC₅₀ were above the combination-isobol line (Fig. 1B and E). After that, we selected two cisplatin doses (3 and 6 µmol/l) that kill ~20-40% of ovarian cancer cells and combined them with 30 µmol/l DAPT. Data showed that pretreatment of tumor cells with DAPT increased the potency of cisplatin-reduced cell viability, e.g., treatment of tumor cells with 3 and 6 µmol/l cisplatin for 72 h caused ~21 and...
38% reduction of viability of A2780/CP70 cells, respectively. In contrast, tumor cell viability was reduced to 51 and 64% when pretreated with DAPT (Fig. 1C and F and Table I), whereas tumor cell viability was only reduced to 37 and 42% for 3 and 6 µmol/l cisplatin, respectively, if cisplatin was administrated before DAPT (Table II).

Furthermore, we also determined the effects of their combination in regulation of tumor cell colony formation capacity. Our data showed that pretreatment of tumor cells with DAPT increased the potency of cisplatin-reduced colony formation in vitro (Fig. 2).

Combination of DAPT with cisplatin arrests cisplatin-resistant ovarian cancer cells in G2 phase of cell cycle. We next assessed the changed cell cycle after their treatment. In untreated control cells, the percentage of cells in G1, S, and G2 phases were 65.42, 22.73 and 11.85%, respectively, while 3 µmol/l cisplatin treatment had no significant effect on changes in the cell cycle distributions, whereas 30 µmol/l DAPT alone caused cell cycle redistribution to G2 phase. In contrast, DAPT-potentiated cisplatin treatment at the above named dose resulted in a pronounced G2 arrest (% of G1, S, and G2 phase cells was 18.90, 21.85 and 59.25%, respectively). We also found a high proportion of sub-G1 phase population (apoptotic cells) in the DAPT-cisplatin treated tumor cells (Fig. 3A and B). Molecularly, DAPT-cisplatin treatment reduced the levels of cyclin B1 protein (Fig. 3C).

DAPT enhances cisplatin-induced apoptosis in ovarian cancer cells. Since DAPT combination with low dose of cisplatin had a high proportion of sub-G1 phase population, we determined apoptosis levels in A2780/CP70 cells. As shown in Fig. 4A, treatment of A2780/CP70 cells with 30 µmol/l DAPT or 3 or 6 µmol/l cisplatin caused negligible increase in tumor cell apoptosis over the background. However, pretreatment of tumor cells with DAPT for 24 h and then with cisplatin for 72 h caused a significant increase in apoptosis (Fig. 4A, columns 5 and 6), whereas cisplatin treatment before DAPT addition at the same dose did not show a significant increase in cell death (Fig. 4A, columns 7 and 8). These findings further confirmed that the combination of DAPT with low dose of cisplatin resulted in induction of apoptosis in A2780/CP70 cells in a drug sequence-dependent manner. At the molecular level, expression of apoptosis-related genes, such as caspase-3 and Bcl-2 proteins was also altered, i.e., caspase-3 was significantly higher and Bcl-2 was lower in DAPT before cisplatin treatment (Fig. 4B, lanes 5 and 6), which were in accordance with the cell death data in Fig. 4A.

Table I. Reduced cell viability (%) by DAPT pretreatment and then cisplatin addition.

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Bold text indicates a significant change after treatment.

Table II. Reduced cell viability (%) by cisplatin pretreatment and then DAPT addition.

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Bold text indicates a significant change after treatment.
DAPT treatment downregulated Notch signaling and its target gene Hes1 expression in ovarian cancer cells. We investigated potential targets of DAPT in ovarian cancer cells. qRT-PCR data revealed that different concentrations of DAPT (30 or 45 µmol/l) resulted in significant downregulation of Notch1 mRNA (Fig. 5A and B) and western blot analysis showed that levels of Notch1 protein were also downregulated by DAPT treatment in A2780/CP70 and OV2008/C13 cells (Fig. 5A and B). These findings indicated that the Notch1 signaling pathway was efficiently suppressed by DAPT treatment in a dose-dependent manner. The concentration of 30 µM was therefore used subsequently to effectively inhibit the Notch1 pathway.

To further determine if DAPT could downregulate expression of Notch1 downstream gene Hes1, we performed qRT-PCR and western blot analyses and found that different concentrations of DAPT (30 and 45 µmol/l) significantly inhibited levels of Hes1 mRNA and protein (Fig. 5C and D). These findings indicated that DAPT treatment downregulated expression of Hes1 dose-dependently.

Moreover, we also performed time course treatments for changed expression of these genes. Our data showed that the altered expression of Notch1 was observed as early as 6 h after DAPT (30 µmol/l) treatment and was more pronounced with a longer period of treatment in A2780/CP70 and OV2008/C13 cells (Fig. 5E and F). These data suggested that the Notch1 signaling pathway was efficiently blocked by DAPT treatment in a time-dependent manner. Similarly, the altered expression of Hes1 gene was observed as early as 6 h after DAPT (30 µmol/l) treatment and was more pronounced with a longer period of treatment in A2780/CP70 and OV2008/C13 cells (Fig. 5G and H).
Figure 3. Effects of DAPT and cisplatin treatment on regulation of ovarian cancer cell cycle distribution. (A and B) A2780/CP70 cells were treated with 30 µmol/l DAPT for 24 h, followed by treatment with 3 µmol/l cisplatin for another 72 h. The cell cycle distribution was analyzed by using flow cytometry. *P<0.05, **P<0.01 compared to the single treatment groups by the Tukey-Kramer Multiple Comparison test (n=3). (C) Western blotting. The duplicated cells were subjected to western blot analysis of cyclin B1 expression.

Figure 4. Effects of DAPT and cisplatin treatment on regulation of ovarian cancer cell apoptosis. (A) Cell Death ELISA. A2780/CP70 cells were treated with DAPT, cisplatin, or combinations and then subjected to ELISA analysis of apoptosis. **P<0.001 compared to the corresponding single treatment groups using the Tukey-Kramer Multiple Comparison test (n=3). (B) Western blotting. The duplicated cells were subjected to western blot analysis of Bcl-2 and caspase-3 expression.
Discussion

Notch signaling is implicated in ovarian cancer tumorigenesis (19,23). Our previous studies showed that inhibition of Notch signaling with γ-secretase inhibitor DAPT resulted in reduced tumor cell viability and induction of apoptosis in ovarian cancer cells (22). In this study, we assessed whether DAPT has the same effect on ovarian cancer cells that are resistant to cisplatin and the underlying molecular events. We found that pretreatment of ovarian cancer cell lines with DAPT and then with cisplatin can synergistically sensitize cisplatin antitumor activity in cisplatin-resistant ovarian cancer cell lines, while cisplatin treatment with a delayed DAPT treatment had an additive or antagonistic effects on these cisplatin-resistant ovarian cancer cells. Similarly, these treatments also inhibited tumor cell colony formation capacity, arrested tumor cells at G2 phase of the cell cycle, but induced apoptosis. Expression of the cell cycle-related gene cyclin B1 and apoptosis-related
gene Bcl-2 was suppressed but apoptosis-related gene caspase-3 was activated by these treatments. Molecularly, pretreatment of ovarian cancer cell lines with DAPT and then with cisplatin downregulated Notch1 and Hes1 expression dose- and time-dependently. This study demonstrated that DAPT pretreatment could sensitize cisplatin-resistant human ovarian cancer cells to cisplatin by downregulation of Notch signaling. Future in vivo study need to confirm our current data before translating into clinical trials.

To date, cisplatin is still widely used in the treatment of various human cancers, including testicular, ovarian, cervical, bladder, head and neck, esophageal and lung cancers (24). In spite of the efficacy of cisplatin-based treatment regimens, long-term cure is difficult to obtain due to drug resistance (3), although great efforts have been made to develop combining chemotherapeutic agents to potentiate the effectiveness of current cytostatic drugs and to overcome chemotherapy resistance (8). Thus, our present study could provide a novel strategy by using combined agents to treat ovarian cancer. Our data showed pretreatment of ovarian cancer cell lines with DAPT and then with cisplatin synergistically sensitized cisplatin antitumor activity in cisplatin-resistant ovarian cancer cell lines. Our treatment regime could reduce the toxic dose of cisplatin but achieved synergistic effects on ovarian cancer cells.

Indeed, γ-secretase is a critical protease for Notch protein activation via nicastrin ectodomain binding to the N-terminus of Notch protein and cleavage of NICD (25). Thus, γ-secretase inhibitors can prevent generation of the intracellular domain of Notch protein and suppress the Notch activity (26). Notch signaling is implicated in ovarian cancer tumorigenesis (19,23). Recently, there has been increased enthusiasm in targeting this pathway using γ-secretase inhibitors for novel and effective cancer therapy strategy (27). For example, treatment of leukemia using this strategy revealed a better efficacy but less side effects of γ-secretase inhibitors (28). A γ-secretase inhibitor GSI could be helpful in treating human T-cell acute lymphoblastic leukemia (T-ALL) by inhibiting the Notch signaling (29) and combination therapy of γ-secretase inhibitors with glucocorticoids improved the anti-leukemic effects of γ-secretase inhibitors and reduced their gut toxicity in vivo (30). It has also been reported that combination of γ-secretase inhibitors with chemotherapy might represent a novel approach for treating metastatic colon cancers (8). In the present study, we investigated the potential use of DAPT to sensitize the cisplatin-resistant ovarian cancer A2780/CP70 and OV2008/C15 cells to cisplatin. We found that DAPT pretreatment reduced cisplatin-resistant ovarian cancer cells to low-dose cisplatin-induced cell death by downregulation of the Notch signaling.

Previous studies reported that DAPT had anti-proliferative activity against several human cancer cell lines (22,31). This anti-proliferative effect was due to induction of apoptosis and cell cycle arrest. Molecularly, Bcl-2 and caspase-3 play an important role in regulation of apoptosis (32,33) and cyclin B1 is one of the key molecules in the cell cycle modulation (34). Thus, our present study revealed an accumulation of G2 cell cycle arrest after DAPT treatment and DAPT-cisplatin treatment in cisplatin-resistant human ovarian cancer A2780/CP70 cells. As G2 arrest is typically linked to DNA damage response, we postulate that DAPT pretreatment-sensitized A2780/CP70 cells to respond to DNA damage. Moreover, overexpression of caspase-3 could sensitize breast cancer cells to drug-induced apoptosis and enhance chemosensitivity (35). In contrast, Bcl-2 expression can confer cisplatin-induced apoptosis in ovarian cancer cells (36). Our current data showed that inhibition of Notch signaling by DAPT sensitized cisplatin-resistant human ovarian cancer cells and enhances chemosensitivity associated with modulation of apoptosis-related gene expression.

Nevertheless, γ-secretase inhibitors can cleave a number of proteins, such as ErbB-4, CD44, E-cadherin and Notch family proteins (37). In the present study, we observed that DAPT resulted in downregulation of Hes1 in ovarian cancer cells in a dose-dependent manner by confirmed DAPT-inhibited Notch expression. However, it is unknown whether there are other signaling pathways or other important events in this process and further study is needed. Apoptosis induced by cisplatin is another feature of cellular response to combined DAPT and cisplatin treatment, which manifests as the synergistic inhibitory effect on cisplatin-resistant ovarian cancer cells, although the underlying molecular mechanisms mediating this synergistic antitumor effect are not completely understood. Further studies are required to better understand how the combination of γ-secretase inhibitors with cisplatin treatment can sensitize cisplatin-resistant ovarian cancer cells, especially the sequential use of DAPT and cisplatin in combination for control of ovarian cancer.

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References