

TAZ overexpression is associated with epithelial-mesenchymal transition in cisplatin-resistant gastric cancer cells

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Abstract. Gastric cancer is one of the common malignant diseases. The poor treatment outcome is mainly due to chemotherapeutic resistance. Therefore, it is important to determine the molecular mechanism of drug resistance in gastric cancer. To explore the mechanisms of cisplatin resistance in gastric cancer cells, several approaches were performed including MTT assay, real-time RT-PCR, western blot analysis, migration and invasion assays, wound healing assay, and transfection. We found that cisplatin-resistant (CR) gastric cancer cells acquired epithelial-mesenchymal transition (EMT) phenotype. The CR cells with EMT features obtained higher migratory and invasive activities. Moreover, we observed that TAZ was highly expressed in CR cells. Consistently, depletion of TAZ caused partial reversal of EMT to MET in CR cells. Our results suggest that TAZ plays a pivotal role in CR-induced EMT. Targeting TAZ could be a potential therapeutic strategy for gastric cancer.

Introduction

Gastric cancer is one of the most common malignancies in the United States. There is a decline in gastric cancer occurrence partly due to a lower prevalence of *Helicobacter pylori* infection because of improved hygiene and a higher consumption of vegetables and lower salt intake (1). However,

it is estimated that 28,000 new cases and 10,960 deaths will occur in 2017 in the US (1). In China 679,100 new gastric cancer cases occurred in 2015, being one of the four most common cancers diagnosed (2). Notably, gastric cancer is the second leading cause of cancer death worldwide. Current therapies in clinic mainly include surgery, chemotherapy, and chemoradiation for gastric cancer (3). Although surgery remains the curative therapy, chemotherapy is the important treatment for gastric cancer (4). Due to intrinsic or acquired resistance to chemotherapeutic drug, chemotherapy often fail to achieve effective treatment benefit for gastric cancer patients. Therefore, it is important to determine the molecular mechanism of drug resistance in gastric cancer (5).

Cisplatin has been known as an effective chemotherapeutic drug for the treatment of human malignancies including gastric cancer (6). Cisplatin and fluoropyrimidine-based chemotherapy and trastuzumab have been widely used for advanced stage patients with epidermal growth factor 2 (EGFR2) positivity (7). Unfortunately, cisplatin resistance often occurred during chemotherapeutic treatment (8). Emerging evidence has suggested that drug resistant tumor cells acquired epithelial-mesenchymal transition (EMT) (9). During EMT progress, epithelial cells transit to mesenchymal phenotype, leading to loss of epithelial cell-cell junction and actin cytoskeleton reorganization (10). Subsequently, the expression of epithelial marker E-cadherin was downregulated, but the expression of mesenchymal markers was upregulated, including Vimentin, Snail, Slug, ZEB1 (zinc-finger E-box binding homeobox 1), and ZEB2 (11). A study revealed that chemoresistance to cisplatin induced EMT in human lung adenocarcinoma cells (12). Similarly, another study identified that EMT is associated with cisplatin resistance in gastric cancer cells (13). Furthermore, EMT transcription factor Snail and Slug directly contribute to cisplatin resistance in ovarian cancer cells (14). Although these studies indicated the cisplatin resistance molecular basis, the precise mechanisms of cisplatin resistance are still elusive.

TAZ (transcriptional co-activator with PDZ-binding motif) has been characterized to be involved in gastric tumorigenesis (15). It is known that YAP1 (Yes-associated protein 1) and its paralog TAZ are key factors which are negatively regulated by the Hippo pathway (16). TAZ exerts

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its oncogenic function through interaction with TEAD transcription factors. Notably, TAZ was highly expressed in gastric cancer samples (17). Strikingly, the expression of TAZ had a close correlation with lymphatic metastasis and tumor TNM (tumor, node, metastasis) stage in gastric cancer (18). Several studies have revealed that TAZ was critically involved in drug resistance in a variety of human cancers (19,20). For instance, taxol resistance is mediated by TAZ and its downstream transcriptional targets *Cyr61* and *CTGF* (connective tissue growth factor) in breast cancer cells (21). Moreover, TAZ promoted EMT and cancer stem cell maintenance in oral cancer (22). Due to TAZ playing a key role in drug resistance and EMT, it is important to determine whether TAZ could regulate drug resistance and EMT in gastric cancer. In the present study, we investigated whether cisplatin-resistant (CR) cells exhibit EMT features in gastric cancer. We also elucidate the role of TAZ in CR-mediated EMT in gastric cancer.

Materials and methods

Reagents. MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide] was obtained from Sigma (St. Louis, MO, USA). Primary antibodies including anti-TAZ, anti-Vimentin, anti-Snail, anti-Slug, anti-ZO1, anti-E-cadherin, anti- β -catenin, and anti-Tubulin were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Lipofectamine 2000 and (Roswell Park Memorial Institute) RPMI-1640 medium were purchased from Invitrogen (Carlsbad, CA, USA). The non-specific control siRNA and TAZ siRNA oligonucleotides were purchased from GenePharma (Shanghai, China). Tumor Invasion Assay System was obtained from BD Biosciences (Bedford, MA, USA). Protein assay kit was purchased from Bio-Rad Laboratories (Hercules, CA, USA).

Cell culture. Human gastric cancer cell lines MGC803 and SGC7901 were cultured in RPMI-1640 medium supplemented with 5% FBS (fetal bovine serum), penicillin, and streptomycin. Cells were maintained in a humidified 5% CO₂ incubator at 37°C. Cells were cultured in medium with increasing concentrations of cisplatin for more than 6 months to establish CR cell lines (23).

MTT assay. The gastric cancer cells were seeded in each well of the 96-well plates. After 24 h incubation, the cells were treated with different concentrations of cisplatin for 72 h. MTT assay was conducted as previously described (24).

Transwell invasion assay. Cell invasion was detected using 24-well inserts with matrigel in 8- μ m pores as previously described (24). The invasive activity of cells was detected by using Transwell Invasion kit following the protocol from the manufacturer as previously described (24). Briefly, cells were cultured in the upper chamber of the inserts and RPMI-1600 medium with 10% FBS was added in the lower chamber. After 12 h at 37°C, cells on the upper side of the Transwell were removed. The invading cells on the lower side were fixed and stained with Giemsa solution as well as photographed under a microscope.

Cell attachment and detachment assay. Cell attachment and detachment assays were performed as previously described (25). Briefly, for attachment assay, gastric cancer cells were added in 24-well plates. After 1 h of incubation, we removed the unattached cells and counted the attached cells. For cell detachment assay, after 24 h incubation, the detached cells by 0.05% trypsin digestion for 2 min were counted. The remaining attached cells were trypsinized and counted. The results are presented as a percentage of the attached or detached cells to total cells.

Wound healing assay. The cells were seeded in 6-well plate. After the cells reached >90% confluency, the scratch wound was generated by a pipette tip. After 16 h, photographic images were taken (26).

Reverse transcription-PCR (polymerase chain reaction) analysis for gene expression. The total RNA was isolated with TRIzol (Invitrogen) according to the manufacturer's protocols. The real-time PCR reactions and the primers used in PCR reaction were previously described (24).

Western blot analysis: Cells were lysed with RIPA buffer supplemented with protease inhibitors. The protein concentrations were measured by the Bio-Rad protein assay kit. The same amount of protein samples were separated by SDS-PAGE (sodium dodecyl sulfate polyacrylamide gel electrophoresis) and then electrotransferred to the membranes. The membranes were immunoblotted with primary antibodies for western blotting as previously described (27).

Transfection. Gastric cells were seeded in 6-well plates and transfected with TAZ siRNA, or control siRNA (GenePharma) using Lipofectamine 2000 as previously described (24). After the transfection, the cells were applied for further investigation as described under the results section.

Statistical analysis. Statistical analyses were performed to evaluate significance between different groups by GraphPad Prism 5.0 (Graph pad Software, La Jolla, CA, USA). All data were expressed as mean \pm SD (standard deviation). $P < 0.05$ was considered to indicate a statistical significant difference.

Results

Establishment of cisplatin-resistant gastric cancer cell lines. To determine the acquired function of drug resistant gastric cancer cells, the cisplatin-resistant (CR) gastric cancer cells were established. MGC803 and SGC7901 cells were exposed to low concentration of cisplatin for several days. The dead cells were removed and increasing concentrations of cisplatin were added in culture medium. After gastric cancer cells were cultured for more than 6 months with increasing concentrations of cisplatin, CR cells (MGC803/DDP and SGC7901/DDP) cells were established (23). Our MTT assay showed that 3 μ M cisplatin led to approximately 50% cell growth inhibition in both MGC803 and SGC7901 cells (Fig. 1A). Moreover, 4 μ M cisplatin treatment caused more than 60% and 80% growth inhibition in MGC803 and SGC7901 cells, respectively (Fig. 1A). However, 2 μ M cisplatin did not inhibit cell growth in

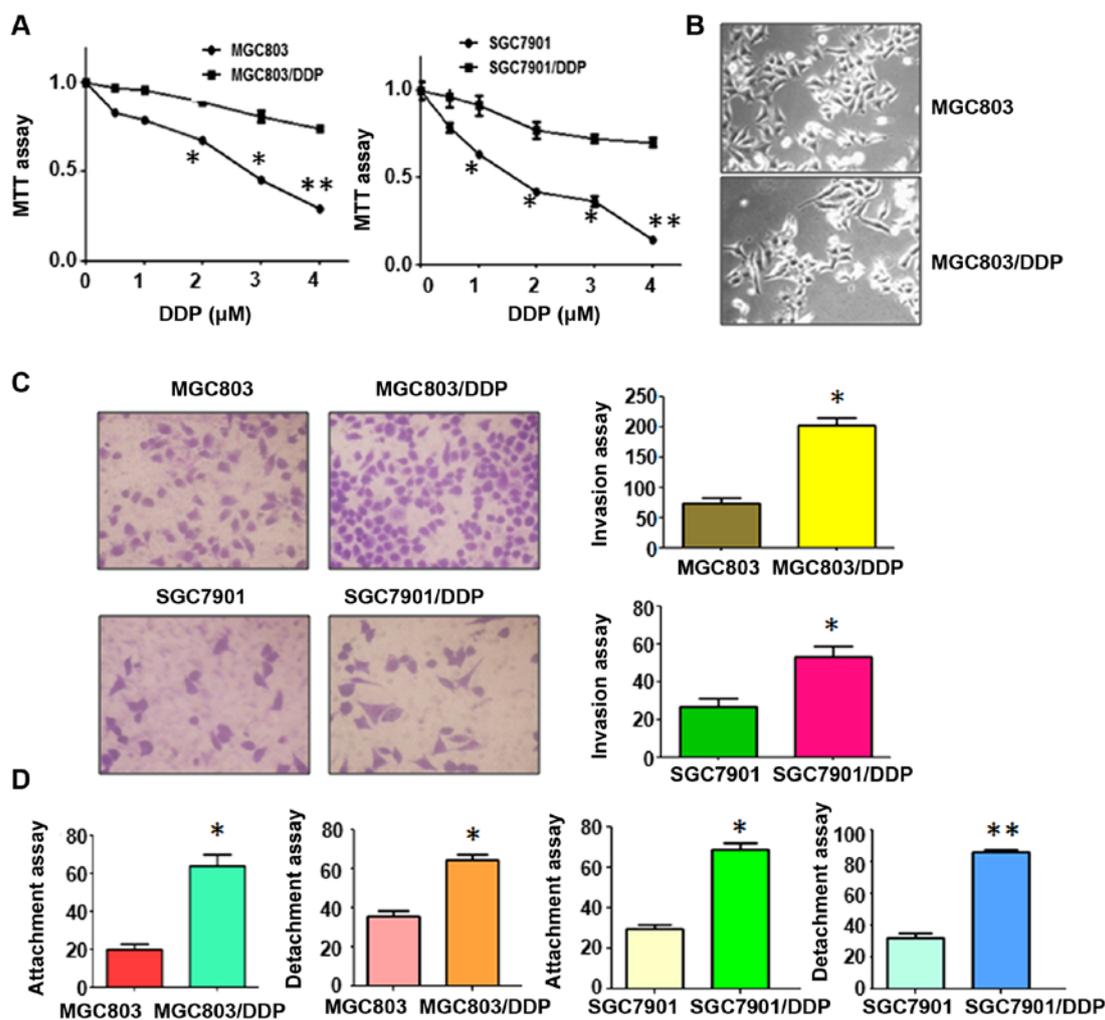


Figure 1. Cisplatin-resistant (CR) cells exhibited EMT phenotype. (A) MTT assay was conducted in parental and CR gastric cancer cells. *P<0.05, **P<0.01 vs. control. (B) Cell morphology was observed by microscopy in parental and CR cells. Parental MGC803 and SGC7901 cells displayed an epithelioid appearance, whereas their CR cells showed elongated, irregular fibroblastoid morphology. (C) Invasion assay was performed to measure the invasive capacity in parental and CR cells. *P<0.05 vs. control. (D) Cell attachment and attachment assays were assessed in parental and CR cells. *P<0.05, **P<0.01 vs. control.

both CR cells (Fig. 1A). The resistant cells were continuously cultured in medium with 2 μM cisplatin for the following study.

CR cells exhibit EMT features. It has been documented that drug resistant cells acquired EMT phenotype (28,29). In line with this concept, MGC803/DDP and SGC7901/DDP cells displayed morphologic changes, such as EMT phenotype. Both MGC803/DDP and SGC7901/DDP cells become elongated, and fibroblastoid in morphology, whereas MGC803 and SGC7901 were a rounded shape (Fig. 1B). EMT-type cells often have aggressive characteristics. Our Transwell invasion assay showed that the numbers of invasive cells were increased in CR cells compared with parental cells (Fig. 1C). Consistently, CR cells exhibited enhanced capacity of detachment and attachment (Fig. 1D). Similarly, our wound healing assay results demonstrated that CR cells acquired increased motility activity (Fig. 2A). Altogether, CR cells acquired EMT characteristics.

CR cells have EMT molecular marker changes. To further confirm whether CR cells acquired EMT, we detected the expression of EMT molecular markers in CR cells and their

parental cells, including N-cadherin, E-cadherin, Vimentin, Slug, and ZO-1. Our real-time RT-PCR results indicated that the expression of E-cadherin and ZO-1 was downregulated in CR cells, whereas the levels of N-cadherin, Vimentin, and Slug were upregulated in CR cells (Fig. 2B). Importantly, our western blotting data validated that epithelial molecules were decreased in CR cells, but mesenchymal markers were increased in CR cells (Fig. 3). These findings suggest that CR cells acquired a mesenchymal feature.

Overexpression of TAZ in CR cells. Several studies have revealed that TAZ is associated with EMT in cancers (30,31). To determine whether TAZ was involved in CR-induced EMT process, real-time RT-PCR and western blotting were performed to measure the transcription and translation levels of TAZ in CR cells. We observed that both mRNA and protein levels of TAZ were significantly upregulated in CR cells (Fig. 4A). Notably, Notch1, a target of TAZ, was also increased (Fig. 4A). We also found that β-catenin is highly expressed in CR cells (Fig. 4A). Therefore, our results displayed that the acquisition of EMT might be due to higher expression of TAZ in CR cells.

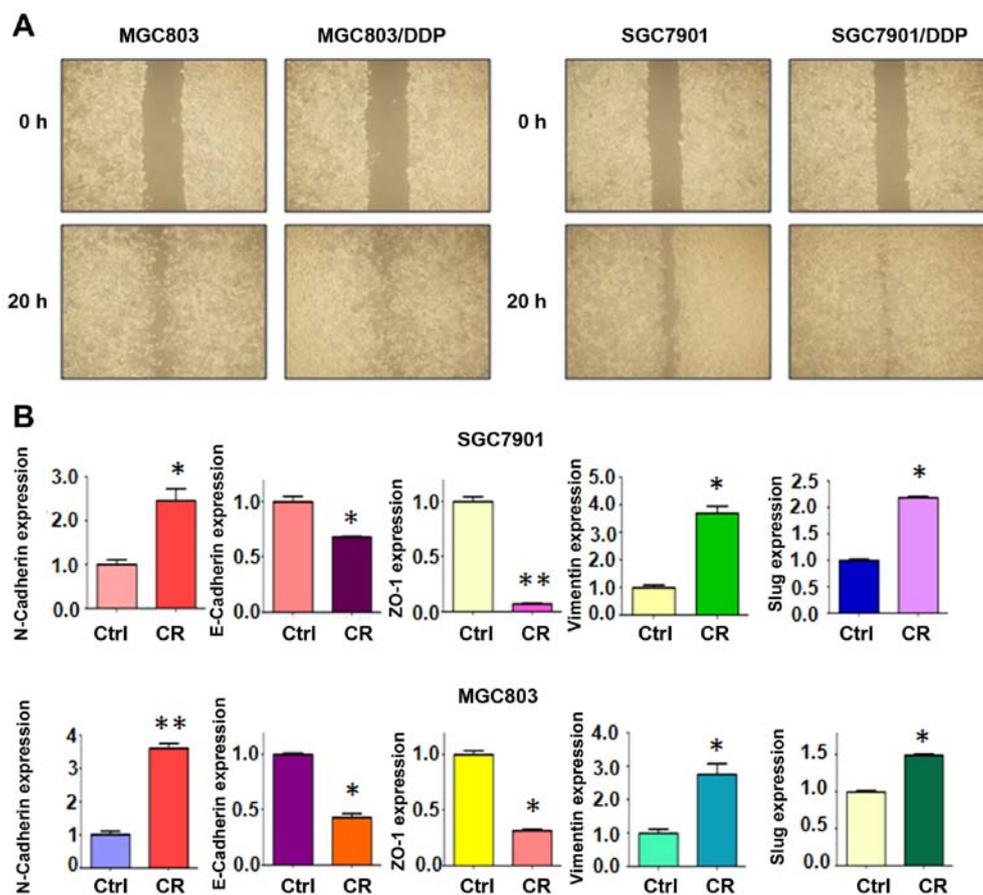


Figure 2. Cisplatin-resistant (CR) cells have enhanced motility activity. (A) Wound assays were performed in parental and CR cells. (B) Real-time RT-PCR assay was conducted to detect the expression of EMT markers in parental and CR cells. * $P < 0.05$ CR, ** $P < 0.01$ vs. control. Ctrl, control.

Depletion of TAZ reverses EMT to MET in CR cells. To further dissect the role of TAZ in CR-triggered EMT, we knocked down the expression of TAZ in CR cells. Our TAZ siRNAs significantly downregulated the expression of TAZ in CR cells (Fig. 4B). We selected TAZ siRNA3 for the following study. We found that depletion of TAZ reverses EMT to mesenchymal-epithelial transition (MET) in CR cells (Fig. 4C). Noteworthy, downregulation of TAZ decreased cell attachment and detachment activities (Fig. 4D). These data suggest that TAZ plays an essential role in CR-induced EMT.

Depletion of TAZ retards motility and invasion in CR cells. To deeper define the function of TAZ in CR cells, Transwell invasion assay was conducted. Our invasion results displayed that downregulation of TAZ significantly inhibited cell invasion in CR cells (Fig. 5A). In support of this, our wound healing assay showed that depletion of TAZ reduced cell motility in CR cells (Fig. 5B). Therefore, TAZ plays an important role in regulation of cell migration and invasion in CR cells.

Depletion of TAZ regulates expression of EMT markers. Next, we explore whether knockdown of TAZ could regulate the expression of EMT markers. To achieve this goal, the mRNA and protein levels of EMT markers were measured by real-time RT-PCR and western blotting, respectively. We found that downregulation of TAZ increased the expression of epithelial markers, but decreased the level of mesenchymal markers

(Figs. 5C and 6A and B). Therefore, these results further confirmed the role of TAZ in regulation of EMT in CR cells.

Downregulation of TAZ enhances CR cells to cisplatin sensitivity. To determine whether depletion of TAZ expression enhances CR cells to cisplatin sensitivity, MTT assay was used to detect the cell growth in TAZ siRNA transfected CR cells. We observed that downregulation of TAZ significantly attenuated cell growth inhibition approximately 20-30% induced by 2 μ M cisplatin (Fig. 6C). These MTT results implied that downregulation of TAZ could enhance CR cells to cisplatin-induced cell growth inhibition.

Discussion

Gastric cancer is the fourth most common cancer worldwide. Since chemotherapy often has treatment failure due to drug resistance, it is pivotal to explore the mechanisms of chemotherapy resistance in gastric cancer. It is clear that reasons of drug resistance are complex and multifactorial. Thus, it is necessary to define the mechanisms of chemoresistance in gastric cancer (23). In the present study, we aimed to elucidate the molecular basis of chemoresistance in gastric cancer. We observed that cisplatin-resistant (CR) gastric cancer cells obtained EMT features, leading to enhanced motility and invasion. Mechanistically, higher expression of TAZ was found in CR cells. Moreover, depletion of TAZ reversed CR-induced

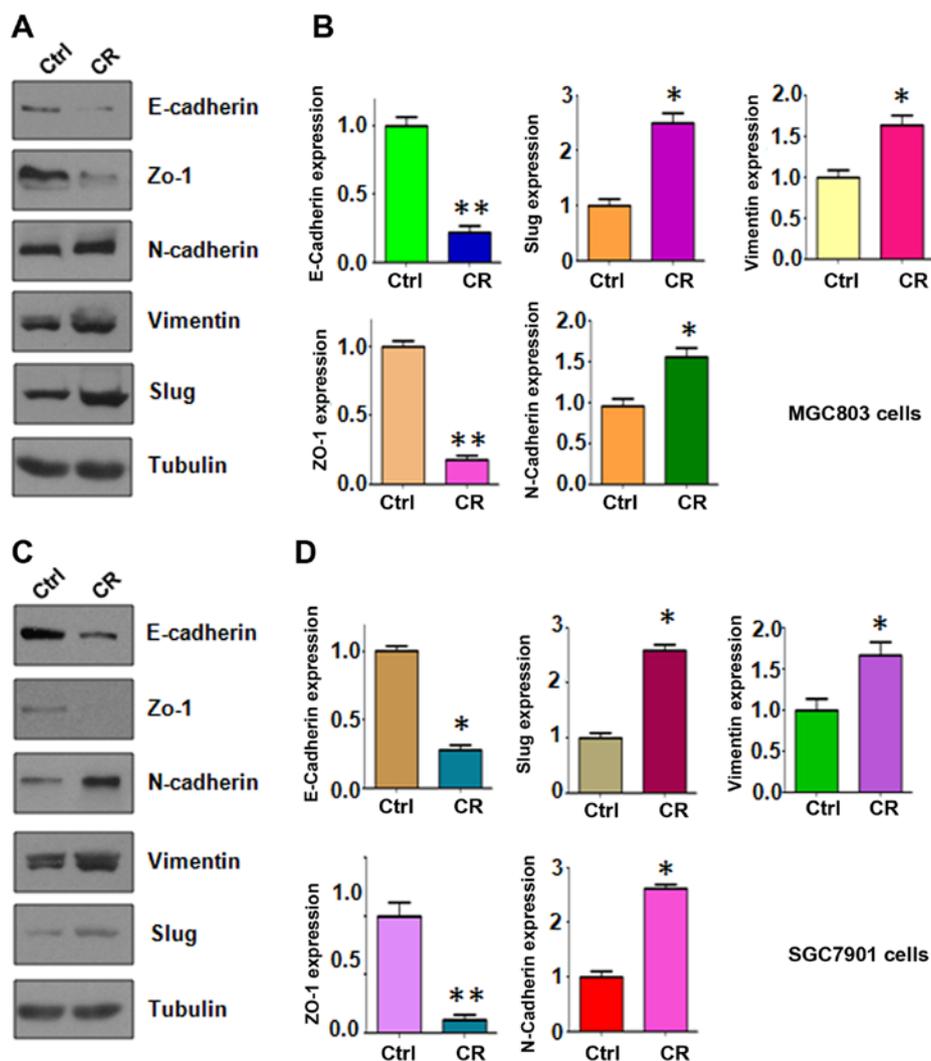


Figure 3. Cisplatin-resistant (CR) cells have EMT marker changes. (A) Western blot analysis was used to detect the expression of E-cadherin, Snail, Slug, and Vimentin in MGC803 and MGC803 CR cells. Ctrl, control. (B) Quantitative results are illustrated for (A). * $P < 0.05$, ** $P < 0.01$ vs. control. (C) Western blot analysis was performed to measure the expression of E-cadherin, Snail, Slug, and Vimentin in SGC7901 and SGC7901 CR cells. (D) Quantitative results are illustrated for (C). * $P < 0.05$, ** $P < 0.01$ vs. control.

EMT to MET. Our findings demonstrated that TAZ plays a critical role in chemoresistance and EMT in gastric cancer cells. Therefore, inhibition of TAZ could be a promising approach to overcome cisplatin resistance in gastric cancer.

Emerging evidence has demonstrated that chemoresistant cancer cells are associated with EMT (32). For example, gemcitabine-resistant pancreatic cancer cells displayed EMT via upregulation of Notch signaling pathway (33). Another study showed that gemcitabine-resistant hepatocellular carcinoma cells underwent EMT process through activation of PDGF-D pathway (24). Cisplatin resistance in gastric cancer cells is correlated with Her2 upregulation-induced EMT (34). CR cervical cancer cells exhibited EMT phenotype and overexpression of Sema4C (35). Consistent with these findings, our current study revealed that CR gastric cancer cells acquired EMT phenotype with decreased E-cadherin and increased expression of mesenchymal markers including Snail, Slug, Vimentin, and N-cadherin. Notably, EMT-type cells exhibited higher activities of migration and invasion. We confirmed that CR gastric cancer cells are associated with EMT.

Multiple studies have elucidated that some signaling pathways controlled cisplatin-induced chemoresistance. Sun *et al* found that NF- κ B signaling played irreplaceable roles in cisplatin-induced chemoresistance and tumor progression in bladder cancer (36). Liu *et al* reported that cisplatin resistance is associated with increased motility and stem-like properties through upregulation of STAT3/Snail axis in tumor cells (37). One study showed that SET-mediated NDRG1 inhibition is involved in cisplatin resistance and EMT in human lung cancer cells (38). Another study demonstrated that Fbw7 upregulation promoted cisplatin cytotoxicity in non-small cell lung cancer cells (39). Wang *et al* reported that Akt/ β -catenin/Snail signaling pathway was associated with CR induced EMT in lung cancer cells (40). Downregulation of Par-4 conferred cisplatin resistance through PI3K (phosphoinositide 3-kinase)/Akt pathway-dependent EMT in pancreatic cancer cells (41). Connexin 43 has been found to reverse the resistance of lung adenocarcinoma cells to cisplatin through inhibition of EMT (42). Further study revealed that prolonged pemetrexed pretreatment enhanced

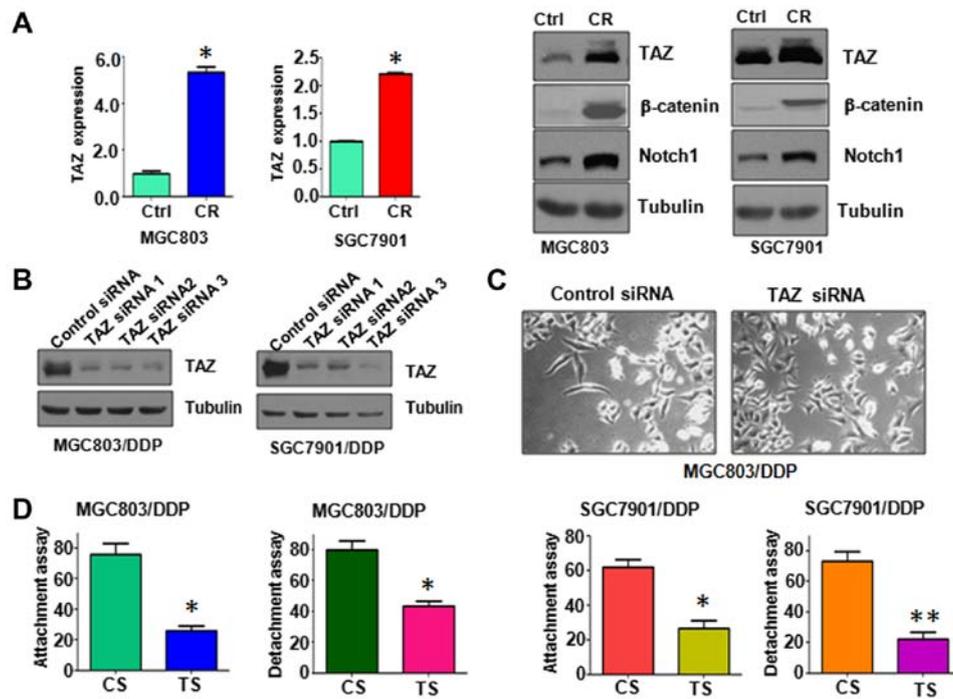


Figure 4. Cisplatin-resistant (CR) cells have high expression of TAZ. (A, left panel) Real-time RT-PCR assay was conducted to detect the expression of TAZ in parental and CR cells. * $P < 0.05$ CR vs. control siRNA. (A, right panel) Western blot analysis was performed to detect the expression of TAZ in parental and CR cells. (B) Western blot analysis was performed to detect the efficacy of TAZ siRNA transfection. (C) Cell morphology was assessed by microscopy in CR cells transfected with TAZ siRNA. (D) Cell attachment and detachment assays were measured in CR cells transfected with TAZ siRNA. * $P < 0.05$, ** $P < 0.01$ vs. control. CS, control siRNA; TS, TAZ siRNA.

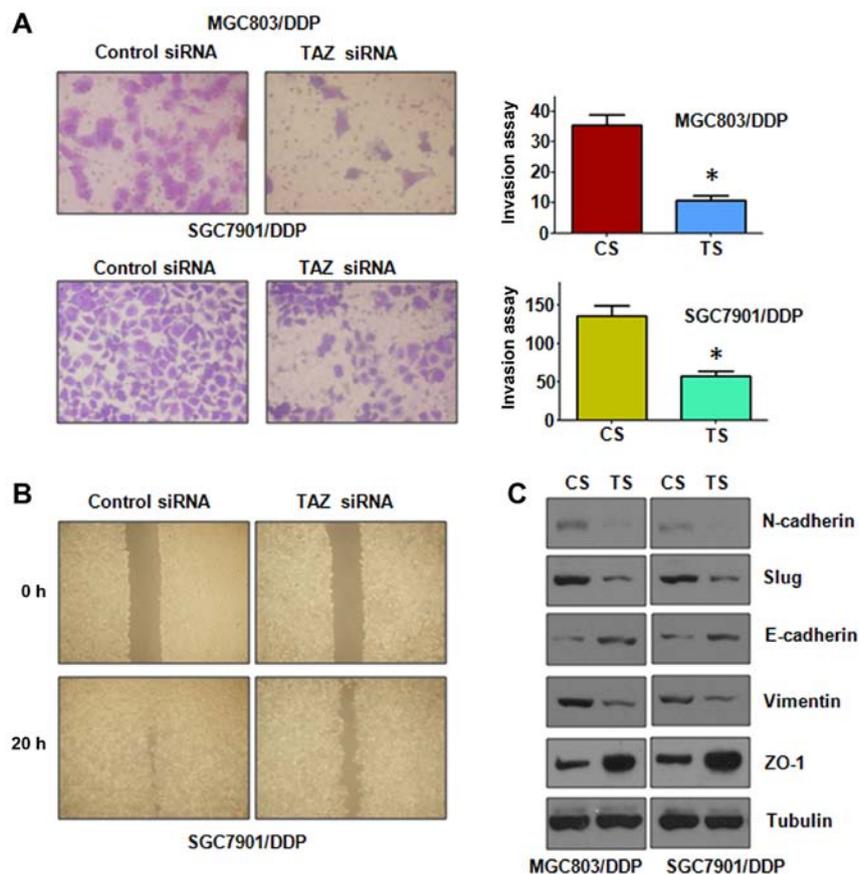


Figure 5. Depletion of TAZ inhibits motility and invasion and regulates expression of EMT markers in cisplatin-resistant (CR) cells. (A, top panel) Invasion assays were performed in CR cells transfected with TAZ siRNA. (A, bottom panel) Quantitative results are illustrated for the top panel. * $P < 0.05$ vs. control siRNA. (B) Wound healing assays were used to detect the motility in CR cells transfected with TAZ siRNA. (C) Western blot analysis was used to detect the expression of EMT markers in CR cells transfected with TAZ siRNA. CS, control siRNA; TS, TAZ siRNA.

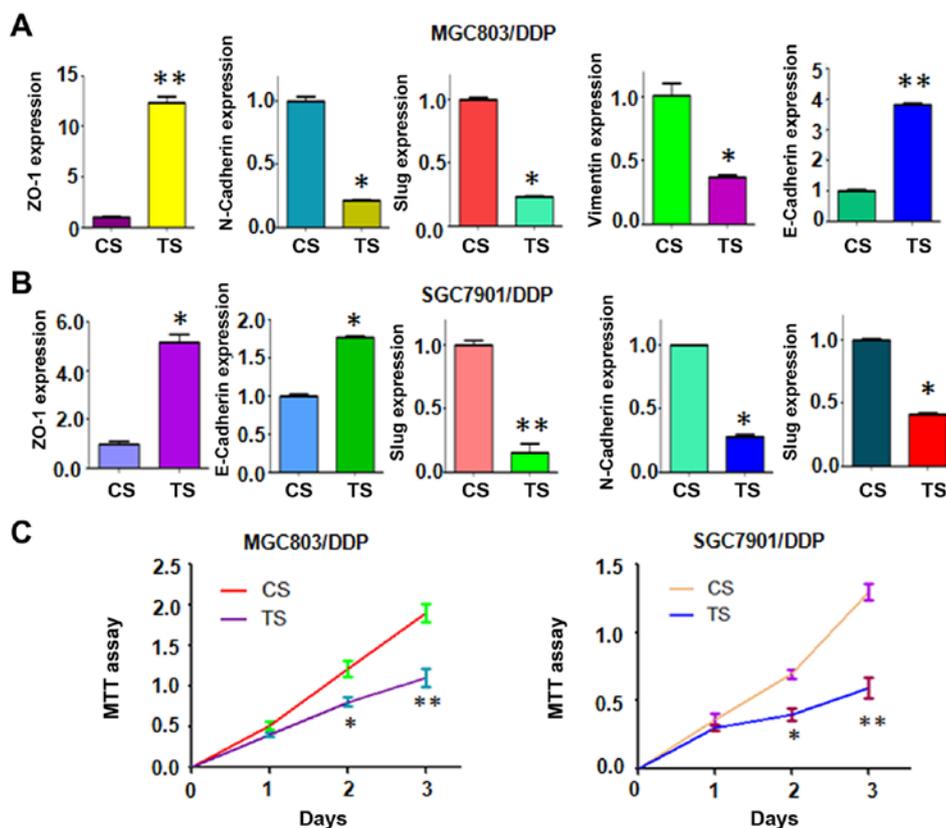


Figure 6. Depletion of TAZ regulates mRNA levels of EMT markers in cisplatin-resistant (CR) cells. (A and B) Real-time RT-PCR was performed to quantify mRNA expression of EMT markers in CR cells transfected with TAZ siRNA. CS, control siRNA; TS, TAZ siRNA. *P<0.05, **P<0.01 compared with control siRNA. (C) MTT assay was performed in CR cells treated with TAZ siRNA. CS: control siRNA; TS: TAZ siRNA. *P<0.05, **P<0.01 compared with control siRNA.

persistence of cisplatin-induced DNA damage and eliminated tumor cells with EMT and cancer stem-like features in lung cancer cells (43). Our study identified that TAZ governed the CR-mediated EMT in gastric cancer cells. Further investigation is required to explore how TAZ controls CR-induced EMT in gastric cancer.

Accumulating evidence has demonstrated that miRNAs play an essential role in drug resistance and EMT in human cancers. For example, it has been observed that miR-20a induced cisplatin resistance via targeting CYLD in human gastric cancer cells (44). Zhao *et al* found that miR-181a suppressed autophagy and sensitized gastric cancer cells to cisplatin (45). In addition, miR-26a enhanced the sensitivity of gastric cancer cells to cisplatin through targeting NRAS and E2F2 (46). Several miRNAs regulated the cisplatin resistance of human gastric cancer cells. Importantly, Chen *et al* reported that miR-206 regulated cisplatin resistance and EMT partly by targeting MET in human lung cancer cells (12). Wang *et al* found that miR-30a modulated EMT and cisplatin sensitivity in gastric cancer cells (13). Zuo *et al* identified that miR-141 inhibited tumor growth and metastasis via directly targeting TAZ in gastric cancer studies (47). These reports indicated that regulation of these miRNAs could be helpful to govern cisplatin-induced EMT.

In summary, our study validated that CR gastric cells acquired EMT in part due to overexpression of TAZ. In addition, consistent with other study (26,48), TAZ was found to increase the expression of Notch-1 and β -catenin. Notch-1

and Wnt/ β -catenin have been characterized as the drivers to induce EMT (33,49). Therefore, we believe that TAZ-induced EMT could be partly through upregulation of Notch-1 and β -catenin. Our results also indicated that downregulation of TAZ could be a useful strategy for restoring sensitivity to cisplatin. Indeed, one study has shown that knockdown of TAZ modified breast cancer cell sensitivity to EGFR inhibitors (50). In line with this, we found that depletion of TAZ sensitized CR cells to cisplatin treatment. Interestingly, one natural compound, curcumin, has been reported to inhibit the expression of TAZ in bladder cancer cells (51) and pancreatic cancer cells (48). Due to non-toxic nature of natural agents, inhibition of TAZ by these compounds could be a safe and effective approach for the prevention and the treatment of gastric cancer.

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