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**MiR-363 Inhibits Cisplatin Chemoresistance of Epithelial Ovarian Cancer by
Regulating Snail-Induced Epithelial–Mesenchymal Transition**

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Epithelial–mesenchymal transition

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Abstract

Chemoresistance is a major barrier to successful cisplatin-based chemotherapy for epithelial ovarian cancer (EOC), and emerging evidences suggest that microRNAs (miRNAs) are involved in the resistance. In this study, it was indicated that miR-363 downregulation was significantly correlated with EOC carcinogenesis and cisplatin resistance. Moreover, miR-363 overexpression could resensitise cisplatin-resistant EOC cells to cisplatin treatment both *in vitro* and *in vivo*. In addition, data revealed that EMT inducer Snail was significantly upregulated in cisplatin-resistant EOC cell lines and EOC patients and was a functional target of miR-363 in EOC cells. Furthermore, *snail* overexpression could significantly attenuate miR-363-suppressed cisplatin resistance of EOC cells, suggesting that miR-363-regulated cisplatin resistance is mediated by *snail*-induced EMT in EOC cells. Taken together, findings suggest that miR-363 may be a biomarker for predicting responsiveness to cisplatin-based chemotherapy and a potential therapeutic target in EOC.

Introduction

Ovarian cancer (OC) is the second most common cancer and a leading cause of death from gynaecologic malignancies in women worldwide (1). Epithelial ovarian cancer (EOC) is the most common type and deadly form of high-grade serous OC (2). The standard treatment for EOC is debulking surgery, followed by platinum-based chemotherapy (3). Cisplatin, a platinum compound, is one of the most common first-line antitumour agents, which bind to and cross-link DNA in cancer cells (4, 5). However, due to cisplatin resistance, the overall 5-year survival rate of EOC patients is about 40% (6). Accumulating evidence demonstrated that processes of the epithelial-mesenchymal transition (EMT), which promote cancer progression and metastasis, played a role in the development of chemoresistance (7). In EOC, Marchini et al. indicated that several genes involved in EMT were associated with overall or progression-free survival, suggesting that EMT was vital to resistance mechanisms (8).

MicroRNAs (miRNAs) are endogenous small non-protein-coding RNA molecules with approximately 22 nucleotides, which usually function as negative regulators of gene expression via binding to targeted mRNA (9). Numerous studies demonstrated that miRNAs have been involved in various biological processes, including cell proliferation, homeostasis, cellular differentiation and tumorigenesis (10-12). It's have been reported that miR-363 acts as a tumor suppressor in thyroid (13), gastric (14), colorectal (15), breast (16), and renal (17) cancers. Besides, miR-363 plays an oncogenic role in prostate cancer (18) and glioma (19). In OC, low miR-363 levels were associated with advanced stage, lymph node metastasis, and poor prognosis, meanwhile, miR-363 could inhibit OC cell growth, migration and invasion (20). Recent evidence indicates that miR-363 play important roles in chemoresistance to multiple anticancer drugs. For example, miR-363 promoted resistance to doxorubicin + cisplatin + 5-FU in gastric cancer via targeting FBW7 (14); In breast cancer, miR-363 reversed the resistance to cisplatin by negative regulating of Mcl-1, which is an anti-apoptotic Bcl-2 family member and often overexpressed in

breast tumors(16); miR-363 reduced oxaliplatin resistance by targeting the 3'-UTR of NR2F1-AS1 and ABCC1 mRNA in hepatocellular carcinoma(21). However, whether miR-363 modulate cisplatin resistance in EOC and the mechanisms underlying the resistance remains to be fully understood.

In this study, qRT-PCR assay was applied to assess miR-363 expression in primary cisplatin-sensitive and cisplatin-resistant EOC patients and tissues. In addition, *in vitro* and *in vivo* functional studies of miR-363 were conducted to determine their potential roles in the regulation of cisplatin resistance. Furthermore, data indicated that miR-363 might directly target Snail, and this interaction played an important role in the regulation of chemoresistance in EOC.

Results

Decreased miR-363 is associated with EOC tumour progression and cisplatin chemoresistance

miR-363 expression levels were evaluated in 107 malignant EOC tissues resected at the time of primary surgery from patients who subsequently received cisplatin-based primary therapy and 29 benign tissue samples. qRT-PCR revealed that miR-363 expression levels were significantly decreased in malignant EOC tissues compared with benign tissues (Fig. 1A). Correlation analysis in 107 patients with malignant EOC further revealed that miR-363 downregulation was significantly correlated with high FIGO stage ($P=0.03$), metastasis ($P=0.011$) and chemoresistance ($P=0.006$) (Table 1). 83.3% of the patients with high miR-363 expression levels and only 56.9% of the patients with low miR-363 expression levels had primary chemosensitivity. miR-363 relative expression levels in chemoresistant group were significantly lower than that of chemosensitive group (Fig. 1B). In order to substantiate the involvement of miR-363 in cisplatin response *in vitro*, miR-363 expression levels and cisplatin sensitivity in chemoresistant OVC cells A2780cp and C13 as well as their chemosensitive counterparts A2780s and OV2008 were evaluated. qRT-PCR data revealed that the levels of miR-363 were significantly decreased in A2780cp and C13

cells compared to that of A2780s and OV2008 cells (Fig. 1C). The half maximal (50%) inhibitory concentration (IC₅₀) values were significantly higher in A2780cp and C13 cells compared with A2780s and OV2008 cells (Fig. 1D). Taken together, data suggest that decreased miR-363 may be associated with EOC carcinogenesis and cisplatin resistance.

miR-363 sensitizes EOC cells to cisplatin treatment *in vitro* and *in vivo*

To investigate the role of miR-363 in regulating cisplatin sensitivity in EOC cells, miR-363 has been restored in the cisplatin-resistant cells A2780cp and C13 and has been knocked down in the cisplatin-sensitive cells A2780s and OV2008, respectively (Fig. 2A). Enhancing the expression of miR-363 in A2780cp and C13, cell migration and invasion were repressed (Fig. S1). MTT assays with different cisplatin doses showed that cisplatin sensitivities were significantly increased after forced miR-363 overexpression in A2780cp and C13 cells (Fig. 2B). Conversely, the cisplatin sensitivities of A2780s and OV2008 cells were decreased after miR-363 silencing (Fig. 2C). To further evaluate the role of miR-363 in regulating cisplatin sensitivity *in vivo*, the cisplatin-resistant cells A2780cp were injected subcutaneously, overexpressing miR-363 into BALB/C nude mice. Thirty-five days after injection, when all mice developed palpable tumours, each mouse was treated with 0.1 ml cisplatin (10 mg/kg) every 5 days and euthanized 10 days later. As shown in Fig. 2D, prior to cisplatin treatment, tumour growth was slightly inhibited in the miR-363 overexpressing group compared to the control group. Notably, upon 10-day cisplatin treatment, tumour growth was more dramatically inhibited in the miR-363 overexpressing group than in the control group (Fig. 2D). Taken together, findings suggest that miR-363 might sensitize EOC cells to cisplatin treatment.

***Snail* is upregulated in cisplatin-resistant EOC cell lines and patients**

Previous reports revealed that induction of EMT may contribute to the decreased efficacy of cisplatin therapy. To examine EMT in response to cisplatin treatment, the expression levels of EMT-related markers (*E-cadherin*, *fibronectin*, *N-cadherin*, *vimentin*) and inducers (*TGF-β1*, *snail*, *slug*, *twist*, *ZEB1*) were evaluated in

chemoresistant A2780cp and C13 cells and their chemosensitive counterparts A2780s and OV2008 cells. qRT-PCR analysis revealed that E-cadherin was significantly downregulated in chemoresistant A2780cp and C13 cells compared with that in their chemosensitive counterparts A2780s and OV2008 cells (Fig. 3A), suggesting that EMT was promoted in response to cisplatin treatment. Among EMT inducers tested, *snail* was most upregulated in A2780cp and C13 cells compared with that in A2780s and OV2008 cells (Fig. 3A). In addition, *snail* expression levels in EOC and benign tissues were detected where Snail relative expression levels were found to be higher in malignant tissues compared with benign tissues (Fig. 3B) and were also higher in cisplatin-resistant patients compared with those in cisplatin-sensitive patients (Fig. 3C). Taken together, these data suggest that *snail* is upregulated in cisplatin-resistant EOC cell lines and patients.

miR-363 directly inhibits *snail* expression in EOC cells

Then *in silico* prediction for miR-363 target genes suggested that *snail* might be a target of miR-363 in EOC cells (Fig. 3D). To validate if miR-363 could directly target 3'UTR of *snail*, a fragment of wild-type or mutant 3'UTR of *snail* was cloned into psi-CHECK2 reporter vector, respectively. Luciferase reporter assays revealed that miR-363 significantly decreased the relative luciferase activity of 3'UTR of *snail* in A2780s and OV2008 cells but had no effect on the mutant 3'UTR of Snail. qRT-PCR analysis further revealed that miR-363 mimics significantly reduced the *snail* expression, and knocking-down of miR-363 increased Snail expression in A2780s and OV2008 cells (Fig. 3E). The reverse relation of miR-363 and Snail was observed in correlation analysis of their expressions in 107 malignant EOC patients ($R^2 = 0.319$, $p < 0.01$) (Figure 3F). Taken together, these data suggest that *snail* is a functional target of miR-363 in EOC cells.

Snail is involved in miR-363-suppressed cisplatin-resistance of EOC cells

To further determine whether miR-363 exerts its effect on regulating cisplatin sensitivity through the downregulation of *snail* in EOC cells, Snail in A2780cp and C13 cells were knocked down with *snail*-specific small interfering RNAs (si-Snail),

and *snail* expression in A2780s and OV2008 cells was upregulated (Fig. 4A and S2). As expected, MTT assay revealed that *snail* knockdown significantly sensitized A2780cp and C13 cells to cisplatin (Fig. 4B), while *snail* upregulation resulted in decreased cisplatin sensitivities of A2780s and OV2008 cells (Fig. 4C). These observations were opposite the effects of miR-363. Then, *snail* expression was rescued in stable miR-363 overexpressing A2780cp or C13 cells by transfecting *snail* expression plasmids lacking 3'UTR. MTT assay revealed that Snail overexpression significantly attenuated miR-363-suppressed cisplatin resistance of A2780cp and C13 cells (Fig. 4D). Taken together, these results suggest that *snail* is a functional target of miR-363, involved in miR-363-suppressed cisplatin resistance of EOC cells.

Discussion

Cisplatin is a first-line antitumour chemotherapeutic agent; however, due to cisplatin resistance of EOC, 25% of patients will develop resistance to this agent within 6 months after chemotherapy (22). To break the therapeutic barrier of cisplatin resistance in EOC, it is urgent to clarify the mechanisms underlying cisplatin resistance and identify new biomarker and therapeutic targets.

Recently, multiple studies have indicated that miRNA dysregulation played an important role in the development of cisplatin resistance in EOC. The human let-7 family is well known to associate with tumorigenesis of different types of cancers, such as lung cancer, breast cancer, prostate cancer and OVC (23-26). It was reported that let-7e expression was significantly reduced in cisplatin-resistant EOC cell line and let-7e overexpression could resensitize cisplatin-resistant EOC cells to cisplatin (27). In this study, we have shown that miR-363 levels were significantly downregulated in malignant EOC tissues compared with benign tissues, and miR-363 downregulation was significantly correlated with high FIGO stage, metastasis and chemoresistance. Moreover, miR-363 overexpression could resensitize cisplatin-resistant EOC cells, and miR-363 knockdown could decrease the cisplatin sensitivities of normal EOC cells both *in vitro* and *in vivo*.

The upregulation of EMT inducers was supposed to be closely associated with

the development of multiple chemotherapeutic drug resistance in different cancers (28, 29). Haslehurst et al. found that the EMT inducer *snail* was upregulated in the chemoresistant EOC cells (30). By evaluating expressions of EMT inducers in both chemoresistant EOC cells and their chemosensitive counterparts, it was found that *snail* was highly upregulated in chemoresistant EOC cells and cisplatin-resistant patients. In addition, as predicted, *snail* might be a target of miR-363 with *in silico* study. The subsequent luciferase reporter assays proved the prediction, and an inverse correlation between miR-363 and Snail expression was observed in both EOC cells and patients. It was reported that knockdown of *snail* leads to cisplatin sensitization in lung adenocarcinoma and head and neck squamous cell carcinoma (28, 31). Results revealed that *snail* knockdown significantly sensitized EOC cells to cisplatin and *snail* upregulation resulted in decreased cisplatin sensitivities. Furthermore, rescued *snail* expression in miR-363 overexpressing EOC cells could significantly attenuate miR-363-suppressed cisplatin resistance of these cells. It was seen that *snail* couldn't fully restoring cell survival under cisplatin treatment condition. As we all known, each miRNA may regulates multiple (even hundreds) of target genes, there may be other target of miR-363 that affects cisplatin resistance in EOC cell lines. Just as reported, FBW7, Mcl-1, ABCC1 were target genes for miR-363 in regulating chemoresistance (14, 16, 21). In OC, miR-363 play a tumor suppressor role by targeting NOB1 (20), these are possible targets of miR-363 which affects cisplatin resistance in EOC, but we need to verify it.

In conclusion, this study provides evidence that miR-363 could sensitize EOC cells to cisplatin treatment *in vitro* and *in vivo*. More importantly, the data demonstrate for the first time that *snail* is a target of miR-363, and MiR-363 inhibits cisplatin resistance of EOC by regulating *snail*-induced EMT. Taken together, findings indicate that miR-363 may be a biomarker for predicting responsiveness to cisplatin-based chemotherapy and a potential therapeutic target in EOC.

Materials and Methods

Patients and samples

One hundred seven patients with OVC were enrolled at Xiangya Hospital (Changsha, Hunan, China) from 2013 to 2016. All patients were given written informed consent, and this study was authorized by the Ethics Committee of Xiangya Hospital, South University. Treatment response was assessed with clinical and radiologic examination evaluated by the same investigator according to RECIST (Response Evaluation Criteria in Solid Tumours, version 1.0). The responders were defined as having either complete or partial response. The non-responders included patients with stable or progressive disease.

Cell culture

Cisplatin sensitive (OV2008, A2780s) and resistant (C13, A2780cp) were generously provided by Drs. Rakesh Goel and Barbara Vanderhyden (Ottawa Hospital Cancer Center, Ottawa, ON, Canada). Cells were cultured at 37°C, 5% CO₂ in either RPMI 1640 (OV2008 and C13) or DMEM-F12 (A2780s and A2780-cp) containing 10% FBS (Invitrogen, USA). Cells (1×10^6) were plated in log growth phase onto 60-mm dishes for 24 h in the above culture medium before the initiation of treatment.

qRT-PCR

Total RNAs were extracted using the TRIzol method (Invitrogen, USA) and reversely transcribed into cDNA using the PrimeScript RT reagent Kit (TaKaRa Bio, Japan) according to the manufacturer's instructions. qRT-PCR was performed on ABI 7500 Sequence Detection System (Life Technologies, USA) using SYBR Green real-time PCR master mix (Toyobo Co., Japan). The specific primers for miRNA-363 and small nuclear U6, which was used as an internal control, were purchased from Guangzhou RiboBio (Guangzhou RiboBio Co., Ltd., Guangzhou, China). Relative expression levels were calculated using the $2^{-\Delta\Delta C_t}$ method. Primers for qRT-PCR were synthesized by Invitrogen (Shanghai, China); the sequences are listed in Table 2.

Cell proliferation assay

The cell proliferation was assessed using 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazoliumbromide (MTT) solution (Sangon Biotech, China). Forty-eight h after transfection, EOC cells were seeded into 96-well plates at an initial density of 5×10^3 cells/well. After 24 h of culture, these

cells were exposed to various concentrations of cisplatin for 24 h, respectively. Then the cells were treated with 10 μ l MTT by adding it into each well. The cells were incubated at 37°C with 5% CO₂ for another 4 h; then the medium was removed carefully, and 150 μ l dimethylsulfoxide (DMSO) solution (MP Biomedicals, USA) was added for 10 min to lyse the cells. Subsequently, the absorbance was measured at 570 nm using a microplate reader Multiskan MK (Thermo Scientific, USA). The survival rate was calculated using the equation: (mean absorbance of drug well/mean absorbance of control wells) \times 100%.

Lentiviral infection

Knocking-down miR-363 by the lentiviral GV-428 vector, encoding the anti-sense of miR-363 (anti-miR-363, based on the described sequence[23]) was designed, synthesized and sequence-verified by the GeneChem Company (Shanghai, China). Lentivirus-expressing miR-363 (Lv-miR-363) and negative control (Lv-NC) were also purchased from the GeneChem Company. To get stably infected cells, the cells were cultured in about 80% of the plates and then added by a concentration of 5.0×10^4 TU/well lentivirus. RTq-PCR was performed to determinate miR-363 expression levels after being infected for 5 days.

Animal treatment

All animal experiments were carried out in accordance with a protocol approved by the Institutional Ethical Committee (Institutional Animal Care and Use Committee of Xiangya Hospital). Five-week-old BALB/c nu/nu mice were purchased from Shanghai Laboratory Animal Center (SLAC, Shanghai, China) and were handled under specific pathogen-free conditions. A2780cp cells overexpressing miR-363 (5×10^6 cells in 0.1 ml of phosphate-buffered saline per mouse) were injected into the proximal tibia of each mouse (n=5 animals per group). Every week post inoculation, the individual tumour was measured with calipers according to the formula: $1/2 \times \text{length} \times \text{width}^2$. Seven weeks after inoculation, when all mice developed palpable tumours, each mouse was treated with 0.1 ml cisplatin (10 mg kg⁻¹) via tail vein injection once a week for 2 weeks. After complete cisplatin treatment, all of the mice were euthanized, and the tumours were excised and imaged under a light microscope.

Dual-luciferase reporter assay

The fragments from 3'UTR of *snail* containing the predicted miR-363 binding site were synthesized and cloned into the luciferase construct psi-CHECK2. The resulted vector *snail*-3'UTR-psi-CHECK2 was called the reporter vector WT-3'UTR. The corresponding mutant was called Mut-3'UTR. The miR-363 mimic or control mimic was co-transfected with the reporter vectors using transfection reagent (Invitrogen, USA). 48 h after transfection, Firefly and Renilla luciferase activities in cell lysates were measured using the Dual-Luciferase Reporter Assay Kit (E1910; Promega,USA).

Statistical analysis

The experiments were repeated at least 3 times, and the data are shown as the mean \pm s.d. Data analyses were performed using Student's t-test for simple comparison of the 2 groups. The difference in results was considered statistically significant when the *p* value was <0.05 .

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Disclosure of Conflicts of Interest

No potential conflicts of interest were disclosed.

Figure Legends

Figure 1. Decreased miR-363 levels in cisplatin-resistant EOC patients.(A) qRT-PCR analysis of miR-363 expression levels in 107 malignant EOC tissues and 29 benign tissues. (B) qRT-PCR analysis of miR-363 expression levels in primary cisplatin-sensitive (n=72) and cisplatin-resistant (n=35) EOC tissues. (C) qRT-PCR

analysis of miR-363 expression levels in chemoresistant OVC cells A2780cp and C13 and their chemosensitive counterparts A2780s and OV2008 cells. (D). MTT assay of cisplatin sensitivity in A2780cp, C13, A2780s and OV2008 cells (**p* value < 0.05)

Figure 2. Restoration of miR-363 sensitizes EOC cells to cisplatin *in vitro* and *in*

***vivo*.** (A) The chemoresistant A2780cp and C13 and chemosensitive A2780s and OV2008 cells were infected with Lv-miR-363 and Lv-NC labelled with GFP reporter, respectively. qRT-PCR was performed to determine miR-363 expression levels. (B, C) The impacts of miR-363 on cisplatin sensitivity at different doses (0, 2, 4, 6, 8 and 10 ng/ml) were determined by MTT assay. (D) BALB/C nude mice were subcutaneously inoculated with Lv-miR-363-infected A2780cp cells (n=5) and Lv-miR-NC-infected A2780cp cells (n=5), respectively. Tumour volume (cm³) was calculated every week. After 7 weeks, each mouse was treated with 0.1 ml cisplatin (10 mg kg⁻¹) via tail vein injection every 5 days and euthanized 10 days later. After complete cisplatin treatment, all mice were euthanized, and the tumours were excised and imaged under a light microscope (**p* value < 0.05).

Figure 3. Snail was upregulated in cisplatin-resistant EOC and its expression was inhibited in EOC cells by miR-363. (A) qRT-PCR analysis of the expression levels

of EMT inducers in chemoresistant OVC cells A2780cp and C13 and their chemosensitive counterparts A2780s and OV2008 cells. (B) miR-363 expression levels in 107 malignant EOC tissues and 29 benign tissues. (C) miR-363 expression levels in primary cisplatin-sensitive (n=72) and cisplatin-resistant (n=35) EOC tissues (**p* value < 0.05). (D) Snail had a predicted miR-363 binding site in 3'UTR, and luciferase activity of the reporter construct containing the wild-type or mutant miR-363 binding site was measured after co-transfection with 50 nM miRNAs. (E) qRT-PCR was performed to determine *snail* expression levels in miR-363 overexpression or knockdown cells (**p* value < 0.05). (F) The reverse relation of miR-363 and *snail* was observed in the correlation analysis of their expressions in 107 patients with malignant EOC.

Figure 4. Snail is involved in miR-363-suppressed cisplatin resistance of EOC

cells. (A) qRT-PCR was performed to determine *snail* expression levels in *snail*

knockdown cells (A2780cp and C13) and snail overexpression cells (A2780s and OV2008). (B, C, D) The impacts of *snail* and miR-363 on cisplatin sensitivity at different doses (0, 2, 4, 6, 8 and 10 ng/ml) were determined by MTT assay (**p* value < 0.05).

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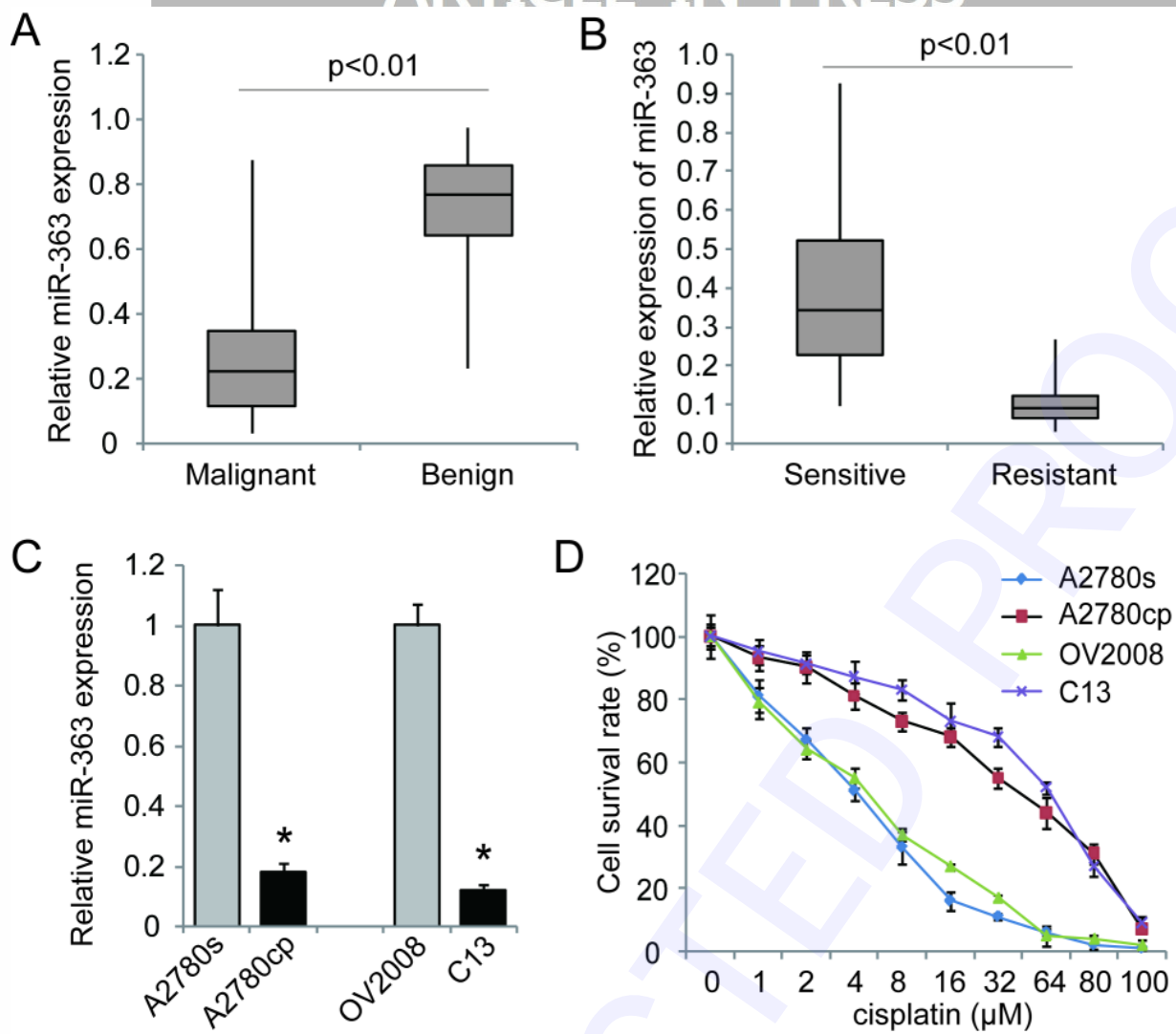


Fig. 1.

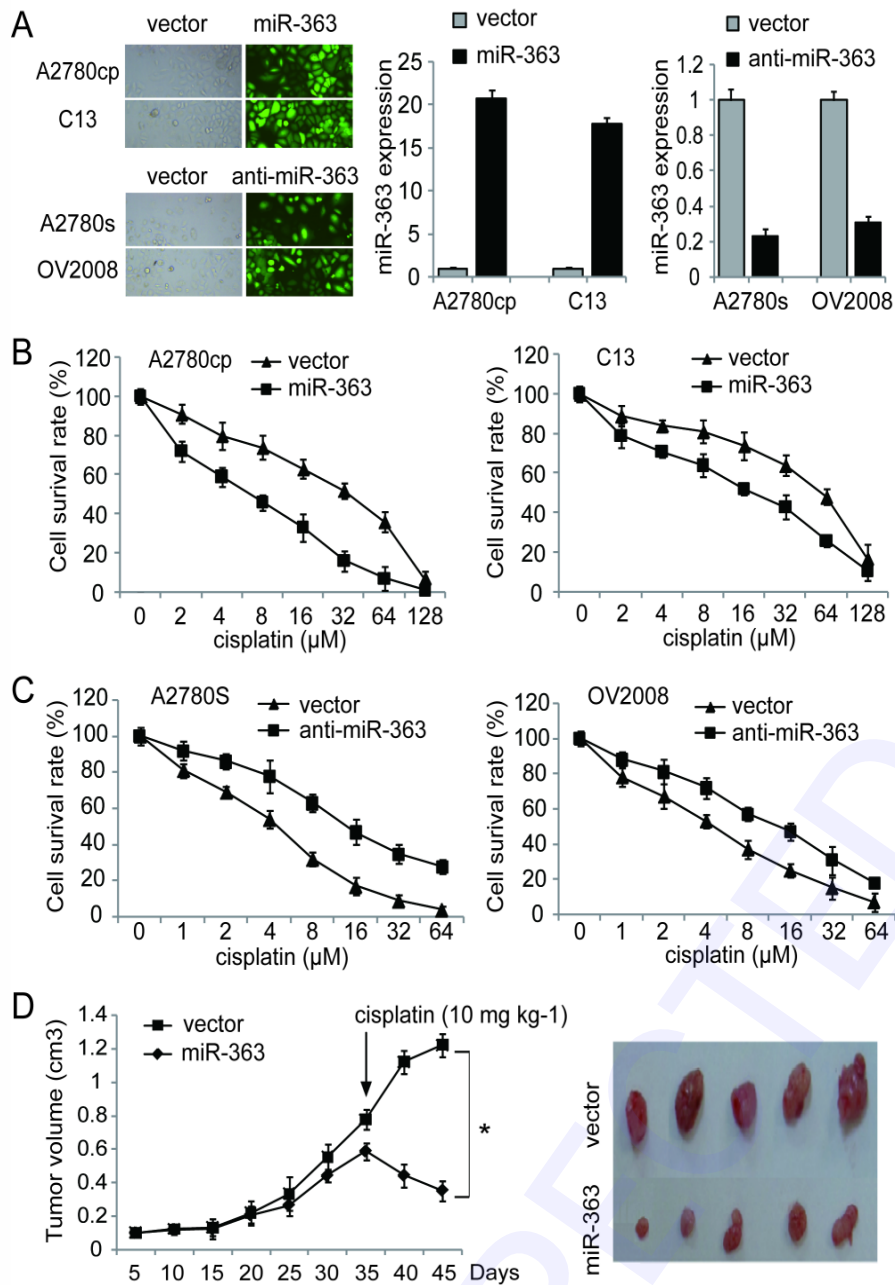


Fig. 2. Revised Figure 2

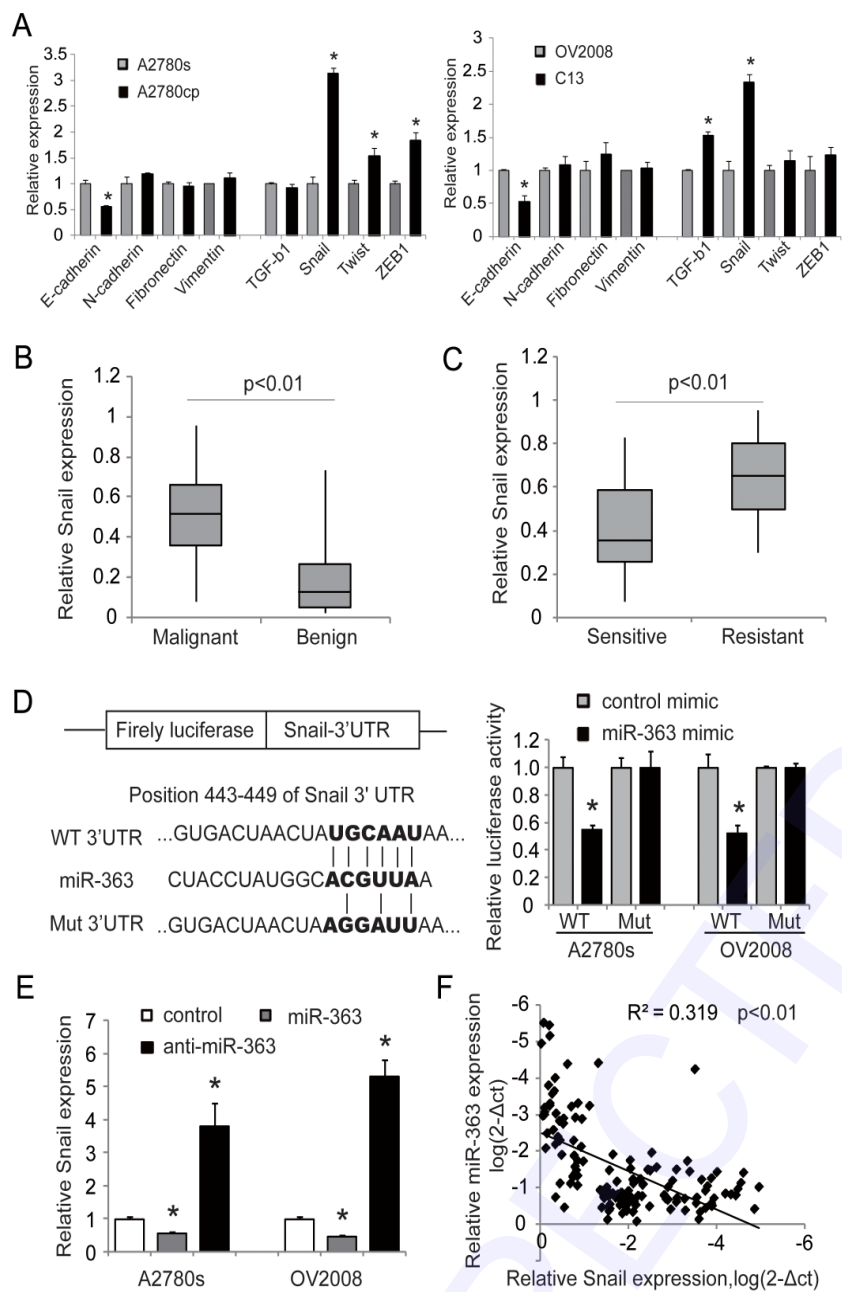


Fig. 3. Revised Figure 3

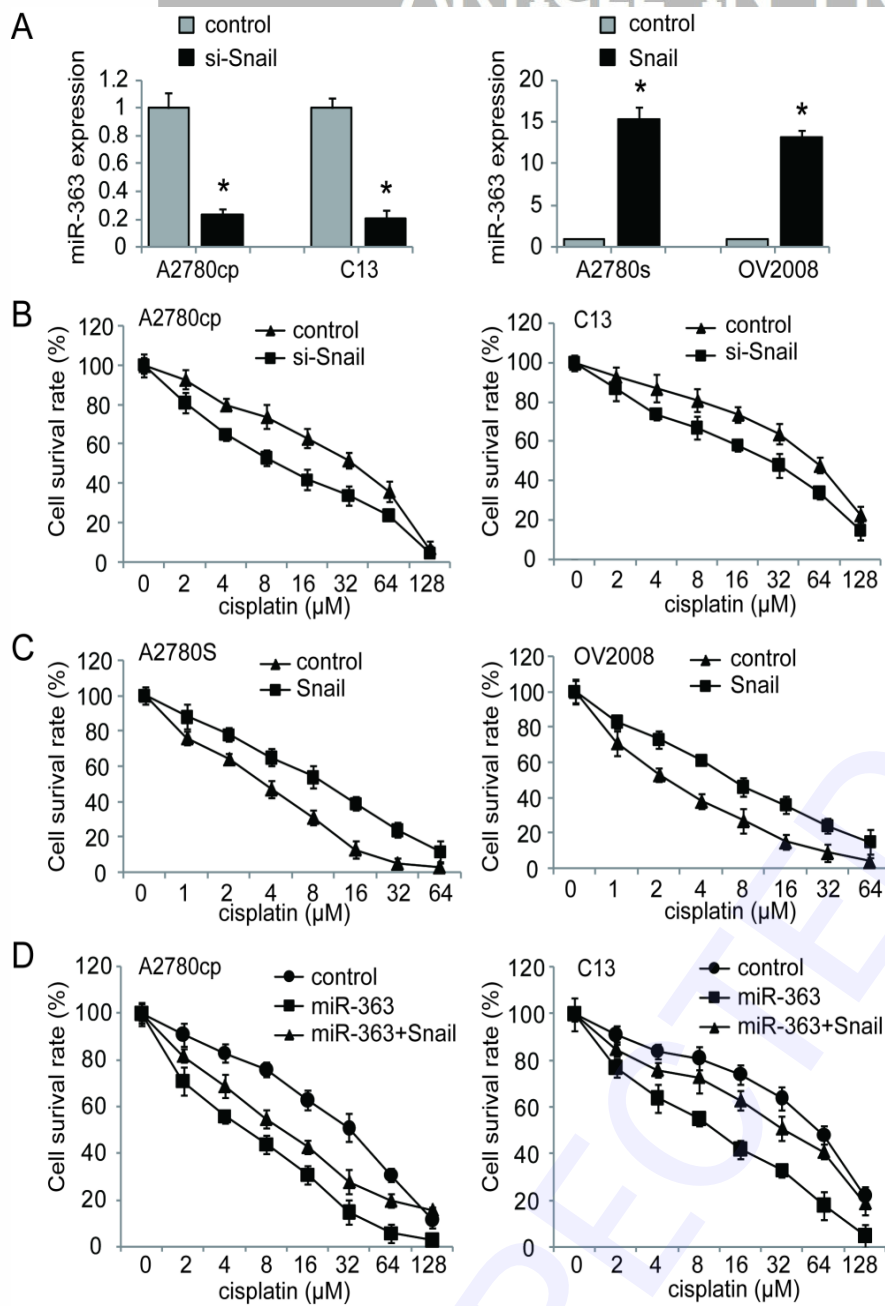


Fig. 4.

Table 1 The correlation between miR-363 expression and clinicopathological parameters of 107 EOC patients

Clinipathological parameters	Number of cases	miR-363 expression		P value
		High (%)	Lowh (%)	
Age (years)				0.844
≥ 55	53	20	33	
< 55	54	22	32	
Serum CA-125 level				0.792
≥ 35	89	34	55	
< 35	18	8	10	
FIGO stage				0.03
I+II	32	18	14	
III+IV	75	24	51	
Differentiation status				0.321
High + Medium	59	26	33	
Low	48	16	32	
Lymph node metastasis				0.011
No	52	27	25	
Yes	55	15	40	
Chemoresistance				0.006
No	72	35	37	
Yes	35	7	28	

Table 2 Primer sequences used in qPCR

Genes	Primer sequences
<i>E-cadherin</i>	F: TGCCCAGAAAATGAAAAAGG R: GTGTATGTGGCAATGCGTTC
<i>N-cadherin</i>	F: ATGACAATCCTCCAGAGTTTA R: ATCCTTATCGGTCACAGTTAG

<i>Fibronectin</i>	F: GGCGACAGGACGGACATCTTTG R: GCACAAGGCACCATTGGAATT
<i>Twist</i>	F: CGCCACAGCCCGCAGACTTCTT R: CCGACGACAGCCTGAGCAACA
<i>Snail</i>	F: GCCTAGCGAGTGGTTCTTCT R: TAGGGCTGCTGGAAGGTAAA
<i>TGF-β1</i>	F: AAGGACCTCGGCTGGAAGTG R: CCGGGTTATGCTGGTTGTA
<i>ZEB1</i>	F: CACTGGTGGTGGCCCATTAC R: TGCACCATGCCCTGAGG
<i>vimentin</i>	F: ACGTCTTGACCTTGAACGCA R: TCTTGGCAGCCACACTTTCA
<i>β-actin</i>	F: AGGGGCCGGACTCGTCATACT R: GGCGGCACCACCATGTACCCT

Materials and Methods

Snail interfering RNA transfection

The oligonucleotides for human *snail* small interfering RNAs (siRNAs) and control were synthesized by Invitrogen. The 19-bp sequences of *snail* siRNA were designed by OligoEngine Workstation software: 5'-GCTGGCAGCCATCCCACCT-3'; 5'-GCACAACAAGCCGAATACA-3' (control). A2780cp and C13 cells were transfected with 200 pmol each siRNAs by Lipofectamine 2000 (Invitrogen, USA) according to the manufacturer's instructions. At 48 h after transfection, the total RNA and protein were extracted.

Western blot

Protein was extracted from the indicated cells using RIPA lysis buffer(). BCA Protein assay reagents (Beyotime, China) were used to measure the protein concentration. A total of 30 µg of protein was separated with 10% SDS-PAGE and transferred to a PVDF membrane, which was blocked in 5% nonfat dried milk in PBS at room temperature for 2 h. The membrane was incubated with goat anti-Snail polyclonal antibody (Abcam, USA) or goat anti-β-actin monoclonal antibody (Abcam, USA) overnight at 4°C and then with donkey anti-goat secondary antibody (Abcam, USA) for 2 h. Enhanced chemiluminescence reagent (Thermo fisher, USA) was used to detect the signal on the membrane. Data were analyzed by densitometry using Image-Pro plus software 6.0 and normalized to β-actin expression.

Cell migration and invasion assay

Cell migration was assessed by wound healing assays. In brief, cells were seeded in 6-well plates and cultured to 100% confluence. Wounds of approximately 1 mm in width were scratched vertically into the cell monolayer using a 20-µL sterile pipette tip, and the detached cells were removed by aspiration. Medium was replaced with RPMI 1640 (C13) or DMEM-F12 (A2780cp) containing 3% FBS. After incubation for 24 h, the wound closure was assessed by Scion Image Software (Scion Corporation,

Frederick, USA). For cell invasion assays, the matrigel invasion chambers (BD Bioscience, USA) were used to assess cell invasion ability. Briefly, 1×10^5 cells were seeded in the upper chamber with media containing 0.1% fetal bovine serum, whereas the lower chamber was filled with media containing 10% fetal bovine serum. After incubation for 48h, the noninvading cells were removed with cotton swabs, and the cells that invaded through the membrane were stained with 0.1% crystal violet and manually counted at $\times 200$ magnification.

Figure Legends

Figure S1. Restoration of miR-363 inhibited EOC cells migration and invasion.

Enhancing the expression of miR-363 in chemoresistant cells A2780cp and C13, the migration (A) and invasion (B) were detected by scratch wound healing assay and Transwell assay respectively(* p value < 0.05).

Figure S2. The expression of *snail* protein was detected by Western blot assay.

Western blot was performed to determine snail protein levels in *snail* knockdown cells (A2780cp and C13) and *snail* overexpression cells (A2780s and OV2008), * p value < 0.05 .

Figure S1

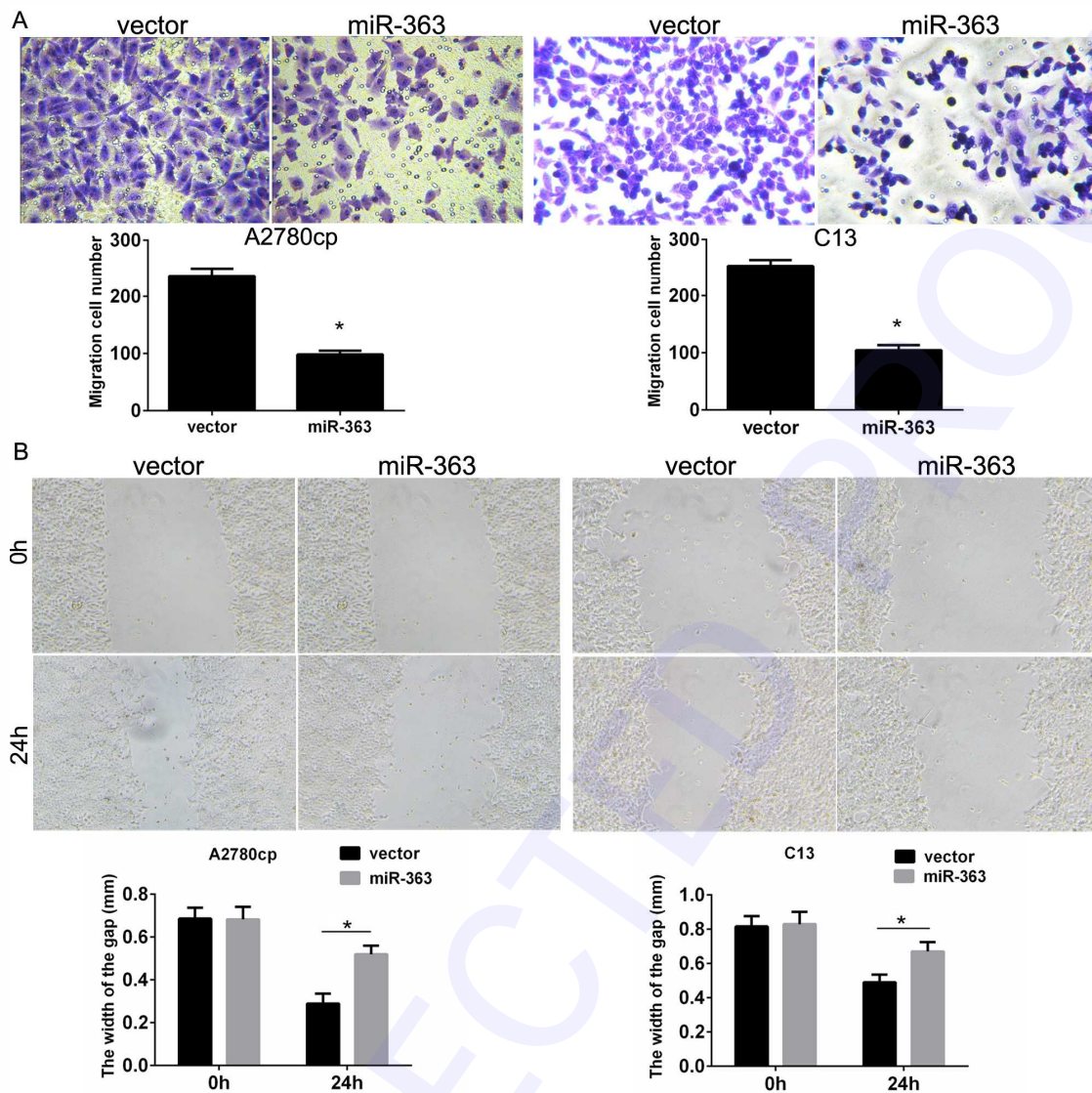


Figure S2

