Research Paper

Solanum Incanum Extract Downregulates Aldehyde Dehydrogenase 1-Mediated Stemness and Inhibits Tumor Formation in Ovarian Cancer Cells

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Received: 2015.05.19; Accepted: 2015.07.20; Published: 2015.08.22

Abstract

Solanum incanum extract (SR-T100), containing the active ingredient solamargine, can induce apoptosis via upregulation of tumor necrosis factor receptor expression and activation of the mitochondrial apoptosis pathway, and has therapeutic effects in patients with actinic keratosis. Here, we evaluate the novel molecular mechanisms underlying SR-T100-regulated stemness and chemoresistance. The concentration of SR-T100 that inhibited 50% cell viability (IC₅₀) was lower in ovarian cancer cells than in nonmalignant cells. Furthermore, the SR-T100 IC₅₀ in chemoresistant cells was similar to the IC₅₀ in chemosensitive cells. Additionally, SR-T100 increased cisplatin and paclitaxel sensitivity in chemoresistant cells. SR-T100 downregulated the expression of stem cell markers, including aldehyde dehydrogenase 1 (ALDH1), Notch1, and FoxM1, and reduced sphere formation in ovarian cancer cells. Using microarray analyses, immunoblotting, luciferase activity, and chromatin immunoprecipitation (ChIP) assays, we showed that SR-T100 suppressed the expression of c/EBPβ and COL11A1, and its promoter activity, in resistant cells, but not sensitive cells. SR-T100, paclitaxel, and cisplatin inhibited the growth of A2780CP70 cells in mouse xenografts, as compared to the vehicle control, and the combination of cisplatin and SR-T100 was more effective than either treatment alone. SR-T100 may represent a potential therapeutic adjunct to chemotherapy for ovarian cancer treatment.

Key words: Solanum incanum, aldehyde dehydrogenase 1, ovarian cancer, stemness, chemoresistance

Introduction

Epithelial ovarian carcinoma (EOC) is the most lethal gynecological malignancy (1). The initial symptoms are vague, and the majority of women have advanced disease at the time of diagnosis. The standard treatment is aggressive cytoreductive surgery, followed by platinum-taxane based chemotherapy. After therapy, a significant number of patients eventually relapse and develop chemoresistance, leading to a very low overall survival rate. Despite the incorporation of new chemotherapies for EOC treatment, only a small increase in overall survival has been achieved (2). Additionally, second-line
chemotherapeutics can cause substantial adverse effects in patients (3). This highlights a need for novel therapeutic approaches to improve EOC treatment.

One approach to overcome this problem is the development of new agents that can be used in combination with existing chemotherapeutics to yield better results than chemotherapeutics alone (4). Accumulating evidence suggests that many natural products, including extracts and isolated chemicals, have the potential to interact with multiple targets in the signaling pathways that regulate cancer progression (5). Therefore, a systemic study of natural products is needed to define their antitumor effects and understand their mechanisms of action to develop new treatments.

The anticancer effects of Solanum species have been known for centuries (6-8). Several active ingredients, including solamargine, solasodine, and solasonine, suppress cancer growth in vitro and in vivo (7-10). Solamargine is the major glycoalkaloid in Solanum incanum. Previous work showed solamargine increases the expression of tumor necrosis factor receptors (TNFR1 and TNFR2)(11-14) and activates the mitochondrial apoptotic pathway in human hepatocellular carcinoma (HepG2 and Hep3B), lung cancer (A549, H441, H520, H661, and H69)(15, 16), and breast cancer (SK-BR3, MCF-7, HBL-100, and ZR-75-1) cell lines (17, 18). Compared with paclitaxel, cisplatin, gemcitabine, and etoposide, solamargine has a superior ability to suppress human lung cancer cell growth (16). Additionally, solamargine inhibits the growth of various cultured human solid tumor cell lines, including HT-29 and HCT-15 (colon), LNCaP and PC-3 (prostate), and T47D and MDA-MB-231 (breast)(19). Recently, S. incanum extract (SR-T100), which is primarily composed of solamargine alkaloid, was shown to induce cutaneous squamous cell carcinoma apoptosis via activation of TNFRs and the mitochondrial apoptotic pathway, curing this cancer in hairless mice and actinic keratosis in human patients (19). Additionally, our preliminary findings showed that SR-T100 exhibits similar killing effects in chemoresistant cancer cells and their parental chemo-naive cells, indicating that SR-T100 may overcome chemoresistance. However, the mechanisms by which SR-T100 modulates chemoresistance remain unclear. In this study, the novel molecular mechanisms underlying SR-T100-regulated chemoresistance were elucidated, providing an understanding of its mechanism of action.

**Materials and Methods**

**Cell lines and cell culture**

The immortalized normal human ovarian surface epithelial cell line, IOSE-398, and human ovarian cancer cell lines ES2, TOV-21G, IGROV1, A2780, A2780CP70, ov2008, and ov2008CP20 were obtained from the American Type Culture Collection (ATCC; Manassas, VA, USA). IOSE-398 and TOV-21G cells were grown in MCDB105/M199ES2 medium with 10% fetal calf serum (FCS). ES2 cells were grown in McCoy’s 5A medium supplemented with 10% FCS. All other cells were grown in RPMI-1640 medium with 10% FCS. Cells were grown at 37°C in a 5% CO₂ atmosphere. Cells were cultured and stored according to the supplier’s instructions and used between passages 5 and 20. Cell lines were routinely authenticated once every 6 months through cell morphology monitoring, growth curve analysis, species verification by isoenzymology and karyotyping, identity verification using short tandem repeat profiling analysis, and contamination checks.

**Western blot analysis**

Cells were washed in PBS, and then lysed in lysis buffer containing 50 mM Tris-HCl (pH 7.5), 150 mM NaCl, 1 mM EDTA, 1 mM MgCl₂, and 0.5% Triton X-100. Lysates were cleared by centrifugation at 13000 g for 20 min at 4°C and analyzed by western blot analysis. Protein samples were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), transferred to a polyvinylidene difluoride membrane, and probing with the indicated antibodies. Proteins were detected by chemiluminescence.

**Antibodies and reagents**

Western blotting was performed using the following antibodies at the indicated dilutions: anti-aldehyde dehydrogenase 1 (ALDH1; 1:1000; BD Biosciences; Franklin Lakes, NJ, USA), anti-Notch1 (1:1000), anti-COL11A1 (1:1000; Abcam; Cambridge, UK), anti-FoxM1 (1:1000; Santa Cruz Biotechnology; Santa Cruz, CA, USA), and anti-β-actin (1:5000; Sigma; St. Louis, MO, USA). Cisplatin (Fresenius Kabi Oncology Ltd, Haryana, India) and paclitaxel (Corden Pharma Latina S.P.A., Sermoneta, Italy) were kindly provided by the Cancer Center at National Cheng Kung University Hospital, Tainan, Taiwan. SR-T100 isolated from S. incanum was provided by G&E Herbal Biotechnology Company (Tainan, Taiwan).
Sphere formation assay

Standard sphere formation assays were performed according to Zhang et al. (21) with minor modifications. Cells (1 × 10^6) were resuspended in serum-free DMEM/F12 medium supplemented with 5 μg/mL insulin (Sigma), 20 ng/mL human recombinant epidermal growth factor (EGF; Invitrogen, Life Technologies, Carlsbad, CA, USA), and 10 ng/mL basic fibroblast growth factor (Invitrogen) in ultra-low attachment plates (Corning Costar, Corning, NY, USA). Spheres that arose within 1–2 weeks were counted. Colony diameters > 50 μm were counted as a single positive colony. The middle field was chosen to count spheres, and two fields for each plate were counted under a dissecting microscope. For all sphere formation experiments, a minimum of eight wells were run for each condition. All data represent the mean ± SEM of three separate experiments and at least 24 different fields.

Microarray analysis and data processing

Total RNA was isolated using the RNeasy Micro Kit (Qiagen; Valencia, CA, USA), quantified, and checked for quality with a Bioanalyzer 2100 system (Agilent; Palo Alto, CA, USA). Labeled probes for Affymetrix DNA microarray analysis were prepared according to the manufacturer’s instructions. Biotin-labeled cRNA, produced by in vitro transcription, was fragmented and hybridized to Affymetrix U133A GeneChip arrays (22,283 probe sets, Santa Clara, CA, USA) at 45°C for 16 h, and then washed and stained using GeneChip Fluidics. The arrays were scanned to a target intensity of 500 by a GeneArray Scanner, and patterns of hybridization were detected as light emitted from the fluorescent reporter groups incorporated into the target and hybridized to oligonucleotide probes. Image acquisition and probe quantification was performed using Affymetrix GeneChip Operating Software. Microarray quality control was performed using R package gcrma (22). The gcrma function in the R package affy was applied to normalize the CEL files using the RMA method (23). Data are MIAME compliant and have been deposited in the Gene Expression Omnibus.

Plasmid construction and site-directed mutagenesis

The COL11A1 polymerase chain reaction product was cloned into the KpnI and XhoI sites in the pGL4 vector. The resulting construct was confirmed by DNA sequencing. The COL11A1 promoter deletion construct, COL11A1-541/-203, was similarly generated using the COL11A1-541/+1 construct as a template.

Luciferase reporter assay

Luciferase assays were conducted using a luciferase reporter assay system (Promega; Madison, WI, USA) 48 h after transfection. Normalized luciferase activity is reported as luciferase activity/β-galactosidase activity.

Chromatin immunoprecipitation (ChIP) assay

Native protein-DNA complexes were cross-linked by treatment with 1% formaldehyde for 15 min. The ChIP assay was performed as previously reported (24). Briefly, equal aliquots of isolated chromatin were subjected to immunoprecipitation with anti-c/EBPβ and IgG monoclonal antibodies.

Xenograft animal model

All animal procedures were reviewed and approved by the Institutional Animal Care and Use Committee at National Cheng Kung University (No. 102267). Female 6-week-old NOD-SCID mice (Charles River Laboratories) were subcutaneously implanted in the flank with A2780CP70 cells (1 × 10^6 cells/100 μL). Tumor dimensions were measured 2 to 3 times per week, and volume was calculated as length × width × height × 0.52. Once tumors reached 20 mm^3, the mice were randomly assigned to 10 groups (n = 6). Animals in each group received the same volume of saline, cisplatin, or SR-T100 in 200 μL by intraperitoneal injection. Treatment frequency was 1 time per day for SR-T100 and 1 time per 3 days for cisplatin. Tumor growth, tumor imaging, and body weights were determined, as described (25). The mice were sacrificed by CO2 inhalation and xenograft tumor tissues were excised.

Statistical analysis

Statistical analysis was performed using the Statistical Package for the Social Sciences software, version 17.0, for Windows (SPSS Inc.). Frequency distributions between categorical variables were compared using the Pearson chi-square test and Fisher’s exact method. Student’s t-test and Mann-Whitney U test were used to compare two groups with normally and non-normally distributed interval data, respectively. One-way ANOVA and Kruskal-Wallis tests were used compare three or more groups with normally distributed and non-normally distributed interval data, respectively.

Results

SR-T100 sensitizes chemoresistant cells to cisplatin and paclitaxel

To illustrate the anticancer effects of SR-T100 in ovarian cancer, we first evaluated SR-T100-induced cytotoxicity using the MTT assay in ovarian cancer
cells and IOSE-398 cells. As shown in Fig. 1A, the concentration of SR-T100 that reduced cell viability 50% (IC₅₀) was lower in malignant cells (IC₅₀ of ES2 = 12.83 µg/ml; IC₅₀ of TOV-21G = 14.88 µg/ml; IC₅₀ of IGROV1 = 14.69 µg/ml) than in nonmalignant cells (IC₅₀ of IOSE-398 = 27.73 µg/ml) (Fig. 1A), suggesting that SR-T100 is less toxic in IOSE-398 cells. The effect of SR-T100 on chemosensitivity was examined in the following paired ovarian cancer cell lines: A2780 and A2780CP70, and ov2008 and ov2008CP20. Chemoresistant cells were less sensitive to cisplatin than chemosensitive cells (Fig. 1B). The IC₅₀ values for SR-T100 treatment in resistant cells (IC₅₀ of A2780CP70 = 12.62 µg/ml; IC₅₀ of ov2008CP20 = 15.94 µg/ml) were similar to those in sensitive cells (IC₅₀ of A2780 = 11.91 µg/ml; IC₅₀ of ov2008 = 15.25 µg/ml) (Fig. 1C). These results suggest that SR-T100 is cytotoxic in both chemosensitive and chemoresistant ovarian cancer cells. More importantly, the IC₅₀ values in TOV-21G cells decreased after treatment with SR-T100 in combination with various concentrations of cisplatin or paclitaxel, as compared with cisplatin or paclitaxel treatment alone (Fig. 1D). These results suggest that SR-T100 sensitizes chemoresistant cells to cisplatin and paclitaxel.

Figure 1. SR-T100 treatment increases cell sensitivity to cisplatin and paclitaxel. Cell viability following SR-T100 (A), cisplatin (B), or combined (C) treatment was measured by MTT assay. (D) Cell viability following combination treatment with SR-T100 and cisplatin or paclitaxel was measured by MTT assay. All experiments were performed in triplicate.
SR-T100 inhibits stem cell marker expression and sphere formation in ovarian cancer cells

Recent studies have shown that subpopulations of cancer cells, cancer stem-like cells (CSCs), are characterized by their enhanced tumor formation ability and drug resistance. CSCs share some markers with stem cells, including ALDH1 and Notch1 (21, 26, 27). Additionally, our recent work has indicated that FoxM1 regulates epithelial-mesenchymal transition, stemness, and chemoresistance in epithelial ovarian carcinoma cells (28). Thus we evaluated whether ALDH1, Notch1, and FoxM1 were inhibited by SR-T100. As shown in Figure 2A, SR-T100 dose-dependently downregulated the expression of ALDH1, Notch1, and FoxM1 in chemoresistant A2780CP70 cells. This decreased expression was not observed in chemosensitive A2780 cells. Sphere formation in A2780CP70 cells was also dose-dependently inhibited by SR-T100 treatment (Fig. 2B).

Downregulation of c/EBPβ and COL11A1 levels by SR-T100 in chemoresistant ovarian cancer cells by expression profiling

To identify the genes regulated by SR-T100, A2780CP70 cells were treated with 2.5 or 5 µg/ml SR-T100, and a gene expression array was performed. Microarray analysis showed that 197 genes (66 up-regulated and 131 downregulated) were differentially expressed between SR-T100-treated and control cells (Tables 1 and 2). Among these genes, c/EBPβ and COL11A1 were among the most downregulated genes. Our recent findings show that COL11A1 plays a dual role in EOC tumor progression and chemoresistance, and that the c/EBPβ binding site on the COL11A1 promoter is a major determinant of cisplatin- and paclitaxel-dependent COL11A1 activation (29). Our results showed that the expression of c/EBPβ and COL11A1 was enhanced by cisplatin, and that this elevated expression was inhibited by SR-T100 in chemoresistant A2780CP70 cells. In contrast, the expression level of c/EBPβ and COL11A1 was largely unaffected in chemosensitive A2780 cells by either cisplatin or SR-T100 treatment (Fig. 3A). In addition, the increased promoter activity by cisplatin in the region between -541 and -203 on COL11A1, which is important for the transcriptional regulation by cisplatin and paclitaxel (29), was decreased in SR-T100-treated A2780CP70 cells, but not in A2780 cells (Fig. 3B). ChIP assays further indicated that the binding of c/EBPβ to the COL11A1 promoter was inhibited by SR-T100 alone and SR-T100 plus cisplatin (Fig. 3C).

Figure 2. SR-T100 treatment reduces stem cells marker expression and sphere formation in chemoresistant ovarian cancer cells. (A) Cells were treated with SR-T100 (2.5 or 5 µg/ml) for 48 h, and cell lysates were collected for western blotting. (B) Cells were treated with SR-T100 (2.5 or 5 µg/ml) and were evaluated by sphere formation assays. Representative images show spheres generated from single-cell cultures after 10 days. The lower panel depicts the relative sphere formation ratio (scale bar = 200 µm).
Figure 3. SR-T100 downregulated c/EBPβ and COL11A1 in chemoresistant ovarian cancer cells. (A) Left panel: Cells were treated with SR-T100 (2.5 or 5 µg/ml) for 48 h, and cell lysates were collected for western blotting. Right panel: COL11A1 and c/EBPβ protein expression in cells treated with SR-T100 (5 µg/ml) and cisplatin (10 µM) for 48 days. (B) A2780CP70 cells transfected with the indicated COL11A1 promoter constructs were treated with SR-T100 and cisplatin for 48 h. The luciferase activity was measured and normalized to Renilla luciferase activity. All experiments were performed in triplicate. (C) A ChIP assay was performed to evaluate the binding of c/EBPβ to the COL11A1 promoter after treatment of A2780CP70 and A2780 cells for 48 h with varying cisplatin concentrations.

Table 1. The 66 genes upregulated in response to SR-T100 treatment in A2780CP70 cells.

| Gene             | Log2 ratio | Gene             | Log2 ratio |
|------------------|------------|------------------|------------|
| HSPA1B           | 2.94       | FABP5L2          | 1.22       |
| LOC399988        | 2.48       | M6PRBP1          | 1.22       |
| HMOS1            | 2.26       | C7ORF68          | 1.22       |
| VGF              | 2.15       | GBPDA1           | 1.21       |
| HSPA8            | 1.99       | FEN1             | 1.20       |
| TFRC             | 1.94       | DNAI1            | 1.20       |
| SNOAR61          | 1.76       | B4GALT1          | 1.20       |
| RPL29            | 1.72       | RAB7L1           | 1.19       |
| IL3              | 1.63       | MDF1             | 1.19       |
| TMEM158          | 1.54       | SLCO5R2          | 1.19       |
| C18ORF55         | 1.52       | RPL36A           | 1.18       |
| HS1D7B1          | 1.51       | LOC85389         | 1.17       |
| MED1             | 1.49       | CKS2             | 1.17       |
| ELOF1            | 1.48       | TAGLN            | 1.15       |
| MA2EL1           | 1.43       | MCM10            | 1.14       |
| LOC100128928     | 1.41       | LOC433567        | 1.14       |
| MRPL36           | 1.38       | MRPL34           | 1.13       |
| FHL2             | 1.38       | NT5B             | 1.13       |
| C6orf156         | 1.38       | UTP11L           | 1.11       |
| CCDC58           | 1.36       | LOC535506        | 1.09       |
| SFA3A            | 1.34       | MBD5             | 1.09       |
| CCDC56           | 1.31       | CDC25A           | 1.09       |
| ITPKA            | 1.31       | C7orf30          | 1.08       |
| TUBB2C           | 1.29       | RABGCTB          | 1.07       |
| SERPIN11         | 1.27       | TSPAN9           | 1.06       |
| LC3C             | 1.27       | LRRK2            | 1.05       |
| SPHK1            | 1.26       | ZC3HAV1          | 1.05       |
| C6orf115         | 1.26       | MAGOH            | 1.04       |
| LSM3             | 1.26       | TM4SF4           | 1.03       |
| AFI3             | 1.25       | EIF4E2           | 1.02       |
| C7orf79          | 1.24       | RPL34            | 1.01       |
| FLJ35767         | 1.23       | FAM103A1         | 1.00       |
| MMP1             | 1.23       | UNG              | 1.00       |

SR-T100 increases cisplatin sensitivity in mouse xenografts

To determine whether SR-T100 could suppress tumor growth, mice were injected subcutaneously with $1 \times 10^6$ A2780CP70 cells and treated with intraperitoneal injection of SR-T100, with or without adding cisplatin (Figs. 4A and 4B) or paclitaxel (Figs. 4D and 4E). When compared to the vehicle controls, single treatment with cisplatin ($P = 0.002$), varying SR-T100 doses (SR-T100 2.5 mg/kg, $P = 0.003$; SR-T100 5.0 mg/kg, $P = 0.002$), or paclitaxel (paclitaxel 2.5 mg/kg, $P = 0.001$; paclitaxel 5.0 mg/kg, $P = 0.001$; paclitaxel 10 mg/kg, $P = 0.001$) significantly inhibited the growth of A2780CP70 cells in mouse xenografts. Moreover, combinations of cisplatin and SR-T100 (SR-T100 2.5 mg/kg + cisplatin 3 mg/kg vs. controls, $P = 0.002$; SR-T100 5.0 mg/kg + cisplatin 3 mg/kg vs. controls, $P = 0.002$) significantly reduced tumor volumes, as compared to the controls. The tumor suppression effect was greater with combination treatment (SR-T100 5 mg/kg and cisplatin 3 mg/kg) than with SR-T100 alone (5 mg/kg alone; $P = 0.046$). An effect was also found between the combination group and cisplatin alone group ($P = 0.032$). However, SR-T100 did not significantly increase the sensitivity to paclitaxel in chemoresistant cells (Fig. 4D). The body weight of animals receiving cisplatin, paclitaxel,
SR-T100, or combined treatment remained relatively unaltered, suggesting negligible toxicity was observed (Figs. 4C and 4F). These results further confirmed that SR-T100 has the potential to kill chemoresistant cancer cells.

Table 2. The 131 genes downregulated in response to SR-T100 treatment in A2780CP70 cells.

| Gene        | Log2 ratio | Gene          | Log2 ratio |
|-------------|------------|---------------|------------|
| FGFR1       | 2.94       | CTORF24       | 1.57       |
| DDIT4       | 2.92       | DDT3         | 1.55       |
| AT1F3       | 2.72       | ZNF627        | 1.55       |
| IN1BR       | 2.69       | ITGA4         | 1.55       |
| TXNIP       | 2.29       | SLC9A14       | 1.54       |
| TSC22D3     | 2.13       | LOC727758     | 1.54       |
| SLC38A2     | 2.06       | IRE2BP2       | 1.53       |
| ARHGEF2     | 2.04       | WARS          | 1.52       |
| FOX03       | 2.03       | CBLB          | 1.52       |
| DDX2        | 2.03       | SLC6A9        | 1.51       |
| PCK2        | 2.01       | TUBB1         | 1.51       |
| ASNS        | 1.90       | SLC38A1       | 1.50       |
| CEBPB       | 1.86       | MUC15         | 1.50       |
| HIF1        | 1.84       | SLC3A2        | 1.49       |
| JIF2        | 1.79       | TRIB1         | 1.49       |
| LOC729779   | 1.79       | TNFRSF19      | 1.49       |
| SLC7A5      | 1.75       | SLC7A1        | 1.47       |
| ITG3A       | 1.73       | CHD9          | 1.46       |
| IL1R1       | 1.71       | HOXG1         | 1.46       |
| NTSC2       | 1.71       | PBRM1         | 1.45       |
| CICL4       | 1.71       | GARS          | 1.45       |
| PSIP1       | 1.70       | CLDN1         | 1.43       |
| PDE4D       | 1.69       | RORC         | 1.43       |
| COG8P1      | 1.69       | PPKDC         | 1.43       |
| AKT2        | 1.69       | CEBPA         | 1.43       |
| SSBX2F1     | 1.68       | TCEA1         | 1.42       |
| FAM129A     | 1.66       | UHRF1BP1      | 1.42       |
| BTG1        | 1.66       | HNRNPD1       | 1.41       |
| PAUI1       | 1.63       | LOC630103     | 1.39       |
| COL11A1     | 1.63       | TNFRSF10B     | 1.38       |
| CALCRL      | 1.62       | GPT2          | 1.37       |
| CNTNAP1     | 1.62       | LRBA          | 1.37       |
| PPP1I15A    | 1.57       | AARS          | 1.37       |
| EVL5L       | 1.57       | KLETD3        | 1.17       |
| TSC30B      | 1.36       | SRSN2         | 1.16       |
| LONP1       | 1.36       | SERPINF1      | 1.16       |
| SREBF1      | 1.35       | SMARCA1       | 1.16       |
| MTHFD2      | 1.35       | TRUB1         | 1.16       |
| PFKCDH      | 1.34       | SARS          | 1.15       |
| ULK1        | 1.34       | VEZT          | 1.14       |
| EPIS        | 1.34       | XBP1          | 1.14       |
| CALD1       | 1.33       | MLT1T1        | 1.14       |
| LOC723242   | 1.33       | KDM5B         | 1.12       |
| SPRY2       | 1.30       | ZZ53          | 1.12       |
| CNOT1       | 1.30       | ZCCHC8        | 1.12       |
| FBXO11      | 1.29       | YY1AP1        | 1.11       |
| KLGF        | 1.29       | ASIP1         | 1.09       |
| GRP5D2      | 1.29       | TNFSF19       | 1.09       |
| ULBP1       | 1.28       | SCRNI          | 1.09       |
| GPR126      | 1.28       | CHAC1         | 1.08       |
| COL3A1      | 1.26       | PLK2          | 1.08       |
| LOC440341   | 1.23       | BNP2          | 1.08       |
| LOC728734   | 1.23       | SERINC1       | 1.06       |
| HERPUD1     | 1.22       | ZFP90         | 1.05       |
| C10orf58    | 1.22       | ETV3          | 1.05       |
| STARD4      | 1.22       | FASN          | 1.05       |
| ZNF643      | 1.22       | UBR4          | 1.05       |
| TSC22D1     | 1.22       | AKNA          | 1.04       |
| C1orf84     | 1.21       | ACL1          | 1.04       |
| SEZ6L2      | 1.20       | IARS          | 1.04       |
| ERRF11      | 1.19       | BRSK1         | 1.03       |
| RAD21       | 1.18       | ABCA1         | 1.02       |
| SLC3A2      | 1.18       | KATNB1        | 1.02       |
| C1orf2p2    | 1.18       | SLCAT5        | 1.02       |
| ZNF627      | 1.18       | DPN5          | 1.01       |
| LOC440333   | 1.17       | LOC650215     | 1.01       |
| DYNC1H1     | 1.17       |                |            |

Discussion

Solamargine effectively induces apoptosis in various cancer cell lines. In contrast, similar concentrations of solamargine did not cause apoptosis in normal cells, such as bone marrow cells, fibroblasts, and normal hepatocytes, suggesting it preferentially kills malignant cells (13-18, 30). Previous studies indicate that solamargine promotes cell death via the activation of TNFRs (11-14) and the mitochondrial apoptotic pathway (15-18). Recently, SR-T100 was shown to induce cutaneous squamous cell carcinoma cells apoptosis via similar mechanisms and suppresses tumor growth in mouse xenografts and patients with actinic keratoses (20). In this study, we reveal the novel molecular mechanism underlying SR-T100-induced cytotoxicity, via suppression of ALDH1-mediated stemness and COL11A1-regulated chemoresistance, thereby suppressing tumor growth in ovarian cancer cells.

Recent studies have shown that CSCs, which are characterised by enhanced tumour formation ability and drug resistance, share some stem cell markers, including ALDH1 and Notch1 (21, 26, 27). Our recent report demonstrates that ALDH1 directly regulates the functions of ovarian cancer cells. ALDH1 expression is closely associated with tumorigenic potential in ovarian cancer cell lines, and FoxM1 and Notch1 are important downstream effectors required for ALDH1-induced cancer stemness (31). Additionally, we further showed that FoxM1 regulates epithelial-mesenchymal transition, stemness, and chemoresistance in epithelial ovarian carcinoma cells (28). In the present study, SR-T100 inhibited CSC marker and FoxM1 expression and sphere formation in ovarian cancer cells. These results indicate that SR-T100 exerts killing effects in chemoresistant ovarian cancer cells and suppresses tumor growth in mouse xenografts.

Chemoresistance often causes patient death, due to a lack of effective treatments. Li et al. (32) reported that solamargine effectively triggers apoptosis in multidrug-resistant tumor cells, which is associated with actin disruption and downregulation of multiple drug resistance 1 expression. In addition to reduced stemness, we showed that SR-T100 regulates cell sensitivity to anticancer drugs via downregulation of c/EBPβ and COL11A1. Our recent report demonstrates that the activation of COL11A1 in ovarian cancer cells by cisplatin and paclitaxel confers chemoresistance by activating the Akt/c/EBPβ pathway and stabilizing phosphoinositol-dependent kinase-1 (29). In the present study, we provided the first evidence that SR-T100 enhances cell sensitivity to cisplatin and paclitaxel by downregulating the binding activity of c/EBPβ to COL11A1 promoter.
In addition to inducing apoptosis, SR-T100 has been shown to sensitize breast cancer cells to cisplatin and epirubicin (18). Liang et al. (16) reported that combination therapy using low concentrations of solamargine and the low-toxicity topoisomerase II inhibitor epirubicin synergistically enhanced cytotoxicity in non-small cell lung cancer cells. In the present study, SR-T100 increased the sensitivity of chemoresistant ovarian cancer cells to cisplatin and paclitaxel in vitro. Furthermore, combination treat-
ment using cisplatin and SR-T100 was more effective in inhibiting the growth of A2780CP70 cells in mouse xenografts than either therapeutic alone. Taken together, these results suggest that SR-T100 can enhance cell sensitivity to anticancer drugs and may have potential against chemoresistant ovarian cancer cells.

Acknowledgments

SR-T100 was a kind gift from Professor Kou-Wha Kuo, Research Center of Herbal Medicines, G&E Herbal Biotechnology, National Tainan Science Park, Tainan 744, Taiwan. This work was supported by the National Science Council under Grant Number: NSC 101-2325-B-006-022, 102-2314-B-006-081, and MOST 103-2314-B-006-069. This study was also supported by the Department of Health under Grant Number: 102TM1050, and in part by the Headquarters of University Advancement at the National Cheng Kung University, which is sponsored by the Ministry of Education, Taiwan, ROC.

Author Contributions

Conceived and designed the experiments: YHW and CYC. Performed the experiments: YHW, WTC and MJY. Analyzed the data: THC and YFH. Wrote the paper: YHW and CYC.

Competing Interests

The authors have declared that no competing interest exists.

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