Therapeutic Discovery

Growth Inhibition of Ovarian Tumor–Initiating Cells by Niclosamide

Yi-Te Yo1,2, Ya-Wen Lin2,3, Yu-Chi Wang1,2, Curt Balch5,6, Rui-Lan Huang1, Michael W.Y. Chan8, Huey-Kang Sytwu3, Chi-Kuan Chen3, Cheng-Chang Chang1, Kenneth P. Nephew5,6,7, Tim Huang9, Mu-Hsien Yu1, and Hung-Cheng Lai1,2

Abstract

A recent hypothesis for cancer chemoresistance posits that cytotoxic survival of a subpopulation of tumor progenitors drives the propagation of recurrent disease, underscoring the need for new therapeutics that target such primitive cells. To discover such novel compounds active against drug-resistant ovarian cancer, we identified a subset of chemoresistant ovarian tumor cells fulfilling current definitions of cancer-initiating cells from cell lines and patient tumors using multiple stemness phenotypes, including the expression of stem cell markers, membrane dye efflux, sphere formation, potent tumorigenicity, and serial tumor propagation. We then subjected such stem-like ovarian tumor-initiating cells (OTIC) to high-throughput drug screening using more than 1,200 clinically approved drugs. Of 61 potential compounds preliminarily identified, more stringent assessments showed that the antihelmintic niclosamide selectively targets OTICs in vitro and in vivo. Gene expression arrays following OTIC treatment revealed niclosamide to disrupt multiple metabolic pathways affecting biogenesis, biogenesis, and redox regulation. These studies support niclosamide as a promising therapy for ovarian cancer and warrant further preclinical and clinical evaluation of this safe, clinically proven drug for the management of this devastating gynecologic malignancy. Mutat Cancer Ther; 11(8); 1703–12. ©2012 AACR.

Introduction

Ovarian cancer is the most lethal gynecologic malignancy and the fifth-most cause of overall cancer death of women in developed countries (1). The majority of women diagnosed with advanced-stage epithelial ovarian cancer experience associated with the development of chemoresistance and platinum-resistant tumor recurrence, a pathology that is uniformly fatal (2). An increasingly accepted cancer stem cell hypothesis regards tumors as caricatures of normal organs, possessing a hierarchy of cell types, at various stages of aberrant differentiation, descended from precursor tumor-initiating cells (TIC) cells that are highly resistant to conventional cytotoxics (3, 4).

Several TICs have now been isolated, in ovarian and other solid tumor malignancies, and our group also successfully isolated ovarian TICs (OTIC) from human tumors that specifically express the surface markers CD44 (a hyaluronate receptor) and CD117 (c-Kit or stem cell factor receptor; ref. 5). Other reported OTIC surface markers include CD133, CD24, CD44, and Myd88 (6–8). Although this approach has proved successful for characterizing a number of solid tumor progenitor cells, shortcomings include uncertainty of the precise roles of the specific surface markers (with regard to tumor potency) and insufficient isolation of cells for use in drug screening assessments (9).

In addition to cell-surface markers, some marker-free methods have also been applied to enrich TICs. Similar to hematopoietic and other somatic tissue stem cells, TICs express various ATP-binding cassette (ABC) transporters, including MDRI, ABCB1, and ABCG2, which confer chemoresistance via active transmembrane drug export (10). To exploit this stem cell attribute, fluorescence-activated cell sorting (FACS) of a side population of fluorescent dye-excluding cells has been used to enrich numerous normal tissue stem cells and TICs (4, 11). The side population-to-stemness link is now well substantiated by its coexistence with numerous other commonly
exploited stem cell attributes. Those progenitor phenotypes include expression of self-renewal and pluripotency-conferring transcription factors, differentiation into multiple, diverse progeny cells, activation of embryonic signaling pathways, and potent in vivo tumor initiation, methods that our group and others have used in the assessment of ovarian tumor progenitors (5, 12, 13). Another feature of stem/progenitor cells is their ability to self-aggregate into substrate-independent spheres, architectural structures originally discovered to be highly enriched in multipotent neural stem cells (14). Subsequently, anchorage-independent sphere formation has now been used for enrichment of stem cells from normal breast (15) and skin (16) tissues, in addition to TICs of brain (17), pancreatic (18), colon (19), and ovarian (5, 13) cancers.

Traditionally, anticancer drug development has been based on the targeting of rapidly proliferating cells, rather than cells having a capacity for self-renewal and/or differentiation, which might be more effective against malignancies possessing a tumor cell hierarchy (4, 10). It is likely that ovarian cancer is one such malignancy, based on its frequent progression to chemo-resistance (20). For drug discovery targeting TICs, cell-surface marker-independent isolation offers distinct advantages in allowing relatively straightforward procedures and less biased selection of cells with specific markers. In this study, for greater stringency, we used 2 stemness selection methods, dye exclusion, and subpopulation of sphere-forming ability, while still obtaining sufficient numbers of ovarian TICs for high-throughput identification of their antagonists.

Materials and Methods

Cells and experimental animals

Human ovarian cancer cell lines OVCAR-3 and SKOV-3 were purchased from American Type Culture Collection (ATCC). The human ovarian cancer cell line A2780 and its cognate cisplatin-resistant CP70 were obtained in 2007 from Dr. Tim Huang’s lab (University of Texas Health Science Center, San Antonio, TX). Earlier passages of all cell lines were maintained in several cryovials, respectively, in liquid nitrogen in our laboratory. The cell lines were used by were tested by Bioresource Collection and Research Center (Hsinchu, Taiwan) for identity verification by DNA profiling of short tandem repeat (STR) sequences. Briefly, 15 STR loci were tested by using the Applied Biosystems AmpFLSTR Identifier Kit (Applied Biosystems Catalog # 4322988). DNA profiles were compared manually to the ATCC and European Collection of Cell Cultures database. Cells were stored and used within 3 months after resuscitation of frozen aliquots. All cell lines were cultured in RPMI-1640 (Invitrogen) supplemented with nonessential amino acid (Invitrogen), sodium pyruvate (Invitrogen), 10% FBS (Biological Industries), and different growth factors, respectively, and plated in standard cell culture dishes (Corning). For in vivo tumor xenograft studies of CP70sps cells, female nonobese dia-betic severe-combined immunodeficient (NOD/SCID) mice were obtained from the Laboratory Animal Center of the National Taiwan University (Taipei City, Taiwan). Six-week-old mice were used for experiments unless indicated otherwise. The rules of the Animal Protection Act of Taiwan were strictly followed and all animal procedures approved by the Laboratory Animal Care and Use Committee of National Defense Medical Center.

Side population spheroid cells

For side population analysis, 106 cells were harvested and stained with 50 μg/mL Hoechst 33342 (Sigma). The dye exclusion phenotype (21), via membrane efflux, was confirmed by an ABCG2-specific inhibitor, GF120918. Hoechst-stained cells were subjected to FACS using a FACSaria (BD Biosciences) for collection of dye-excluding (side population) cells, which were then cultured in Ultra Low attachment plates (Corning) with serum-free Dulbecco’s Modified Eagle’s Medium/F12 medium containing 5 μg insulin (Sigma), 0.4% bovine serum albumin (Sigma), 10 ng/mL basic fibroblast growth factor (Invi-trogen), and 20 ng/mL human recombinant EGF (Invitrogen). Side population cells were then observed for the stem cell phenotype of tumour sphere formation (22) under suspension-cultured conditions, and those spheroid-forming cells were then referred to as side population spheroid (SPS) cells. For differentiation studies, CP70sps cells were cultured in standard coated dishes (Corning) with normal media and 10% FBS.

Characterization of stem cell properties of CP70sps cells

Stem cell phenotypes of the parental CP70 and side population CP70sps cells were assessed by the expression of specific stem cell gene markers (Supplementary Table S1). SPS cells also subjected to intracytoplasmic proteins staining, following fixation by paraformaldehyde and permeation by Triton X-100. Cells were stained with antibodies against human OCT4 (Abcam), NANO G (Abcam), NESTIN (Abcam), ABCG2 (Santa Cruz Biotechnology) and then stained with fluorescence-conjugated secondary antibodies (The Jackson Laboratory) or assess the ALDH activity using ALDEFLUOR Kit (StemCell Technologies) according to the manufacturer’s instructions as described previously (23) and analyzed by flow cytometry (BD Biosciences). Cells subjected to surface marker staining were incubated with CD34, CD44, or CD133 fluorescence-conjugated monoclonal antibodies (Abcam), followed by flow cytometry analysis.

Xenograft transplantation of CP70sps cells

CP70sps (P0) cells were injected intraperitoneally into NOD/SCID mice. After tumor formation, tumor nodules were harvested, mechanically dissected, enzymatically digested, and cultured in serum-free suspension to form spheroids (P1) as previously described (5). NOD/SCID mice were inoculated intraperitoneally with various numbers of CP70, CP70sps passage 0 (P0), and CP70sps...
passage-1 (P1) cells. Recipients were monitored every week. Mice were euthanized when the abdomen showed swelling to observe ascites and tumor formations.

Large-scale screening of bioactive compounds

For primary screening, $1 \times 10^3$ CP70sps cells were seeded per well in a 96-well culture dish (Corning). Twenty-four hours later, cells were treated with 30 $\mu$mol/L of more than 1,200 compound from a LOPAC library (Sigma). Three days later after treatment, the viability of treated cells was measured by CellTiter-Glo Luminescent Cell Viability Assay Kit (Promega), according to the manufacturer’s instructions. Candidate compounds were then subjected to a secondary screen using a 10-fold lower dose (3 $\mu$mol/L), using 5,000 CP70sps cells seeded per well in 96-well culture dish for 24 hours.

In vitro chemosensitivity and apoptosis analysis

Ovarian cancer cells treated with various doses of cisplatin for 24 hours were used to define the chemoresistance capacity. The IC$_{50}$ value was calculated by the cell growth using MTS-based CellTiter 96 Cell Proliferation Assay Kit (Promega). For investigation of cytochrome c release and detection of short functional apoptosis-inducing factor (AIF) fragment, niclosamide-treated CP70sps cells were harvested 48 hours later and subjected to detection using FlowCellect Cytochrome C Kit (Millipore) according to the manufacturer’s instructions or to immunoblot using AIF antibody (Millipore) staining. CP70sps cells were treated with niclosamide for 72 hours and then total cells were harvested and stained with propidium iodide (PI). Samples were then analyzed for DNA content using a FACSCalibur (BD Biosciences), and relative cell-cycle distribution was analyzed using the CellQuest software (Verity Software House).

Preclinical therapeutic effects of niclosamide

NOD/SCID mice, 5 or 4 per group, were injected intraperitoneally with $1 \times 10^7$ CP70sps or $1 \times 10^6$ CP70 cells at day 0 and the experimental group then treated with niclosamide intraperitoneally (5 or 10 mg/kg/daily) for 3 weeks, followed by biweekly treatments for another 4 weeks. Mouse bodyweights were monitored every 3 days, and mice were euthanized to observe tumor nodule formation 7 weeks later, upon the development of ascites. Stem-like OTICs from patients were then peritoneally derived, as described previously (5). Five thousand peritoneally derived OTICs were then seeded in 96-well dish and treated with 3 $\mu$mol/L of niclosamide and viability detected 3 days later by an MTS-based assay, according to the manufacturer’s instructions. To test the therapeutic effects in vivo, NOD/SCID mice were injected intraperitoneally with $1 \times 10^3$ of CP70sps and then treated with niclosamide intraperitoneally (10 mg/kg/daily) for 3 weeks and then biweekly for another 12 weeks. Mice were euthanized upon development of measurable ascitic fluid, and tumor formation was assessed. All studies were approved by the Institutional Review Board of the Tri-Service General Hospital, National Defense Medical Center.

Quantitative RT-PCR array, global gene expression array, and bioinformatics

CP70 and CP70sps RNA was isolated by RNeasy Mini Kits (Qiagen), suspended in the RNase-free water, and following reverse transcription to cDNA, subjected to RT$^2$ Profiler PCR Array Systems (SABioScience) for detection of stem cell gene expression using an ABI 7500 Real-Time PCR System (Applied Biosystems). For global gene expression analysis, total RNA (2 $\mu$g) from niclosamide-treated CP70 and CP70sps were isolated after various periods (2, 4, and 6 hours) and then qualified and sent to a core service unit (Genetech Biotech Co.) for whole genome expression analysis. We detected gene expression profiles on Illumina HumanWG-6 v3.0 Expression Beadchip (Illumina). The gene expression data were deposited in Gene Expression Omnibus (GEO) DataSets (accession number GSE36259). After quintile normalization, the significance of analysis microarray (SAM) algorithm was used to statistically differentiate expression patterns using a MeV 4.6.2 tool (24). PathVisio 2.0.11 (25) was then used to determine functional pathways impacted by changes in gene expression patterns, using the WikiPathways and KEGG pathways database.

Statistical analyses

Data were expressed as mean ± SD. Differential chemoresistance in various cell clines was analyzed using the nonparametric Mann–Whitney test. P values less than 0.05 were considered to be statistically significant. All analyses were carried out using the statistical package in R (R version 2.13.0).

Results

Isolation and characterization of stem-like ovarian cancer cells

We first determined the side population fractions of a panel of OC cell lines by staining with Hoechst 33342 dye, followed by the isolation of Hoechst-negative cells (Fig. 1A). An ABC transport-specific inhibitor, GF120918, confirmed Hoechst dye efflux via active membrane transport (Supplemental Fig. S1A). Sphere-forming ability of the side population cells was then evaluated. CP70 SP cells were screened for spheroid formation (Supplementary Fig. S1B), and those cells were thus designated CP70sps. We next compared CP70sps and bulk culture CP70 cells for expression of various embryonic and previously reported cancer stemness markers. As shown in Fig. 1B, CP70sps cells upregulated numerous stem cell marker genes, including NANOG (5.9-fold), ALDH1 (2.2-fold), NESTIN (2.3-fold), and CD44 (1.3-fold). However, we did not observe CP70sps upregulation of CD133, a previously reported marker of ovarian cancer and/or tumor-associated endothelial, stem cells (6, 26–28). To more broadly assess gene expression in CP70sps cells, reverse
transcriptase PCR (RT-PCR) target arrays were used, and enrichment of several stem cell gene signatures, including the \textit{Wnt} signal cascade (Supplementary Table S1), was observed.

As high capacity for aberrant differentiation is a foundation for the cancer stem cell hypothesis (3, 4), we next assessed whether CP70sps cells lose their stemness phenotype when exposed to differentiation-associated conditions. As shown in Fig. 1C, those culture conditions resulted in a time-dependent decrease of side population cells, from 37.6%, after one week, to 1.3%, after 6 weeks, thus showing asymmetric cell division and differentiation potency in culture. Platinum chemoresistance was shown in CP70sps cell, with IC\textsubscript{50} dose sensitivities being 2.0- and 20.6-fold higher than the IC\textsubscript{50} values for unselected bulk CP70 cells and their A2780 cells chemosensitive parent cell line (Fig. 1D). Analogously, after 4 to 6 weeks under differentiating culture conditions, CP70sps cells lost their chemoresistance phenotype, with an approximately 40% loss of viability (Fig. 1E).
As the gold standard for the definitive identification of cancer stem–like cells is exponentially increased tumorigenic potential (3, 4), we directly assessed the tumorigenic capacity of CP70sps cells \textit{in vivo}. Currently, tumorigenicity of 100 to 5,000 cells is considered sufficient to define tumor-initiating ability, with prior studies showing 100 CD117\textsuperscript{+} (5), 100 to 500 CD133\textsuperscript{+} (6), 5,000 PKH\textsubscript{high/low} (29), 5,000 CD24\textsuperscript{+} (8), and 1,000 ALDH1\textsubscript{br/low} cells (23) to sufficiently indicate ovarian cancer stemness. In this study, 1,000 CP70sps and 100 CP70sps P1 cells grew tumors, fulfilling the current definition of TICs. We did not test the tumorigenicity of one single cell in this study. Indeed, the tumorigenicity also depends on the immune background of mice (30). To more precisely recapitulate the physiologic niche of ovarian cancer (5, 20), we injected serially diluted CP70sps cells intraperitoneally into recipient NOD/SCID mice. Although only one of 3 mice grew intraperitoneal cavity tumors when injected with 10\textsuperscript{4} and 10\textsuperscript{5} CP70 cells, 3 of 3 injections of only 10\textsuperscript{3} CP70sps stem-like cells were tumorigenic (Table 1). Intraperitoneal CP70sps xenograft tumors were then harvested and subcultured for 7 days to reform spheroids (passage P-1), which were subsequently dissociated and reinjected intraperitoneally into fresh mice. This serial transplantation resulted in tumors from injection of only 100 cells (3/3), and after just one such serial passage, we found tumor latency of 10\textsuperscript{3} CP70sps cells to decrease from 70 to 41 days, although the same number (10\textsuperscript{3}) of unselected CP70 cells were not tumorigenic, even at nearly twice that time period (Fig. 2A). Also, in accordance with another TIC requisite (reproducible recapitulation of the original tumor histology; refs. 3, 4), all xenograft tumors were similarly categorized as poorly differentiated grades 3 to 4 (Fig. 2B). On the basis of above results, CP70sps meet the currently accepted criteria for stem-like, OTICs and were then used in subsequent drug screening experiments.

**Table 1. Tumorigenicity of CP70 OVCA cells and CP70sps OTICs**

<table>
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<th>Cell type</th>
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<th>10\textsuperscript{6}</th>
<th>10\textsuperscript{5}</th>
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<td>1/3 (60 d)</td>
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<tr>
<td>CP70sps P0</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>3/3 (66 d)</td>
<td>3/3 (70 d)</td>
<td>–</td>
</tr>
<tr>
<td>CP70sps P1</td>
<td>–</td>
<td>–</td>
<td>3/3 (21 d)</td>
<td>3/3 (33 d)</td>
<td>3/3 (41 d)</td>
<td>3/3 (65 d)</td>
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Growth inhibition of CP70sps was used to screen a library of 1,258 bioactive compounds (LOPAC; Sigma), using the procedure summarized in Fig. 3A. That initial screen identified 61 anti-OTIC compounds (Fig. 3B). Further evaluation of those compounds in growth inhibition assays using lower concentrations resulted in the selection of 2 candidate anti-OTIC agents, of niclosamide and rottlerin, which were then tested using CP70sps and CP70 (Fig. 3C). Although both agents displayed activity, niclosamide was selected for further study, based on our findings of poor \textit{in vivo} activity (data not shown) and previous anticancer studies showing rottlerin to be a poor inhibitor of protein kinase-C isoforms (31). OTIC-selective targeting by niclosamide was further shown by growth inhibition of OTICs derived from CP70 cell line and...
OVARIAN CANCER PATIENTS (PERITONEALLY DERIVED OTICS) IN VITRO (Figs. 3D AND 4A).

TO ASSESS NICLOSMAMIDE ACTIVITY AGAINST TUMOR PROGENITOR CELLS IN VIVO, GROUPS OF MICE (N = 3, EACH GROUP) WERE INOCULATED WITH 1 × 10⁶ CP70Sps CELLS INTRAPERITONEALLY AND THEN TREATED WITH NICLOSMAMIDE (10 mg/kg/daily), ADMINISTRATION TRIPLETHELONALLY, STARTING ONE DAY AFTER OTIC INOCULATION. BODY WEIGHTS REMAINED CONSTANT DURING NICLOSMAMIDE TREATMENT (SUPPLEMENTARY FIG. S2), SHOWING A LACK OF SYSTEMIC TOXICITY, WHEREAS BODY WEIGHTS OF MOCK-TREATED MICE WERE SIGNIFICANTLY INCREASED BECAUSE OF ABDOMINAL TUMOR GROWTH (FIG. 4B). AT LOWER DOSES (5 mg/kg/daily), NICLOSMAMIDE INHIBITED OTIC GROWTH WITHIN THE OVARY AND PARTIALLY INHIBITED GROWTH OF OTICS IN THE PERITONEAL CAVITY (FIG. 4C). WE ALSO EXAMINED WHETHER NICLOSMAMIDE POSSESSED ACTIVITY AGAINST OTIC IN THE PRESENCE OF MORE DIFFERENTIATED TUMOR CELLS AND OBSERVED SIMILAR TUMOR GROWTH INHIBITION OF CP70 CELL Xenografts IN MICE TREATED WITH NICLOSMAMIDE (SUPPLEMENTARY FIG. S3).

DISRUPTION OF OTIC METABOLISM BY NICLOSMAMIDE

Genome-wide gene expression array was subjected to elucidate the mechanism of niclosmamide-induced cell growth inhibition of our cancer stem-like OTICS (R2C1). Significant changes of gene expression in 2,928 genes were identified after niclosmamide treatment for different time periods (Fig. 5A). Because uncoupling of mitochondrial oxidative phosphorylation is believed to be its anti-helminthic mechanism of action (32), we hypothesized that niclosamides antagonistic effects on OTICS could, in part, be due to its disruption of metabolism. Consequently, we focused on genes participating in metabolic pathways (oxidative phosphorylation, glycolysis, and fatty acid biosynthesis; Fig. 5B). Our results showed that genes participating in protein complexes of oxidative phosphorylation were downregulated. The shift from oxidative phosphorylation to aerobic glycolysis (i.e., the Warburg effect) is the best characterized metabolic phenotype of cancer (33) and hypothesized to divert carbon into biosynthetic pathways required for high rates of cell proliferation (34). Although gene array results can be informative, functional validation (including enzymatic metabolic assessments) will be necessary for further conclusions with regard to niclosamide metabolic effects.

IN PARTICULAR, THE M2 ISOFORM OF PYRUVATE KINASE (PK-M2), EXPRESSED IN NEARLY ALL CANCEROUS BUT NOT NORMAL CELLS...
PK-M2 facilitates the conversion of phosphoenolpyruvate to pyruvate, thus regulating glycolytic intermediates for possible diversion into biosynthetic pathways. Interestingly, enzymes important in these biosynthetic pathways were downregulated after niclosamide treatment. Oxidative stress is another important metabolic regulator and as shown in Fig. 6A, niclosamide treatment resulted in a more than 20% increase in reactive oxygen species (ROS) in cultured OTICs. As elevated ROS is a predominant inducer of mitochondrial apoptosis, we subsequently showed that niclosamide treatment significantly induced apoptosis in human OTICs, as shown by 3 different assays: (i) cytosolic cytochrome C release; (ii) cleavage of AIF1; and (iii) DNA fragmentation determined by PI staining (Fig. 6B and D). Taken together, niclosamide disrupts bioenergetics, biogenesis, and redox regulation in cancer stem–like OTICs.

Discussion

In this study, we carried out a drug screening approach for compounds with activity against cancer stem–like cells designated as OTICs. We discovered that niclosamide, which has proved to be safe and effective for the past 2 decades against numerous parasites, inhibited OTIC growth both in vitro and in vivo. Our results further suggest that niclosamide represses metabolic enzymes responsible for bioenergetics, biosynthesis, and redox regulation specifically in OTICs, presumably leading to mitochondrial intrinsic apoptosis pathways, loss of tumor stemness, and growth inhibition. In apoptosis cascades, although we found only minor mitochondrial release of cytochrome C, marked release of AIF was observed, suggesting that AIF plays a major role in niclosamide-associated apoptosis. However, other mechanisms leading to cell death cannot be excluded. Our study further strengthens the concept for therapeutic targeting of cancer stem cells in solid tumors.

Niclosamide is believed to inhibit mitochondrial oxidative phosphorylation (32), and recent research indicates that targeting mitochondria could be an efficient strategy to cancer chemotherapy (36). Niclosamide was reported to inactivate NF-κB, causing mitochondrial damage and the generation of ROS, leading to apoptosis of leukemic stem cells (37). In another study, both rottlerin and niclosamide were identified in a screen for mTOR-signaling inhibitors (38). In addition, the tuberous sclerosis complex (TSC)–mTOR was reported to maintain stemness properties of HSCs by inhibiting mitochondrial biogenesis and ROS levels (39), implying that mTOR inhibitors (such as niclosamide) may interfere with mitochondria and various metabolic pathways in TICs via disruption of antioxidant responses.

Although mTOR is an important contributor to cancer metabolism (33), in general, the metabolism associated with cancer stemness remains largely unexplored. However, a recent study in lung cancer suggests that TICs possess high mitochondrial membrane potential, low glucose/oxygen consumption, and low ROS and ATP content (40). Thus, interfering with energy production, biosynthesis, and oxidative stress regulation may represent a potential approach for targeting TICs. Compared with our study, biosynthetic diversion of glycolytic intermediates was not examined, and ROS elevation did not evoke apoptosis, but rather autophagy, ERK activation, and caspase-independent cell death (partially mediated by the energy sensor AMPK; ref. 41). Our results, for the first time, show that repression of genes encoding...
multiple metabolism enzymes leads to mitochondrial intrinsic apoptosis pathways and inhibition of tumor stemness. As the metabolism of TICs may be different from the bulk population of tumor cells and normal tissues, it may be possible to achieve a therapeutic window for niclosamide, with TIC-specific cytotoxicity and low normal cell toxicity (42, 43).

Embryonic signaling pathways such as Wnt, Hh, and Notch pathways have also been proposed as TIC therapeutic targets (see review ref. 44). We observed Wnt hyperactivity in OTICs, in agreement with previous hypotheses of Wnt inhibitor effectiveness as an ovarian cancer therapy (45). In addition, niclosamide has now been independently identified in screens for Wnt inhibitors (46), via the promotion of Frizzled-1 endocytosis, downregulation of Dishevelled-2, and inhibition of Wnt3A-induced β-catenin stabilization (40, 46), in addition to downregulation of the Wnt/β-catenin target oncoproteins survivin and c-Myc (47, 48). Likewise, we also observed the downregulation of multiple Wnt ligands/receptors and Wnt target genes survivin and c-Myc (data not shown). Several components of another embryogenesis-associated pathway strongly implicated in ovarian carcinogenesis, the cell-to-cell signaling pathway Notch (8), were also suppressed by niclosamide (data not shown). These results agree with another recent niclosamide study in leukemia (49), and it has been widely hypothesized that disruption of Notch signaling may represent a highly effective therapy for ovarian and other solid tumors, via its essentiality to maintaining TIC stemness (50).

In summary, we show that niclosamide has strong inhibitory activity against CP70bps and primary OTICs directly explanted from patient tumors. As a clinically approved drug, further extension of niclosamide to clinical trials may be warranted, allowing clinical assessment of the concept of targeting OTICs, in human ovarian cancer patients. This proof-of-principle study strongly strengthens the rationale for developing highly effective TIC-targeted therapeutics for the management of highly drug-resistant solid tumors, such as epithelial ovarian cancer.

Figure 5. Genome-wide gene expression analysis in niclosamide-treated OTICs. A, the 2,928 genes with significant changes after niclosamide treatment for 2, 4, and 6 hours were shown. The red indicates activation and the green indicates suppression. B, genes participating in metabolism including fatty acid biosynthesis, glycolysis and gluconeogenesis, and oxidative phosphorylation were highlighted.
Disclosure of Potential Conflicts of Interest
The patent of new indications for cancer therapy using niclosamide is pending. H.-C. Lai is the inventor of the patent.

Authors’ Contributions
Conception and design: M.W. Y. Chan, K.P. Nephew, T. Huang, H.-C. Lai
Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): Y.-T. Yo, C. Balch, R.-L. Huang, C.-K. Chen, K.P. Nephew, M.-H. Yu, H.-C. Lai
Writing, review, and/or revision of the manuscript: Y.-T. Yo, C. Balch, M.W. Y. Chan, K.P. Nephew, H.-C. Lai
Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): Y.-W. Lin, Y.-C. Wang, R.-L. Huang, C.-K. Chen, H.-C. Lai
Administrative, technical, or material support (i.e., reporting or organizing data, constructing databases): Y.-W. Lin, C. Balch, H.-K. Sytwu, C.-C. Chang
Study supervision: M.-H. Yu, H.-C. Lai

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