Cholestane-3β, 5α, 6β-triol Suppresses Proliferation, Migration, and Invasion of Human Prostate Cancer Cells

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Abstract
Oxysterols are oxidation products of cholesterol. Cholestane-3β, 5α, 6β-triol (abbreviated as triol) is one of the most abundant and active oxysterols. Here, we report that triol exhibits anti-cancer activity against human prostate cancer cells. Treatment of cells with triol dose-dependently suppressed proliferation of LNCaP CDXR-3, DU-145, and PC-3 human prostate cancer cells and reduced colony formation in soft agar. Oral administration of triol at 20 mg/kg daily for three weeks significantly retarded the growth of PC-3 xenografts in nude mice. Flow cytometric analysis revealed that triol treatment at 10–40 μM caused G1 cell cycle arrest while the TUNEL assay indicated that triol treatment at 20–40 μM induced apoptosis in all three cell lines. Micro-Western Arrays and traditional Western blotting methods indicated that triol treatment resulted in reduced expression of Akt1, phospho-Akt Ser473, phospho-Akt Thr308, PDK1, c-Myc, and Skp2 protein levels as well as accumulation of the cell cycle inhibitor p27Kip. Triol treatment also resulted in reduced Akt1 protein expression in PC-3 xenografts. Overexpression of Skp2 in PC-3 cells partially rescued the growth inhibition caused by triol. Triol treatment suppressed migration and invasion of DU-145, PC-3, and CDXR-3 cells. The expression levels of proteins associated with epithelial-mesenchymal transition as well as focal adhesion kinase were affected by triol treatment in these cells. Triol treatment caused increased expression of E-cadherin protein levels but decreased expression of N-cadherin, vimentin, Slug, FAK, phospho-FAK Ser722, and phospho-FAK Tyr861 protein levels. Confocal laser microscopy revealed redistribution of β-actin and α-tubulin at the periphery of the CDXR-3 and DU-145 cells. Our observations suggest that triol may represent a promising therapeutic agent for advanced metastatic prostate cancer.

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Introduction
Prostate cancer is the second most frequently diagnosed cancer of men and the fifth most common cancer overall in the world. In 2008, more than 899,000 new cases were diagnosed (GLOBOCAN 2008 database, version 1.2). In Western countries, prostate cancer is the most common non-cutaneous carcinoma of men. According to the statistics of Surveillance Epidemiology and End Results (SEER) of the National Cancer Institute, more than 240,000 men were diagnosed with and more than 28,000 men died of prostate cancer in 2012 in the United States. Although surgery is often successful for organ-confined prostate cancer, androgen ablation therapy is the primary treatment for metastatic prostate cancer. Unfortunately, most prostate cancer patients receiving androgen ablation therapy will ultimately develop recurrent, castration-resistant tumors within 1–3 years after treatment. The median overall survival time is 1–2 years after cancer relapse [1,2]. No effective standard therapy exists for patients that relapse with advanced prostate cancer. Chemotherapy is often used to treat metastatic hormone-refractory prostate cancer [2,3]. However, chemotherapies generally show little effect on prolonging survival. Therefore, new treatments for advanced prostate cancers are needed.
Oxysterols are oxidation products of cholesterol. Oxysterols play essential roles in regulating cholesterol homeostasis, platelet aggregation, apoptosis, and protein prenylation [4]. However, oxysterols are associated with development of atherosclerosis, neurological disease, and cancers [4]. Certain oxysterols have been reported to exhibit anticancer effects, possibly via modulation of cholesterol efflux, Akt, or liver X receptors (LXRs) [5,6]. For example, treatment with 22(R)-hydroxycholesterol, 24(S)-hydroxycholesterol, 7α-hydroxycholesterol, 7β-hydroxycholesterol, 25-hydroxycholesterol, and 5α,6α-epoxycholesterol suppressed the proliferation of human prostate, breast, colon, lung, and leukemia cancer cells [7–14]. These oxysterols caused either G1 cell cycle arrest [7–11] or apoptosis in cancer cells [12–14]. Therefore, oxysterols with cytotoxic activity might be a potential therapeutic agent for advanced prostate cancers.

Cholestane-3β, 5α, 6β-triol (abbreviated as triol) is one of the most abundant oxysterols. Triol is derived from cholesterol by oxidation via formation of 5α, 6α-epoxycholesterol and 5β, 6β-epoxycholesterol [15,16] as intermediates. Previously, 5α, 6α-epoxycholesterol was reported to exhibit anti-cancer activity [13]. In this study, we examined the ability of triol to suppress the proliferation of advanced human prostate cancer cell lines both in vitro and in vivo. We applied Micro-Western Arrays, a recently developed antibody-based, high-throughput Western blotting assay [17–20], to study the signaling proteins affected by triol in advanced prostate cancer cells. Our observations suggested that triol may represent a promising therapeutic agent for advanced metastatic prostate cancer.

**Materials and Methods**

**Materials**

Cholestane-3β, 5α, 6β-triol was purchased from Sigma (St. Louis, MO, USA). Matrigel was purchased from BD Bioscience (San Jose, CA, USA).

**Cell Culture**

Prostate cancer cell lines PC-3, DU-145, and LNCaP sublines were a gift from the laboratory of Professor Shutung Liao (The University of Chicago, IL, USA). Matrigel was purchased from BD Bioscience (San Jose, CA, USA).
medium) as previously described [28]. The PZ-HPV-7 cells were purchased from Bioresource Collection and Research Center (BCRC, Hsinchu City, Taiwan) and were cultured in a keratinocyte-serum free medium (Gibco/Invitrogen) with 5 ng/ml human recombinant epidermal growth factor and 50 μg/ml bovine pituitary extract.

**Cell Proliferation Assay**

Cells were seeded at a density of $3 \times 10^3$ cells/well in 100 μl complete medium in 96-well plates. Proliferation assays were performed under maintenance conditions (DMEM with 10% FBS for PC-3 and DU-145; DMEM with 10% CS-FBS for LNCaP CDXR-3). Relative cell number was analyzed by measuring the DNA content of cell lysates with the fluorescent dye Hoechst 33258 (Sigma, St. Louis, MO, USA) as described previously [8,10,11,21,23,24,27,28]. All readouts were normalized to the average of the control condition (no triol treatment) in each individual experiment. The experiment was repeated three times. Ten wells were utilized for each condition. The mean and standard deviation represented the results from all 30 wells in the three experiments.

**Cell Viability Assay**

Cells were seeded at a density of $3 \times 10^3$ cells per well in a 96-well plate (BD Bioscience). After 24 hrs, the cells were treated with increasing concentrations of triol for 48 hrs or 96 hrs. Cell viability was assessed by an MTT (3,4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) assay [29]. The amount of formazan was determined by measuring the absorbance at 560 nm using a Tecan GENios™ plate reader (Tecan group Ltd, Mannedorf, Switzerland) [29]. All results were normalized to the average of the control condition (no triol treatment) in each individual experiment. The experiment was repeated three times. Each time ten wells were utilized for each condition. The mean and standard deviation represented the results from all 30 wells in the three experiments.

**Flow Cytometric Analysis**

Cells were seeded at a density of $5 \times 10^5$ cells in 10-cm dishes in 10 mL of media and cultured for 24 hrs before addition of triol. After 48 hrs of culture in the presence of various concentrations of triol, cells were removed with trypsin and fixed in 70% ethanol in phosphate buffered saline (PBS) overnight at $-20^\circ$C. Fixed cells were washed with PBS, treated with 0.1 mg/mL RNase A in PBS for 30 min, and then suspended in PBS containing 50 μg/mL.
propidium iodide. Cell cycle profiles and distributions were determined by flow cytometric analysis of cells using a BD Facsscan flow cytometer (BD Biosciences). The cell cycle distribution was analyzed using ModFit LT software (Verity Software House, Topsham, ME, USA) as described [10,23,24,26,27].

Soft Agar Colony Formation Assay

PC-3 and LNCaP CDXR-3 cells (8x10³) were suspended in 0.3% low melting agarose (Lonza, Allendale, NJ, USA) containing DMEM with 10% FBS or 10% charcoal-stripped FBS (CS-FBS), respectively, and then layered on top of 3 ml of solidified 0.5% low melting agarose containing DMEM medium with 10% FBS (PC-3) or 10% CS-FBS (CDXR-3) in 6 cm dishes. Cells were allowed to grow at 37°C with 5% CO2 for 14 days (PC-3) or 17 days (CDXR-3). The plates were stained with 0.005% crystal violet in 30% ethanol for 6 hrs to detect cell colonies.

TUNEL Assay

Cells were cultured on cover slips in 24 wells and were treated with 0, 20, and 40 μM triol for 48 or 96 hrs. Cells were rinsed twice with PBS and subjected to the TUNEL assay using ApoAlert DNA Fragmentation Assay Kit (catalog no. 630108 from Clontech, Mountain View, CA, USA) according to the manufacturer’s instruction. The TUNEL-stained cells were observed with Olympus confocal microscope at 200 x (FV300, Olympus, Tokyo, Japan).

Ethics Statement

This study was carried out in strict accordance with the recommendations in the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health (Taiwan). The protocol was approved by the National Health Research Institutes Institutional Animal Care and Use Committee (NHRI-
**Figure 4. Effect of triol on cell cycle distribution of prostate cancer cells.** LNCaP CDXR-3 (A) and DU-145 cells (B) were treated with 0 or 10 μM triol for 48 hrs. PC-3 cells (C) were treated with 0, 10, or 20 μM triol for 48 hrs. (D) DU-145 cells were treated with 0, 20, and 40 μM triol for 48 hrs. Cell cycle distribution was determined by flow cytometry. Asterisk (*, **, ***) represents statistically significant difference ($p < 0.05$, $p < 0.01$, and $p < 0.001$, respectively) between the treated group and the control group.

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**Figure 5. Triol induces apoptosis in prostate cancer cells.** LNCaP CDXR-3, DU-145, and PC-3 cells were treated with 0, 20, and 40 μM triol for 48 hrs. Cell morphology was determined by light microscopy. TUNEL assay was performed as described in Materials and Methods to determine the apoptotic cell population. Green fluorescent light indicated apoptotic cells stained with TUNEL assay. Images were viewed at 200× with Olympus confocal microscope.

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Mice were kept under controlled environmental conditions (22°C, 12 hrs alternate light-dark cycles, 50% humidity, food and water ad libitum). Anesthesia (Isoflurane 1%) was given to reduce pain in mice during inoculation of prostate cancer cells. Animal care was conducted in accordance with the standard ethical guidelines (National Institutes of Health’s “Guide for the care and use of Laboratory animals” and Clinical laboratory animal medicine (K. Hrapkiewicz, L. Medina, and D.D. Holmes, 1998, Iowa State University Press). The experiment was designed to minimize the number of nude mice used.

Tumor Xenografts in Athymic Mice

Experiments involving mice were approved by National Health Research Institutes Institutional Animal Care and Use Committee (NHRI-IACUC-101046-A). Male Balb/c nu/nu mice (NCI Frederick, MD), 6–8 weeks of age, were injected subcutaneously in both flanks with 5×10⁶ PC-3 cells suspended in 0.5 ml of Matrigel (BD Bioscience, Franklin Lakes, NJ, USA). Monitoring of tumor growth was started one week after tumor inoculation. Mice were separated into control and treatment groups. 5 mice carrying 10 tumors comprised the control group and 5 mice carrying 8 tumors comprised the treatment group. Three weeks after inoculation of PC-3 cells, mice in the treatment group were treated orally 5 days/week with 20 mg/kg triol. Triol was administered in a vehicle containing 1% Tween 20, 25% dimethylsulfoxide in PBS. Mice in the control group were gavaged with vehicle only. Triol treatment was started 21 days after inoculation of cells and continued for 21 days. Tumors were measured weekly using calipers and volume was calculated using the formula volume = length × width × height × 0.52 [7,21–23].

Figure 6. Micro-Western Array analysis of protein expression in CDXR-3, DU-145, and PC-3 cells treated with 0 or 20 μM triol. CDXR-3, DU-145, and PC-3 cells were treated with 0 or 20 μM of triol for 48 hrs. Cell lysates were collected according to Micro-Western Array (MWA) protocol [18] and MWAs were performed to measure the changes in abundance and modification of indicated proteins. The six samples printed in each well from left to right were CDXR-3 (control), CDXR-3 (20 μM triol), DU-145 (control), DU-145 (20 μM triol), PC-3 (control), PC-3 (20 μM triol) (Figure S2). The right half blot (well A7-H12) was the technical duplicate of the left half blot (well A1-H6). Antibody list is shown in Table S2. Red and green were artificial color for anti-rabbit and anti-mouse 2nd antibody, respectively. The blot is the same size as a standard 96-well microtiter plate. Relative protein abundance of each signaling protein in each cell line is listed in Table S3.
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Luciferase-reporter Assay

PC-3 cells were seeded at 2 × 10^5 cells/well in a 12-well plate in DMEM containing 10% FBS. 24 hrs after plating, PC-3 cells were transfected with pRL-TK-Renilla luciferase plasmid (normalization vector; 5 ng/well), pSG5RXRa and pSG5LXRa (400 ng/well), 4xDR4A56fos pGL3 [reporter gene vector; 500 ng/well] using the PolyJet™ in vitro DNA transfection reagent (SigmaGen Laboratories, Rockville, MD, USA). 24 hrs after transfection, cells were treated with increasing concentrations of triol or T0901317. After an additional 24 hrs, cells were lysed in 100 μl passive lysis buffer (Promega, Madison, WI, USA) and luciferase activity was measured using a Dual-Luciferase kit (Promega) in a 20/20luminometer Turner Biosystems.

Western Blotting Analysis

Cells samples were lysed in SDS lysis buffer (240 mM Tris-acetate, 1% SDS, 1% glycerol, 5 mM EDTA pH 8.0, 0.1 M diithiothreitol) containing the protease inhibitors 1 mM 4-[2-aminomethyl]benzenesulfonyl fluoride (AEBSF), 0.8 μM aprotinin, 40 μM bestatin, 14 μM E-64, 20 μM leupeptin, 15 μM Pepstatin A (Sigma-Aldrich, P8340) and the phosphatase inhibitors cantharidin, bromotetramisole and microcystin LR (Sigma-Aldrich, P2850). Tissues samples were prepared from tissue homogenized in 2x Laemmli buffer as previously described [7,21–24]. Expression of proteins including Akt1, Skp2, phospho-Akt Ser473, phospho-Akt Thr308, PDK1, phospho-p44/42 MAPK Thr202/Tyr204, CDK2, CDK6, fatty acid synthase (FASN), GSK-3α, GSK-3β, Rb, cyclin D1, phospho-p38 MAPK Thr180/Tyr182, phospho-PDK1 Ser241, phospho-Rb Ser780, FAK, and MMP9 was detected using antibodies from Cell Signaling Technology (Danvers, MA, USA). E-cadherin, N-cadherin, and p27Kip1 antibodies were from BD BioSciences (San Jose, CA, USA). Antibodies for detection of c-Myc, caspase 3, caspase 8, caspase 9, PTEN were from Epitomics (Burlingame, CA, USA). Antibodies detecting Slug antibody was purchased from Abgent (San Diego, CA, USA). Vimentin and β-tubulin were from Millipore (Billerica, MA, USA). Antibodies for Skp2, p21, p27, MMP9, phospho-AKT (Thr308), phospho-AKT (Ser473), caspase 3, caspase 8, caspase 9, and β-tubulin were from Santa Cruz (Dallas, TX, USA). Horseradish peroxidase-conjugated secondary antibodies were from Santa Cruz (Dallas, TX, USA). The signal of horseradish peroxidase labeled secondary antibodies was detected by enhanced chemiluminescence reaction (ECL Western Blotting detection kit) (PerkinElmer, Waltham, MA, USA). GAPDH, β-tubulin, and β-actin were used as loading controls.

Micro-Western Arrays

CDXR-3 and DU-145 cells were treated with vehicle (DMSO) or 10 μM triol for 48 hrs. Micro-Western Arrays were performed to measure protein expression and modification as previously described [17,18,20,24].

Skrp2 Overexpression in PC-3 Cells

Ectopic expression of Skp2 was achieved by infecting PC-3 cells with a retrovirus generated from the LPCX plasmid containing wild-type human Skp2 cDNA as described [10]. Puromycin-resistant colonies were expanded and screened for increased Skp2

Figure 7. Protein expression profile of Micro-Western Array results. Protein expression profiles of CDXR-3, DU-145, and PC-3 cells treated with triol are displayed in a heat map format. Black color indicated no change, while green and red colors indicated decrease and increase of protein expression level. Grey color indicated signal below detection threshold. Protein abundance was normalized to the average of α-tubulin, GAPDH, and β-actin. Proteins in different cell lines were normalized to the control condition (no triol treatment) of that cell line. doi:10.1371/journal.pone.0065734.g007
Figure 8. Effect of triol treatment on the abundance and phosphorylation status of signaling proteins. Protein expression of Akt1, phospho-Akt Thr308, phospho-Akt Ser473, PDK1, phospho-p44/42 MAPK Thr202/Tyr204, c-Myc, Skp2, p27^kip1, FASN, GAPDH, β-actin, and α-tubulin in CDXR-3, DU-145, and PC-3 cells treated with 0, 10 or 20 μM triol for 48 hrs were assayed by Western blotting. Values represent relative protein abundance.

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Real-time quantitative PCR was performed using the Reverse Transcription SystemRevertAid™ H Minus First Strand cDNA Synthesis Kit (Fermentas/Thermo Scientific, cat. No. 74104, Qiagen, Venlo, The Netherlands) following the manufacturer’s instructions. The genomic DNA was removed by DNase-on-column treatment supplied with the mini kit. RNA concentration was determined spectrophotometrically at 260 nm. Equal amounts of RNA were used in the cDNA synthesis reactions using the Reverse Transcription SystemRevertAid™ H Minus First Strand cDNA Synthesis Kit (Fermentas/Thermo Scientific, Waltham, MA, USA). Real-time quantitative PCR was performed as previously described [7,8,21–23,28] using SYBR Green system/primers in an optical 96-well plate and cycling conditions consisting of 2 min at 50°C, 10 min at 95°C, 40 cycles of 15 sec at 95°C, and 60 sec at 60°C on an ABI Prism system (Applied Biosystems, Foster City, CA, USA). The sequences of primers for ATP-binding cassette transporter Al (ABCA1) were TTGTCAGTCAGTAATGCTTC (forward) and AAGCGATATGGTCCCGATT (reverse). The transcript level of ABCA1 was determined in PC-3 and DU-145 cells following treatment with 0, 10, and 20 μM triol for 48 hrs and was normalized to GAPDH levels in each sample.

Transwell Migration Assay
Migration assays with PC-3 cells were performed with a transwell kit from BD Bioscience (catalog number 353097). PC-3, DU-145, and CDXR-3 cells were treated with 0, 10, and 20 μM triol for 48 hrs. Cells were then removed from tissue culture plates with trypsin, and washed twice with PBS. Triol-treated cells (1×10^6) in 250 ul of DMEM without serum were placed in the upper invasion chamber and the lower compartment was loaded with DMEM containing 10% FBS. The cell migration chamber was inserted into the lower compartment and incubated for either 6 (PC-3, DU-145) or 24 (CDXR-3) hrs at 37°C. Cells on the topside of the filter were removed with a cotton swab. Cells attached to the filter were then fixed with methanol for 10 min. Cells attached to the filter were then stained with Giemsa stain (5%) for 1 hour. Filters were de-stained by washing with water and the number of cells attached to the filter was then quantified by enumerating cells in photographs of the stained filters.

Transwell Invasion Assay
An invasion assay with PC-3 cells was performed with Growth Factor Reduced BD BioCoat Matrigel invasion chambers (BD Biosciences) according to the manufacturer’s instructions. PC-3 cells were treated with different concentrations of triol for 48 hrs. Cells were then trypsinized and washed twice with PBS. Cells from each condition were seeded at 4×10^4 cells per well in serum-free DMEM in the upper compartment, and DMEM medium with 10% FBS was placed in the lower compartment of the chamber as a chemo-attractant. After 24 hrs of incubation, the non-invading cells on the upper side of the chamber were removed, and the membranes were fixed with methanol, washed, and stained with Giemsa’s solution. Invasiveness was evaluated by counting the invading cells under a light microscope. All experiments were conducted in triplicate.

Wound Healing Assay
CDXR-3 cells were pre-treated with 0, 10, and 20 μM triol for 24 hrs. Cells were then removed from tissue culture plates with trypsin, and washed twice with PBS. Cells were seeded at a concentration of 3.5×10^4 cells/well in 12-well plates. After 24 hrs, a wound healing assay was performed by scraping the cells with a 200-μl pipette tip. Cells were observed at 0, 8, and 24 hrs after scraping and photographed with a live imaging microscope (Leica AF 6000 LX, Leica, Wetzlar, Germany). The migration distance was automatically measured by the program inside live imaging microscope.

Confocal Microscopy
CDXR-3 and DU-145 cells were treated with 0 or 10 μM triol for 24 hrs. Cells were then washed 3 times with PBS buffer for 5 min per wash at room temperature (RT), fixed with 4% paraformaldehyde for 15 min at RT, and rinsed with PBS 3 times.

protein expression by Western blot analysis. PC-3 cells infected with an empty LPCX retrovirus were used as controls [10]. Cells were lysed in Laemmli buffer without bromphenol blue dye.

Real-Time Quantitative Polymerase Chain Reaction
RNA was extracted from the PC-3 and DU-145 cells treated with 0, 10, and 20 μM triol for 48 hrs using the RNeasy Minikit (cat. No. 74104, Qiagen, Venlo, The Netherlands) following the manufacturer’s instructions. The genomic DNA was removed by DNase-on-column treatment supplied with the mini kit. RNA concentration was determined spectrophotometrically at 260 nm. Equal amounts of RNA were used in the cDNA synthesis reactions using the Reverse Transcription SystemRevertAid™ H Minus First Strand cDNA Synthesis Kit (Fermentas/Thermo Scientific, Waltham, MA, USA). Real-time quantitative PCR was performed by Western blotting in PC-3 LPCX control and PC-3 LPCX Skp2 cell lines. PC-3 cells overexpressing Skp2 with LPCX vector and control PC-3 cells (empty LPCX vector) were treated with increasing concentrations of triol for 48 hrs (B) or 96 hrs (C) and analyzed by the 96-well proliferation assay. Asterisk * and *** represent statistically significant difference (p<0.05 and p<0.001) between the PC-3 control cells and PC-3 overexpressing Skp2- cells.

Figure 9. Effect of overexpression of Skp2 protein on growth inhibition by triol treatment. (A) Protein expression of Skp2 assayed by Western blotting in PC-3 LPCX control and PC-3 LPCX Skp2 cell lines. PC-3 cells overexpressing Skp2 with LPCX vector and control PC-3 cells (empty LPCX vector) were treated with increasing concentrations of triol for 48 hrs (B) or 96 hrs (C) and analyzed by the 96-well proliferation assay. Asterisk * and *** represent statistically significant difference (p<0.05 and p<0.001) between the PC-3 control cells and PC-3 overexpressing Skp2- cells.
for 5 min per wash and permeabilized for 10 minutes at RT with 0.1% Triton X-100 in PBS. Cells were blocked for non-specific protein binding with 2% bovine serum albumin in PBS for 30 min, rinsed with PBS three times for 5 min each. Cells were incubated with β-actin or α-tubulin antibody for one hour. After washing, cells were incubated with secondary FITC-conjugated antibody for one hour. After washing, cells were mounted under glass slides and sealed with Permount. Images of cells were observed with an Olympus confocal microscope at 400× (ocular lens 10×; objective lens 40×) (FV300, Olympus, Tokyo, Japan).

Data Analysis

Data are presented as the mean +/- SD of at least three experiments or are representative of experiments repeated at least three times. Student’s t test (two-tailed, unpaired) was used to evaluate the statistical significance of results from the proliferation assay experiments. A Microsoft Excel add-in program ED50V10 was used for calculating half maximal effective concentration (EC50).

Results

Triol Suppressed the Proliferaion of Human Prostate Cancer Cells

We first sought to determine the effect of triol treatment on the viability and proliferation of three commonly used human prostate cancer cell lines (Fig. 1). LNCaP CDXR-3 cells are androgen receptor (AR)-positive, relapsed, androgen-independent cells derived from parental AR-positive androgen-dependent LNCaP 104-S cells [27]. DU-145 and PC-3 both are AR-negative androgen-insensitive cells established from brain- and bone-derived metastasis, respectively. An MTT assay and a Hoechst dye-based proliferation assays indicated that triol suppressed both cell viability and cellular proliferation (Fig. 1). The inhibitory effect on proliferation was dose-dependent and increased over time (Fig. 1, Table S1). The EC50 for triol in the MTT assay was similar to the EC50 measured by Hoechst dye-based proliferation assay (Table S1), suggesting that inhibition of cell proliferation was responsible for the reduction of viable cells caused by triol treatment in all three human advanced prostate cancer cell lines.

Triol Treatment Inhibited Colony Formation of PC-3 and CDXR-3 Cells in Soft Agar

Treatment of PC-3 and LNCaP CDXR-3 cells with 25 μM and 50 μM triol markedly inhibited the formation of PC-3 and LNCaP CDXR-3 colonies in soft agar, confirming the anti-cancer activity of triol. DU-145 cells grew too slowly in soft agar to detect sufficient colonies for further analysis (Fig. 2).

Triol Treatment Retarded Growth of Androgen-insensitive Prostate Cancer Xenografts in Nude Mice

To determine if triol could suppress tumor growth in vivo, we performed a pilot study using DU-145 xenografts in nude mice. Intraperitoneal (i.p.) injection of 1 mg (50 mg/kg) of triol daily for 14 days caused a 36% reduction in the average volume of DU-145 xenografts growing in nude mice (Fig. S1). Triol treatment for 14 days did not affect the body weight of the mice (data not shown).

To determine if lower oral doses of triol could inhibit the growth of PC-3 prostate xenografts, we orally administered triol at 20 mg/kg five times a week. Oral administration of 20 mg/kg triol (5 times per week) from the 3rd to 6th week caused a 65% reduction in average tumor volume (p = 0.0002) in the treatment group compared to tumor volume in the vehicle control group (Fig. 3A). Body weight of both vehicle and triol-treated groups gradually decreased (Fig. 3B). This may be due to cachetic effects of PC-3 xenografts. These observations confirmed that oral administration of triol could retard the growth of prostate tumors in vivo.

Triol Treatment Caused G1 Cell Cycle Arrest in Prostate Cancer Cells

We next performed flow cytometric analysis to determine if cell cycle progression of prostate cancer cells was affected by triol. Treatment with 10 μM triol for 48 hrs caused an increase in the G1 phase cell population and a decrease in S and G2/M phase populations of LNCaP CDXR-3 and DU-145 prostate cancer cells.
Micro-Western Array Revealed Signaling Proteins being Affected by Triol Treatment

We hypothesized that triol may reduce cell viability through its effect on protein signaling networks. In order to determine what signaling proteins might be affected by triol treatment, we used Micro-Western Arrays (MWAs), a high-throughput Western blotting assay [17–20,24], to screen for signaling proteins affected by treatment with 10 μM triol in CDXR-3 and DU-145 cells. PTEN is frequently deleted in prostate cancer, resulting in activation of PI3K/Akt signaling [32]. PI3K/Akt signaling plays an important role in survival and progression of prostate cancer cells [32,33]. Up-regulation of PI3K/Akt activity is associated with poor clinical outcome of prostate cancer [34]. We used 48 antibodies that are capable of detecting proteins regulating cell cycle progression, cell survival and apoptosis, Akt-related signaling pathways, and several epithelial-mesenchymal transition (EMT) markers (Figure S2, Table S2) for the screening portion of our MWA study (Fig. 6). Differences in the protein expression profile in the absence and presence of 20 μM triol is shown in Fig. 7. Each of the three cell lines had a unique protein expression response to triol treatment. The protein profiles of CDXR-3 and PC-3 cells following treatment with 20 μM triol were more similar to each other than to DU-145 cells. Triol caused reduction of Akt1, cyclin E2, cyclin B1, phospho-c-Myc Thr38/Ser62, c-Myc, and phospho-Akt Ser473 in all three prostate cancer cell lines. Triol treatment at 20 μM resulted in at least a 20% increase in the protein abundance of phospho-p38 MAPK Thr180/Tyr182 (CDXR-3), DCK4 (DU-145), cyclin E1 (DU-145), cyclin D1 (DU-145), and p53 (PC-3); but caused at least a 20% decrease in the protein abundance of cyclin D1 (CDXR-3), phospho-c-Myc Thr58/Ser62 (CDXR-3), phospho-Rb Ser780 (CDXR-3), c-Myc (CDXR-3, PC-3), phospho-Akt Thr308 (CDXR-3, PC-3), phospho-Akt Ser473 (DU-145, PC-3), cyclin A (DU-145), cyclin B1 (DU-145, PC-3), phospho-GSK3β Ser21 (DU-145), NF-kB p65 (PC-3), and fatty acid synthase (FASN) (PC-3) (Table S3).

Triol Decreased the Level or Phosphorylation Status of Signaling Proteins Involved in Cell Cycle and Akt Signaling

The MWA result suggested that triol treatment altered the abundance and phosphorylation status of signaling proteins involved in cell cycle and Akt signaling in prostate cancer cells. We performed conventional Western blotting analysis to confirm these results. Triol treatment caused a decrease of Akt1 (DU-145, CDXR-3, phospho-Akt Thr308 (DU-145), phospho-Akt Ser473 (DU-145, CDXR-3, PC-3), PDK1 (DU-145, CDXR-3, PC-3), phospho-p44/42 MAPK Thr202/Tyr204 (DU-145, CDXR-3, PC-3), c-Myc (DU-145, CDXR-3), and S-phase kinase-associated protein 2 (Skp2) (DU-145, CDXR-3, PC-3). Triol treatment also resulted in elevated cell cycle inhibitor p27kip in all three cell lines. FASN expression was slightly increased in DU-145 but slightly decreased in CDXR-3 and PC-3 cells treated with triol (Fig. 8).
Overexpression of Skp2 Partially Rescued the Inhibitory Effect of Triol Treatment in PC-3 Cells

Since triol treatment caused a significant decrease in Skp2 expression and an increase in p27^Kip expression in all three prostate cancer cell lines (Fig. 8), we next wished to determine if overexpression of Skp2 would block the growth inhibition caused by triol in PC-3 cells. Indeed, overexpression of Skp2 partially rescued the cell proliferation rate of PC-3 cells under triol treatment (Fig. 9). Because overexpression of Skp2 did not

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**Figure 12. Effect of triol treatment on cytoskeleton morphology in CDXR-3 and DU-145 cells.** Imaging of α-tubulin and β-actin in CDXR-3 and DU-145 cells treated with vehicle or 10 μM triol for 48 hrs was determined by confocal microscopy.
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**Figure 13. Effect of triol treatment on migration and invasion of PC-3, DU-145, and CDXR-3 cells.** PC-3, DU-145, and CDXR-3 cells were pre-treated with 0, 10, or 20 μM triol for 48 hrs. Cells were then counted and seeded at identical number. Cancer cell migration ability (A) and invasion ability (B) was determined by transwell migration assay (A) and transwell invasion assay 6 hrs (A) and 24 hrs (B) after cell seeding, respectively. (C) Migration of CDXR-3 cells was determined by transwell migration assay 24 hrs after cell seeding. There was no triol in medium during the migration or invasion assay. Asterisk *** represents statistically significant difference (p<0.001) between the control group and triol treatment group.
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completely reverse the growth inhibitory effects, we hypothesized that other additional pathways or mechanisms may be influenced by triol to cause cell growth inhibition of prostate cancer cells.

**Triol Treatment Suppressed Akt1 in PC-3 Xenograft**

We examined the protein expression of Akt1, c-Myc, FASN, Ki-67, and α-tubulin in 5 tumors xenografts from the vehicle-treated group and 8 tumor xenografts from the triol-treated group from Fig. 3. Triol treatment resulted in significantly reduced Akt1 expression in PC-3 xenografts (p < 0.001, Fig. 10). Tumors in the treatment group tended to express lower c-Myc and FASN levels, although the p value was larger than 0.05. We also determined the protein levels of E-cadherin, N-cadherin, MMP-9, vimentin, slug, snail, p27Kip, and phospho-Akt Ser473 in these xenografts. The variation between individual tumors within both control and treatment groups of these proteins was large and no significant difference in specific protein abundances between control and treatment groups (data not shown).

**Triol is as a Weak Liver X Receptor (LXR) Agonist**

LXRs are ligand-activated transcriptional factors that belong to the nuclear receptor super family. LXRs are important regulators of cholesterol, fatty acid, and glucose homeostasis [9]. Two LXR isoforms exist: LXRα expression is most abundant in liver, kidney, intestine, fat tissue, macrophages, lung, and spleen; LXRβ is expressed ubiquitously [9]. A specific group of oxysterols, including 22(R)-hydroxycholesterol, 20(S)-hydroxycholesterol, 24(S)-hydroxycholesterol, and cholestenoic acid are natural ligands for LXRs [9,10]. We previously reported that LXR agonists, such as T0901317, 22(R)-hydroxycholesterol, and 24(S)-hydroxycholesterol suppress the proliferation of prostate and breast cancer cells via induction of G1 cell cycle arrest [6–10]. Since triol is an oxysterol, it might produce some effects as an agonist for LXR. To determine if triol is an LXR agonist, we determined if triol could activate LXR and induce expression of a luciferase reporter gene containing a LXR response element. Triol only slightly increased the transcriptional activity of LXRα (Fig. 11A). However, compared to the well known LXR agonist T0901317, triol was a very weak LXRα agonist (Fig. 11A). Treatment of DU-145 cells...
with 20 μM triol resulted in a slight increase in expression of ABCA1 mRNA expression, a target gene of LXRα activity (Fig. 11B), confirming the agonistic activity of triol for LXRα. The agonistic activity of triol for LXRα may partially contribute to the growth inhibition and G1 cell cycle arrest of prostate cancer cells.

Triol Treatment Altered Actin and Tubulin Distribution in Prostate Cancer Cells

Triol was previously reported to cause disruption of actin microfilaments [35], redistribution of vimentin [36], and loss of vinculin [35] in 73/73 endothelial cells. We next sought to determine if the cytoskeleton was affected by triol treatment of prostate cancer cells. Confocal laser microscopy revealed a modification of the β-actin and α-tubulin microfilaments, especially redistribution at the periphery of the cells following 48 hrs of treatment with 10 μM triol in CDXR-3 and DU-145 cells (Fig. 12). We did not observe significant changes in β-actin and α-tubulin microfilaments in PC-3 cells treated with 10 μM triol (data not shown).

Figure 15. Effect of triol on expression levels of EMT marker and FAK proteins in PC-3, DU-145, and CDXR-3 cells. Protein expression of E-cadherin, N-cadherin, vimentin, Slug, FAK, phospho-FAK Ser722, phospho-FAK Tyr861, β-actin, and α-tubulin in PC-3, DU-145, and CDXR-3 cells treated with 0, 10, or 20 μM triol for 48 hrs were assayed by Western blotting. Values represent relative protein abundance.

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suppression of proliferation of PZ-HPV-7 cells was 110.5 μM (Fig. 16). This suggested the possibility that non-cancerous human prostate cells may be more resistant to triol treatment compared to prostate cancer cells. However, the higher resistance of PZ-HPV-7 cells to triol treatment might be due to the lower proliferation rate of these non-cancerous cells compared to that of cancer cells. Further studies are needed to determine if triol will adversely affect normal prostate cells.

Discussion

Oxysterols are more polar and more readily diffusible through cell membranes than cholesterol, and possesses higher pro-inflammatory and cytotoxic effects than cholesterol in certain cells and tissues [13,37]. Oxysterols have been reported to be potentially involved in the initiation and progression of atherosclerosis, neurodegeneration, diabetes, kidney failure, and ethanol intoxication [13]. In cholesterol-rich foods, the most commonly found oxysterols include 7α-hydroxycholesterol, 7β-hydroxycholesterol, 7-ketocholesterol, 5α,6α-epoxycholesterol, and 5β,6β-epoxycholesterol [13]. Both 5α, 6α-epoxycholesterol and 5β, 6β-epoxycholesterol are unstable and are converted to cholestane-3β, 5α, 6β-triol [15,16]. Oxysterols have been shown to suppress the proliferation and survival of human cancer cells in vitro [7–14]. The cytotoxicity of oxysterols may cause growth inhibition and cell death in cancer cells if applied locally to tumors.

We observed that triol treatment inhibited tumor growth, reduced cell proliferation, induced G1 cell cycle arrest, and altered cytoskeletal morphology of human prostate cancer cells. Oral administration of triol at 20 mg/kg for three weeks reduced the average tumor volume of AR-negative androgen-insensitive PC-3 prostate cancer cells by 65% (Fig. 3). PC-3, one of the most commonly used prostate cancer cell lines, was established from bone-derived metastases [31]. Prostate cancer has a predilection to metastasize to bone and more than 80% of patients who died from prostate cancer developed bone metastases [38–40]. Our animal study suggested that triol treatment might be a potential treatment for prostate cancer metastasis. Indeed, we demonstrated that triol suppressed migration and invasion of DU-145, PC-3, and LNCaP CDXR-3 cells. The inhibition of migration and invasion of prostate cancer cells was not due to reduction of cell proliferation because the triol was given as a pre-treatment only and the same number of prostate cancer cells was seeded in both control and pre-treatment groups for the transwell assay (Fig. 9). Cancer cells in certain circumstance recapitulate normal developmental processes to enhance cell migration and invasion, a procedure called epithelial-to-mesenchymal transition (EMT). During the EMT, polarity is lost, motility is increased, mesenchymal markers are up-regulated, and differentiation markers are down-regulated [41,42]. This process was linked to cancer progression, in which epithelial cells acquire increased capacity for migration, stromal invasion, and metastasis [41,42]. During EMT of prostate cancer cells, the expression of transcription factors such as Snail, Slug, Twist, E47, ZEB1, and ZEB2 is increased, which leads to increased expression of N-cadherin and vimentin and concomitantly decreased expression of E-cadherin and cytokeratins [42–45]. Our study suggested that triol treatment resulted in decreased abundance of the proteins Slug, vimentin, and N-cadherin proteins while resulting in increased expression of the protein E-cadherin protein (Fig. 10). Focal adhesion kinase (FAK) is a non-receptor tyrosine kinase localized in focal contacts that becomes tyrosine phosphorylated. FAK signaling pathway is involved in the regulation of prostate cancer cell migration [46–49]. We observed that triol treatment resulted in reduced abundance of FAK in PC-3 cells as well as phospho-FAK Ser722 in DU-145 cells and phospho-FAK Tyr861 in PC-3 cells. This new finding indicated that FAK signaling is involved in the suppression of prostate cancer cell migration and invasion by triol treatment.
that triol treatment targeted both EMT markers and FAK signaling proteins and can thus be a potential therapeutic agent for treatment of prostate cancer metastasis. We observed that non-cancerous human prostate epithelial PZ-HPV-7 cells were more resistant to triol treatment compared to LNCaP, DU-145, and PC-3 prostate cancer cells (Fig. 11). This result suggested that triol has selectivity against prostate cancer cells versus normal cells.

Micro-Western Arrays (MWAs) represent a new method for proteomic analyses [19]. The MWA is a modification of the antibody-based modified reverse phase protein array. A non-contact microarray (such as the GenTARGET Nanoarray), a flatbed electrophoresis apparatus (such as the GE multiphor), and a near infrared scanner (such as the Licor Odyssey) comprise the MWA analysis system [18]. The MWA system allows detection of the abundance of modification status of up to 384 different proteins with as many as 15 samples simultaneously. The method has been shown to be effective in characterizing the EGFR pathway under multiple growth factor concentrations and in network modeling of complex receptor tyrosine kinase (RTK) cross-talk after stimulation [18]. The MWA has a technical advantage over other current technologies for assaying protein modifications such as Reverse Phase Lysate Arrays, quantitative Mass Spectroscopy, and Lumexin through its ability to efficiently and accurately measure hundreds of chosen phosphosites simultaneously, expanding the number of pathways surveyed in a single experiment. We applied this technology to screen signaling proteins being affected by triol in prostate cancer cells. In our MWA screening using 48 antibodies against proteins involved in Akt signaling and cell cycle regulation, we observed that triol caused reduction of Akt1, Akt2, cyclin E2, cyclin B1, phospho-c-Myc Thr58/Ser62, c-Myc, and phospho-Akt Ser473 in all three prostate cancer cell lines. Treatment with 20 μM triol caused at least a 20% increase in protein abundance of the following proteins: phospho-p38 MAPK Thr180/Tyr182, CDK4, cyclin E1, cyclin D1, and p53; while triol decreased in protein expression of cyclin D1, phospho-c-Myc Thr58/Ser62, phospho-Rb Ser780, c-Myc, phospho-Akt Thr308, phospho-Akt Ser473, cyclin A, cyclin B1, phospho-GSK3β Ser21, NF-kB p65, and FASN (Table S3). These findings were confirmed by conventional Western blotting. Western blotting experiments also showed that the protein levels of Akt1, phospho-Akt Ser473, phospho-Akt Thr308, PDK1, c-Myc, Skp2 in prostate cancer cells was decreased by triol treatment (Fig. 8). PI3K/Akt signaling plays an important role in survival and progression of prostate cancer cells [32,33,34,50]. Akt is a serine/threonine protein kinase encoded in mammals by three unique genes: Akt1, Akt2, and Akt3 [31,52]. Akt1 is involved in cellular survival pathways and inhibits apoptotic processes. Akt contains at least two important phosphorylation sites: threonine 308 and serine 473, which regulate its activity. Phosphorylation of threonine 308 on Akt is activated by PDK1 [53]. Phosphorylation of serine 473 is activated by mTOR kinase, its associated protein rictor, and SIN1/MIP1 [54,55]. Up-regulation of the PI3K/Akt activity is associated with poor clinical outcome of prostate cancer [33,34,50]. c-Myc protein, a well-known proto-oncoprotein, is an important transcriptional regulator of the androgenic response and cell proliferation in prostate and other cancers [25,36]. Expression of both c-Myc mRNA and protein increase during the progression of prostate cancer [57,58]. Decrease in c-Myc protein abundance results in decreased prostate cancer cell proliferation and tumor growth [23,25,27,59]. Forced expression of FASN promoted proliferation while siRNA knockdown of FASN induced apoptosis in prostate cancer cells [60]. Because triol treatment suppressed Akt1, phospho-Akt Ser473, phospho-Akt Thr308, PDK1, c-Myc, and FASN, reduction of Akt signaling, c-Myc, and FASN may all contribute to growth inhibition caused by triol. The only difference in results that we observed between MWAs and conventional Western blots was in the detection of phospho-p44/42 MAPK Thr202/Tyr204, possibly due to the low abundance of the Erk1/2 proteins in these prostate cancer cells and the relatively weak signal of the reporter antibody for this protein. Western blotting also indicated the induction of p27Kip1 under triol treatment. The p27Kip1 is a cell cycle inhibitory proteins and accumulation of p27Kip1 can result in cell cycle arrest and reduced cell proliferation of prostate cancer cells [23,26,61]. Skp2 is a member of the F-box protein family that is responsible for ubiquitination and down-regulation of p27Kip1 and other proteins [62,63]. Our observation that triol treatment resulted in decreased Skp2, increased p27Kip1, and G1 cell cycle arrest was consistent with the known functions of Skp2 and p27Kip1 (Fig. 4, Fig. 7, Fig. 8). Overexpression of Skp2 partially rescued the growth inhibition induced by triol (Fig. 9), indicating that other pathways or mechanisms, such as LXR signaling (Fig. 11) or cytoskeletal signaling (Fig. 12), were probably involved in the growth inhibition caused by triol in prostate cancer cells.

Our study suggested that triol treatment causes both G1 cell cycle arrest and apoptosis (Fig. 4, Fig. 5) in CDXR-3, DU-145, and PC-3 in vitro. Administration via gavage of 20 mg/kg/day triol five times a week retarded the growth of PC-3 xenografts (Fig. 3). The actual concentration of triol in serum will require further examination. However, the fact that Ki-67 was not significantly reduced in PC-3 tumors suggested the possibility that bioavailability of triol might be poor via oral administration. Local injection of triol might be a more efficient way to administer triol in future studies.

Triol was found to exert oxidative stress in rat hepatocytes as well as to increase levels of superoxide dismutase, catalase, and glutathione peroxidase activity [64]. Components of the cytoskeletal network, such as the actin microfilament meshwork or intermediate filaments, are known targets for oxidative and thiol-depleting agents that induce cell injury. Treatments with 20 μM triol, 7-ketocholesterol, or 25-hydroxycholesterol for 6 hrs caused progressive disruption of actin microfilaments, loss of vinculin, redistribution of vimentin, induction of cell detachment, and stimulation of apoptosis in 73/73 endothelial cells [35,36]. In our study, confocal laser microscopy revealed a modification of the β-actin and α-tubulin microfilaments, especially redistribution at the periphery of the cell following 48 hrs of treatment with 10 μM triol in CDXR-3 and DU-145 cells (Fig. 8). Alterations of actin and tubulin distribution have been reported to cause cell cycle arrest in human cancer cells [65,66]. Abnormal distribution of cytoskeletal components caused by triol may contribute to the induction of cell cycle arrest in human prostate cancer cells.

Different signaling proteins in different prostate cancer cells responded differently to triol treatment (Fig. 6–8, 15). Several proteins in DU-145 responded oppositely to 20 μM triol treatment compared to PC-3 and CDXR-3 cells (Fig. 7). This observation might be due to the different characteristics of PC-3, DU-145, and CDXR-3 cells. Although triol treatment suppressed cell proliferation, migration, invasion, and tumor growth in all three of these different prostate cancer cells, it may target different signaling proteins involved in the same pathways (such as Akt signaling) or similar functions (such as regulation of cell cycle, EMT, and cell structure) to produce similar results in different cell lines.

Conclusions
Our study provided systems-level insight into the molecular mechanism of triol action that resulted in suppression of proliferation, migration, and invasion of prostate cancer cells. Our results suggested that triol might be a potential therapeutic...
agent for treatment of advanced metastatic prostate cancer. In addition, our study highlights the power of the MWA approach for rapidly examining the systems level impact of small molecules on protein signaling networks and relating these changes to cell phenotypes.

Supporting Information

Figure S1 Effect of triol on the growth of DU-145 xenografts in nude mice. Male Balb/c nu/nu mice, 6-8 weeks of age, were injected subcutaneously in both flanks with 1x10^6 DU-145 cells suspended in 0.5 ml of Matrigel (BD Bioscience, Franklin Lakes, NJ, USA). Tumors were measured daily using calipers and volume was calculated using the formula volume = length x width x height x0.52. Tumor volume was allowed to increase to larger than 100 mm^3 before treatment started. Both control and treatment groups had 3 mice carrying 5 xenografts in nude mice. Male Balb/c nu/nu mice, 6–8 weeks of age, were injected subcutaneously in both flanks with 6x10^6 DU-145 cells suspended in 0.5 ml of Matrigel (BD Bioscience, Franklin Lakes, NJ, USA). Tumors were measured daily using calipers and volume was calculated using the formula volume = length x width x height x0.52. Tumor volume was allowed to increase to larger than 100 mm^3 before treatment started. Both control and treatment groups had 3 mice carrying 5 xenografts in nude mice.

Table S1 EC_{50} of triol to suppress viability and proliferation of prostate cancer cells. LC50P CDXR-3, DU-145, and PC-3 cells treated with triol for 48 hrs or 96 hrs were assayed with MTT assay or Hoechst dye-based proliferation assay to determine the EC_{50} of triol to cause growth inhibition.

Table S2 List of antibodies used in the MWA study. Information of antibodies and location of antibodies on the antibody plate for MWA blot incubation is shown. The α-tubulin, β-actin, and GAPDH proteins were detected as loading controls.

Table S3 Relative expression level of signaling proteins in prostate cancer cells assayed with MWA. Relative abundance of signaling proteins in CDXR-3, DU-145, and PC-3 cells treated with 0 or 20 µM triol for 48 hrs was determined by MWA and shown as list. N/A represents signaling being too weak to be detected. Protein abundance was normalized to the average of α-tubulin, GAPDH, and β-actin. Proteins in different cell lines were normalized to the control condition (no triol treatment) of that cell line.

Author Contributions

Conceived and designed the experiments: CPC. Performed the experiments: CYL CH LKK HPL YWH YLW LCS YVK JCT RAH JMK CPC. Analyzed the data: YF YWH JMK RBJ CCY HHF YHK LC SDY WYC KKT JYC. Contributed reagents/materials/analysis tools: YF RAH JMK CPC. Wrote the paper: CPC RAH JMK RBJ WYC LC.

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