

Axitinib Targeted Cancer Stemlike Cells to Enhance Efficacy of Chemotherapeutic Drugs via Inhibiting the Drug Transport Function of ABCG2

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Stemlike cells have been isolated by their ability to efflux Hoechst 33342 dye and are called the side population (SP). We evaluated the effect of axitinib on targeting cancer stemlike cells and enhancing the efficacy of chemotherapeutic agents. We found that axitinib enhanced the cytotoxicity of topotecan and mitoxantrone in SP cells sorted from human lung cancer A549 cells and increased cell apoptosis induced by chemotherapeutic agents. Moreover, axitinib particularly inhibited the function of adenosine triphosphate (ATP)-binding cassette subfamily G member 2 (ABCG2) and reversed ABCG2-mediated multidrug resistance (MDR) *in vitro*. However, no significant reversal effect was observed in ABCB1-, ABCC1- or lung resistance-related protein (LRP)-mediated MDR. Furthermore, in both sensitive and MDR cancer cells axitinib neither altered the expression of ABCG2 at the mRNA or protein levels nor blocked the phosphorylation of AKT and extracellular signal-regulated kinase (ERK)1/2. In nude mice bearing ABCG2-overexpressing S1-M1-80 xenografts, axitinib significantly enhanced the antitumor activity of topotecan without causing additional toxicity. Taken together, these data suggest that axitinib particularly targets cancer stemlike cells and reverses ABCG2-mediated drug resistance by inhibiting the transporter activity of ABCG2.

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INTRODUCTION

Axitinib is an oral, potent, small-molecule adenosine triphosphate (ATP)-competitive multitargeted tyrosine kinase inhibitor (TKI). It inhibits cellular signaling by blocking vascular endothelial growth factor receptor 1 (VEGFR-1), VEGFR-2 and VEGFR-3; platelet derived growth factor receptor (PDGFR); and c-KIT (CD117) (1–3). These receptor TKs are transmembrane proteins at the cell surface that play critical roles in the transduction of extracellular signals to

the cytoplasm. It has been reported that these receptors are important in signaling pathways and the development of a number of tumors (4,5). Inhibition of these TKs blocks signal transduction pathways that affect many of the processes involved in tumor cell proliferation, progression, metastasis and angiogenesis. In preclinical and clinical studies, axitinib has been shown to inhibit angiogenesis, vascular permeability and blood flow. In phase II studies, axitinib showed single-agent activity in a variety

of tumor types, including non-small cell lung cancer (6), advanced renal cell carcinoma (7) and thyroid cancer (8).

ATP-binding cassette (ABC) drug transporter proteins can use the energy derived from ATP hydrolysis to extrude numerous structurally and mechanistically unrelated anticancer drugs, which play a key role in the development of multidrug resistance (MDR). Overexpression of ABC transporters is a significant hindrance to successful cancer chemotherapy (9). There are 49 ABC transporter family genes in the human genome, which are divided into seven subfamilies on the basis of amino acid sequence similarities and phylogeny (10). Among them, the ABC transporter-subfamily B member 1 (ABCB1/MDR1/P-glycoprotein, P-gp), subfamily C member 1 (ABCC1/ MRP1) and subfamily G member 2 (ABCG2/BCRP) are considered to be the most important transporters to confer MDR to tumor cells.

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ABCG2/BCRP, also called MXR (mitoxantrone resistance associated protein) and ABCP (placenta-specific ABC transporter), was identified independently from drug-selected human breast cancer cells (MCF-7/AdrVp) (11), human colon carcinoma cells (S1-M1-80) (12) and human placenta (13). The human *ABCG2* gene is located on chromosome 4, band 4q21-4q22 and encodes a 72.6-kD membrane protein composed of 655 amino acids (14). *ABCG2* can transport a wide range of anticancer agents such as doxorubicin (Dox), topotecan, SN-38, mitoxantrone and methotrexate as well as fluorescent dyes such as Hoechst 33342 (15). Wild-type *ABCG2*, with an arginine at position 482 (R482), facilitated efficient transport of mitoxantrone, but not rhodamine 123 or Dox. MCF7/AdVp3000 and S1-M1-80 cells expressing R482T and R482G variants of BCRP/*ABCG2*, respectively, transported rhodamine 123 and Dox while also maintaining their ability to transport mitoxantrone (16,17). Therefore, certain kinds of single-nucleotide polymorphisms of *ABCG2* can alter its function, and consequently affect the disposition of substrate drugs.

Malignant stemlike cells have been identified in various malignant tumors, ranging from leukemia to solid tumors. Like normal stem cells, these cancer stemlike cells (CSCs) are able to self-renew, differentiate and proliferate extensively. The cancer mass originates from rare stemlike cells that can transfer the disease to immunodeficient mice. This finding suggests that these CSCs are responsible for the relapse of cancer following conventional or targeted cancer therapy and that eradication of these CSCs might be necessary to cure the disease permanently. However, it appears likely that CSCs are not effectively ablated by most current therapeutic strategies, leaving the potential for disease progression or relapse. Several recent studies have provided insight into the signaling pathways underlying the CSC phenotype and have also suggested approaches to eliminate CSCs (18,19). The side population (SP) phenotype cells, believed to be CSCs, are

present in diverse tumor types and overexpress *ABCG2*, producing inherent drug resistance (20,21). Currently, *ABCG2* is considered to be a molecular marker for the SP cells (22).

ABCG2 is an ideal target for development of chemosensitizing agents for better treatment of drug-resistant cancers. However, very few compounds have been identified as specific inhibitors of *ABCG2*. Fumitremorgin C (FTC), a mycotoxin from *Aspergillus fumigatus*, was reported first (23). However, FTC neurotoxicity prevented any clinical use. Although analogues of FTC, such as Ko132 and Ko143, have been developed with low toxicity, the efficacy and safety of these modulators are still under evaluation in *in vivo* studies (24). Another less well characterized, but promising, strategy is the application of TKIs, small molecule hydrophobic compounds, which are designed to arrest aberrant signaling pathways in malignant cells. It has been recently shown that the TKIs interact with and modulate the function of ABC transporters such as *ABCB1*, *ABCC1* and *ABCG2*. The BCR-Abl TKIs imatinib (Gleevec) and nilotinib (Tasigna) interact with *ABCB1* and *ABCG2* transporters and significantly inhibit their transport activity (25,26). Gefitinib, an epidermal growth factor receptor (EGFR) TKI, has been observed to directly inhibit the function of *ABCB1* in MDR cancer cells (27) and reverse *ABCG2*-mediated MDR *in vitro* (28). In animal models, gefitinib affected the oral absorption of chemotherapeutic agents by modulating the function of *ABCB1* and *ABCG2* (29). In our previous study, we also found that lapatinib and sunitinib antagonized *ABCB1*- and/or *ABCG2*-mediated MDR (30,31).

Although axitinib was effective as an oral agent in earlier stages of clinical development, its interaction with ABC drug transporters has not been characterized. The objective of this work was to investigate the interaction of axitinib with *ABCB1*, *ABCC1*, *ABCC4*, *ABCG2* and lung resistance-related protein (LRP). We show here that axitinib targets CSCs, in-

creases the efficacy of chemotherapeutic agents and reverses *ABCG2*-mediated drug resistance by inhibiting the drug efflux function of *ABCG2* and increasing the intracellular accumulation of cytotoxic agents in *ABCG2*-overexpressing cells and SP cells.

MATERIALS AND METHODS

Chemicals and Reagents

3-(4,5-Dimethylthiazol-yl)-2,5-diphenylaplatinibrazolium bromide (MTT), vincristine, topotecan, mitoxantrone, Dox, rhodamine 123, verapamil, MK571 and FTC were products of Sigma Chemical Co (St. Louis, MO, USA). Axitinib was purchased from Selleck Chemicals (Houston, TX, USA). Dulbecco's modified Eagle's medium (DMEM) and RPMI 1640 were products of Gibco BRL (Gaithersburg, MD, USA). Anti-mitogen-activated protein kinase (anti-MAP) 1/2 (extracellular signal-regulated kinase [ERK]1/2) (KC-5E01), P-ERK (KC-5E04), P-AKT (KC-5A04) and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) antibodies were purchased from Kangchen Co. (Shanghai, China). AKT antibody (#4685) was a product of Cell Signaling Technology Inc. (Danvers, MA, USA). *ABCG2* antibody (MAB4146) was obtained from Chemicon International, Inc. (Temecula, CA, USA). Other routine laboratory reagents were of analytical grade and obtained from commercial sources.

Cell Culture

The following cell lines were cultured in DMEM or RPMI 1640 containing 10% fetal bovine serum at 37°C in the presence of 5% CO₂: the human colon carcinoma cell line S1 and its mitoxantrone-selected derivative *ABCG2*-overexpressing S1-M1-80 cell line were kindly provided by SE Bates (National Cancer Institute, National Institutes of Health [NIH], Bethesda, MD, USA) (32); the human oral epidermoid carcinoma cell line KB and its vincristine-selected derivative *ABCB1*-overexpressing cell line KBv200 were a gift from Xu-Yi Liu (Cancer Hospital of Beijing, Beijing,

China) (33); the human leukemia cell line HL60 and its Dox-selected derivative *ABCC1*-overexpressing cell line HL60/ADR were purchased from the Institute of Hematology & Blood Diseases Hospital, Chinese Academy of Medical Sciences & Peking Union Medical College, Tianjin, China; the human lung squamous carcinoma cell line SW1573 and its Dox-selected derivative LRP-overexpressing SW1573/2R120 were provided by EAC Wiemer (Department of Medical Oncology, Erasmus Medical Center, Rotterdam, the Netherlands) (34); the murine fibroblasts cell line NIH3T3 and the *ABCC4*-transfected *ABCC4* stable expressing NIH3T3/MRP4-2 cells were kindly provided by Z-S Chen (St John's University, Queens, NY, USA) (35). The human lung cancer A549 cell line was purchased from ATCC (American Type Culture Collection, Manassas, VA, USA). HEK293/pcDNA3.1, *ABCG2*-482-G2 and *ABCG2*-482-T7 cells were established by selection with G418 after transfecting HEK293 with either an empty pcDNA3.1 vector or a pcDNA3.1 vector containing full-length *ABCG2* coding either glycine (G) or threonine (T) at the amino acid 482 position, respectively (36). These cells were obtained from SE Bates (National Cancer Institute, NIH) and were cultured in medium with 2 mg/mL G418.

Cell Cytotoxicity Test

The MTT assay was used to assess the sensitivity of cells to drugs as previously described (37). Briefly, cells were distributed evenly into 96-well microtiter plates and then various concentrations of axitinib were added to the wells. After 68 h of incubation, MTT (5 mg/mL, 20 μ L/well) was added into the cells for 4 h (37°C). Afterward, the medium was discarded, and 200 μ L of dimethylsulfoxide was added to dissolve the formazan product from the metabolism of MTT. Optical density was measured at 540 nm with background subtraction at 670 nm by use of the Model 550 Microplate Reader (BIO-RAD, Hercules, CA, USA). The concentration required to inhibit cell

growth by 50% (IC_{50}) was calculated from survival curves by use of the Bliss method (38). For reversal experiments, axitinib was added to the medium with full range concentrations of (a) topotecan, mitoxantrone and cisplatin in S1 and S1-M1-80; (b) Dox and cisplatin in KB and KBv200; (c) Dox and cisplatin in HL60 and HL60/ADR; (d) Dox and cisplatin in SW1573 and SW1573/2R120; and (e) 6-mercaptopurine (6-MP) and cisplatin in NIH3T3 and NIH3T3/MRP4-2 cells. Fold of resistance was calculated by dividing the IC_{50} for the MDR cells by that for the parental sensitive cells. The degree of reversal of MDR (fold reversal) was calculated by dividing the IC_{50} for cells with the anticancer drug in the absence of axitinib by that obtained in the presence of axitinib.

Animals

Athymic nude mice (BALB/c-nu/nu) of both sexes, 5 to 6 wks old and weighing 18 to 22 g, were bred at the Center of Experimental Animals, Sun Yat-Sen University (Guangzhou, China), and were used for the S1 and S1-M1-80 cell xenografts. Male nonobese diabetic/severe combined immunodeficiency (NOD/SCID) mice, 4–5 wks old, were purchased from Beijing HFK Biotechnology Co. Ltd (Changping District, Beijing, PR China) and were used for the tumorigenicity experiments. All animals received sterilized food and water. All experiments were conducted with the approval of the Sun Yat-Sen University Institutional Animal Care and Use Committee.

Tumor Xenograft Experiments

The S1-M1-80 cell xenograft model was established as previously described (39) with slight modification. Briefly, 1×10^7 S1-M1-80 cells were injected subcutaneously into the posterior flank region of the nude mice. The mice were randomized into four groups after the tumors reached a mean volume of about 100 mm³, and then received various treatments: (a) saline (every 4 d \times 9, intraperitoneally [IP]); (b) topotecan

(every 4 d \times 9, IP, 3 mg/kg); (c) axitinib (every 4 d \times 9, orally [PO], 25 mg/kg); (d) topotecan (every 4 d \times 9, IP, 3 mg/kg) plus axitinib (every 4 d \times 9, PO, 25 mg/kg) (axitinib was given 1 h before topotecan administration). The whole administration was divided into three cycles with a 10-d drug-free recovery period between every two cycles.

For the S1 cell xenograft model, 1×10^7 S1 cells were injected subcutaneously into the posterior flank region of the nude mice. The mice were randomized into four groups after the tumors reached a mean diameter of 0.5 cm, and then received various treatments: (a) saline (every 4 d \times 4, IP); (b) topotecan (every 4 d \times 4, IP, 3 mg/kg); (c) axitinib (every 4 d \times 4, PO, 25 mg/kg); (d) topotecan (every 4 d \times 4, IP, 3 mg/kg) plus axitinib (every 4 d \times 4, PO, 25 mg/kg) (axitinib was given 1 h before topotecan administration). Tumor volumes (*V*) were calculated by the following formula (38):

$$V = \frac{\pi}{6} \left(\frac{A+B}{2} \right)^3$$

In the formula, A is the longer diameter and B is the diameter perpendicular to A. The mouse weight, tumor volume, feeding behavior and activity were recorded every 4 d. Mice were killed when the mean of tumor weights was over 1 g in the control group, and tumor tissue was excised from the mice and weighed. The ratio of growth inhibition (IR) was estimated according to the following formula:

$$IR (\%) = \frac{1 - \text{Mean tumor weight of experimental group}}{\text{Mean tumor weight of control group}} \times 100.$$

SP Analysis and Sorting

We labeled the cell suspensions with Hoechst 33342 dye using the methods described by Goodell *et al.* (40) with modifications. Briefly, A549 cells were re-suspended at 1×10^6 /mL in prewarmed DMEM with 2% fetal calf serum and 10 mmol/L HEPES (4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid) buffer. Hoechst 33342 dye was added at a final concentration of 5 μ g/mL in the presence or absence of FTC (10 μ mol/L), and the

Table 1. Effect of axitinib on reversing ABCG2-mediated drug resistance.^a

Compounds	IC ₅₀ ± SD, μmol/L (Fold reversal)			
	S1		S1-M1-80(ABCG2)	
Topotecan	0.250 ± 0.014	(1.00)	12.79 ± 0.241	(1.00)
+0.25 μmol/L Axitinib	0.249 ± 0.020	(1.01)	6.692 ± 0.504 ^b	(1.85)
+0.5 μmol/L Axitinib	0.255 ± 0.017	(0.98)	4.964 ± 0.320 ^b	(2.47)
+1.0 μmol/L Axitinib	0.199 ± 0.006	(1.26)	3.020 ± 0.242 ^b	(4.11)
+0.5 μmol/L FTC	0.244 ± 0.013	(1.02)	1.195 ± 0.108 ^b	(10.7)
Mitoxantrone	0.193 ± 0.021	(1.00)	14.668 ± 0.636	(1.00)
+0.25 μmol/L Axitinib	0.194 ± 0.057	(0.99)	7.992 ± 0.675 ^b	(1.84)
+0.5 μmol/L Axitinib	0.201 ± 0.011	(0.96)	4.140 ± 0.301 ^b	(3.54)
+1.0 μmol/L Axitinib	0.185 ± 0.011	(1.12)	2.904 ± 0.482 ^b	(5.05)
+0.5 μmol/L FTC	0.188 ± 0.011	(1.03)	1.272 ± 0.056 ^b	(11.5)
Cisplatin	14.877 ± 1.126	(1.00)	14.496 ± 1.112	(1.00)
+1.0 μmol/L Axitinib	14.206 ± 1.460	(1.05)	14.360 ± 1.812	(1.01)
	KB		KBv200(ABCB1)	
Doxorubicin	0.024 ± 0.002	(1.00)	2.204 ± 0.108	(1.00)
+0.25 μmol/L Axitinib	0.024 ± 0.002	(1.00)	2.022 ± 0.185	(1.08)
+0.5 μmol/L Axitinib	0.025 ± 0.001	(0.96)	2.153 ± 0.146	(1.02)
+1.0 μmol/L Axitinib	0.015 ± 0.004 ^c	(1.63)	1.250 ± 0.215 ^c	(1.76)
+1.0 μmol/L Verapamil	0.024 ± 0.003	(1.00)	0.109 ± 0.008 ^b	(20.2)
Cisplatin	1.552 ± 0.121	(1.00)	2.364 ± 0.241	(1.00)
+1.0 μmol/L Axitinib	1.514 ± 0.117	(1.03)	2.412 ± 0.225	(0.98)
	HL60		HL60/ADR(ABCC1)	
Doxorubicin	0.055 ± 0.011	(1.00)	5.725 ± 0.203	(1.00)
+0.1 μmol/L Axitinib	0.054 ± 0.006	(1.02)	5.719 ± 0.591	(1.00)
+0.2 μmol/L Axitinib	0.056 ± 0.006	(0.98)	5.759 ± 0.659	(0.99)
+0.4 μmol/L Axitinib	0.049 ± 0.005	(1.13)	5.476 ± 0.384	(1.04)
+40 μmol/L MK571	0.054 ± 0.005	(1.02)	1.108 ± 0.182 ^b	(5.17)
Cisplatin	1.456 ± 0.187	(1.00)	1.658 ± 0.118	(1.00)
+0.4 μmol/L Axitinib	1.516 ± 0.125	(0.96)	1.524 ± 0.175	(1.08)
	SW1573		SW1573/2R120(LRP)	
Doxorubicin	0.108 ± 0.012	(1.00)	1.266 ± 0.121	(1.00)
+0.25 μmol/L Axitinib	0.096 ± 0.004	(1.12)	1.214 ± 0.174	(1.04)
+0.5 μmol/L Axitinib	0.098 ± 0.004	(1.10)	1.236 ± 0.122	(1.02)
+1.0 μmol/L Axitinib	0.094 ± 0.006	(1.14)	1.138 ± 0.088	(1.11)
Cisplatin	2.473 ± 0.167	(1.00)	9.659 ± 1.073	(1.00)
+1.0 μmol/L Axitinib	2.790 ± 0.147	(0.89)	9.180 ± 0.881	(1.05)
	NIH3T3		NIH3T3/MRP4-2(ABCC4)	
6-MP	0.298 ± 0.050	(1.00)	11.852 ± 1.564	(1.00)
+0.25 μmol/L Axitinib	0.285 ± 0.034	(1.04)	11.145 ± 1.048	(1.06)
+0.5 μmol/L Axitinib	0.318 ± 0.045	(0.94)	10.709 ± 1.637	(1.11)
+1.0 μmol/L Axitinib	0.265 ± 0.011	(1.12)	12.319 ± 1.704	(0.96)
Cisplatin	6.140 ± 0.728	(1.00)	5.668 ± 0.690	(1.00)
+1.0 μmol/L Axitinib	5.033 ± 0.296	(1.21)	5.842 ± 0.691	(0.97)

^aCell survival was determined by MTT assays as described in Materials and Methods. Data are the mean ± standard deviation (SD) of at least three independent experiments performed in triplicate. The fold reversal of MDR (values given in parentheses) was calculated by dividing the IC₅₀ for cells with the anticancer drugs in the absence of axitinib by that obtained in the presence of axitinib.

^bP < 0.01; ^cP < 0.05 versus the values obtained in the absence of inhibitor.

cells were incubated at 37°C for 90 min with intermittent shaking. At the end of the incubation, the cells were washed with ice-cold phosphate-buffered saline (PBS), centrifuged down at 4°C, and re-suspended in ice-cold PBS. Propidium iodide at a final concentration of 2 μg/mL was added to the cells to gate viable cells. The cells were filtered through a 40-μm cell strainer to obtain a single-cell suspension before sorting. Analyses and sorting were done with fluorescence-activated cell sorting (FACS). The Hoechst 33342 dye was excited at 357 nm and its fluorescence was dual-wavelength analyzed (blue, 402–446 nm; red, 650–670 nm).

Tumorigenicity Experiments

Sorted SP and non-SP cells from A549 cells were subcutaneously injected into the NOD/SCID mice. Groups of mice were inoculated with SP or non-SP cells at 1 × 10⁶, 1 × 10⁵, 1 × 10⁴ and 1 × 10³ (six mice per group). The mice were killed 44 d after tumor cell injection.

Detection of Cell Surface Expression of ABCG2 and ABCB1 by Flow Cytometer

SP cells were collected and washed three times with an isotonic PBS buffer (supplemented with 0.5% bovine serum albumin [BSA]). For ABCG2 expression analysis, APC-conjugated anti-human Bcrp1/ABCG2 (R&D Systems, Minneapolis, MN, USA) reagent were mixed with 25 μL of Fc-blocked cells (1 × 10⁶ cells). After incubating for 45 min at 4°C, the cells were washed twice with PBS buffer (supplemented with 0.5% BSA) and re-suspended in 400 μL PBS buffer for flow cytometric analysis. Isotype control samples were treated in an identical manner with allophycocyanin (APC)-labeled mouse immunoglobulin G2b (IgG2b) antibody. For ABCB1 flow cytometric analysis, 1 × 10⁶ cells (100 μL) were incubated at 4°C for 30 min with 10 μL of CD243-PE conjugated antibody (Beckman Coulter, Fullerton, CA, USA), cells were then washed and resuspended in PBS. Isotype control samples were treated with mouse

Table 2. Effect of axitinib on reversing ABCG2-mediated MDR in transfected cell lines.^a

Compounds	IC ₅₀ ± SD (μmol/L) (fold reversal)					
	HEK293/pcDNA3.1		ABCG2-G482-G2		ABCG2-482-T7	
Mitoxantrone	0.029 ± 0.0015	(1.00)	1.176 ± 0.241	(1.00)	1.195 ± 0.301	(1.00)
+0.25 μmol/L Axitinib	0.031 ± 0.0024	(0.94)	0.488 ± 0.085 ^b	(2.41)	0.338 ± 0.028 ^b	(3.54)
+0.5 μmol/L Axitinib	0.041 ± 0.0029	(0.71)	0.247 ± 0.076 ^b	(4.76)	0.206 ± 0.013 ^b	(5.80)
+1.0 μmol/L Axitinib	0.042 ± 0.0033	(0.70)	0.079 ± 0.007 ^b	(14.9)	0.076 ± 0.005 ^b	(15.7)
+0.5 μmol/L FTC	0.040 ± 0.0011	(0.73)	0.138 ± 0.005 ^b	(8.52)	0.096 ± 0.003 ^b	(12.4)
Cisplatin	4.069 ± 0.258	(1.00)	5.141 ± 0.197	(1.00)	4.324 ± 0.176 ^b	(1.00)
+1.0 μmol/L Axitinib	5.449 ± 0.269	(0.75)	4.512 ± 0.224	(1.14)	5.078 ± 0.209 ^b	(0.85)

^aCell survival was determined by MTT assays as described in Materials and Methods. Data are the mean ± standard deviation (SD) of at least three independent experiments performed in triplicate. The fold reversal of MDR (values given in parentheses) was calculated by dividing the IC₅₀ for cells with the anticancer drugs in the absence of axitinib by that obtained in the presence of axitinib.

^b*P* < 0.01 versus the values obtained in the absence of inhibitor.

IgG2a antibody in parallel. Tests and controls were analyzed with a flow cytometer.

Apoptosis Assay

Cells were seeded onto a six-well plate at a density of about 2.0×10^5 cells/well. After treatment with different concentrations of axitinib in the presence of 0.2 μmol/L topotecan or mitoxantrone for 48 h, both floating and attached cells were collected and washed with ice-cold PBS twice. Cells were resuspended in 100 μL of $1 \times$ binding buffer, and the Alexa Fluoro 488 annexin V (5 μL) and propidium iodide (PI) (1 μL) were added before incubation at room temperature for 15 min. After the incubation period, we added 400 μL $1 \times$ binding buffer, mixed gently and analyzed via FACS.

Doxorubicin and Rhodamine 123 Accumulation

The effect of axitinib on the intracellular accumulation of Dox and rhodamine 123 was performed as previously described (41). Briefly, the cells were treated with axitinib of various concentration or vehicle at 37°C for 3 h. Subsequently, 10 μmol/L Dox or 5 μmol/L rhodamine 123 was added and the incubation was continued for an additional 3 h or 0.5 h, respectively. The cells were then collected, centrifuged and washed three times with cold PBS. Finally, the cells were analyzed with flow cytometric anal-

ysis (Cytomics FC500, Beckman Coulter). FTC was used as a control inhibitor of ABCG2 in S1 and S1-M1-80 cells (42).

ATPase Assay of ABCG2

The vanadate-sensitive ATPase activity of ABCG2 in cell membrane prepared from High-Five insect cells (cat. no. 453270) was measured by using the BD Gentest ATPase assay kit (both cells and kit from BD Biosciences, San Jose, CA, USA) according to the manufacturer's instructions. Briefly, the ATPase reaction was initiated by the addition of 12 mmol/L Mg-ATP into a total reaction mixture of 60 μL. After an incubation at 37°C for 10 min, the reactions were terminated by the addition of 30 μL of 10% sodium dodecyl sulfate (SDS) solution. The liberation of inorganic phosphate was detected by its absorbance at 800 nm and quantified by comparing the absorbance to a phosphate standard curve.

Western Blot Analysis

To determine whether axitinib affects the expression of ABCG2, the cells were incubated with 0.25, 0.5, 1.0 and 2.0 μmol/L axitinib for 48 h or with 1.0 μmol/L axitinib for 24, 48 and 72 h. To test whether axitinib blocks AKT or ERK1/2 phosphorylation, the cells were incubated with different concentrations of axitinib (0.25–2.0 μmol/L) for different periods of time (6–24 h). After drug treatment, cells were lysed after washing two times with ice cold PBS. Equal

amounts of protein were resolved by SDS–polyacrylamide gel electrophoresis and transferred to nitrocellulose membranes. Then, the protein–antibody complexes were visualized by the enhanced Phototope TM-HRP (horseradish peroxidase) detection kit (Cell Signaling) and exposed to a Kodak medical x-ray processor (Kodak, Rochester, NY, USA) (31).

Reverse Transcription- and Quantitative Real-Time-Polymerase Chain Reaction

After drug treatment, total cellular RNA was isolated by use of a Trizol Reagent RNA extraction kit according to the manufacturer's instructions (Molecular Research Center, Cincinnati, OH, USA). The first strand of cDNA was synthesized by oligo dT primers with reverse transcriptase (Promega Corp., Madison, WI, USA). Oligonucleotide primers for ABCG2 and GAPDH were synthesized commercially (Invitrogen, Beijing, China). polymerase chain reaction (PCR) primers were 5'-TGG CTG TCA TGG CTT CAG TA-3' (forward) and 5'-GCC ACG TGA TTC TTC CAC AA-3' (reverse) for ABCG2 and 5'-CTT TGG TAT CGT GGA AGG A-3' (forward) and 5'-CAC CCT GTT GCT GTA GCC-3' (reverse) for GAPDH. Using the GeneAmp PCR system 9700 (PE Applied Biosystems, Norwalk, CT, USA), reactions were carried out at 94°C for 2 min for initial denaturation, and then at 94°C for 30 s, 58°C for 30 s and 72°C for 1 min. After 35 cycles of

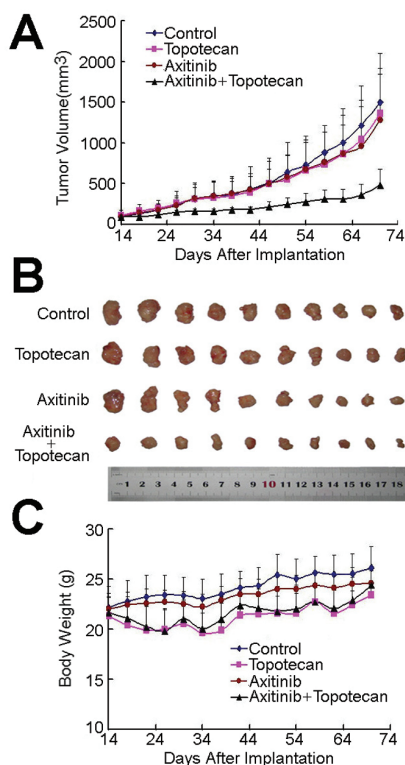


Figure 1. Potentiation of the antitumor effects of topotecan by axitinib in an S1-M1-80 cell xenograft model in athymic nude mice. (A) Changes in tumor volume with time. Each point represents the mean \pm standard deviation (SD) of tumor volumes from 10 mice in the group. (B) Tumor size. The picture was taken on the 70th d after implantation. (C) Changes in body weight with time after tumor cell inoculation. Each point represents the mean \pm SD of body weight from 10 mice in the group. The various treatments were as follows: (a) saline (every 4 d \times 9, IP); (b) topotecan (every 4 d \times 9, IP, 3 mg/kg); (c) axitinib (every 4 d \times 9, PO, 25 mg/kg); (d) topotecan (every 4 d \times 9, IP, 3 mg/kg) and axitinib (every 4 d \times 9, PO, 25 mg/kg) (axitinib was given 1 h before topotecan administration). The whole administration process was divided into three cycles with a 10-d drug-free recovery period between every two cycles.

amplification, additional extensions were carried out at 72°C for 10 min. Products were resolved and examined by 1.5% agarose gel electrophoresis. The expected length of PCR products for *ABCG2* is 235 bp and that for *GAPDH* is 475 bp.

Real-time PCR was performed with an ABI PRISM 7500 sequence detection system (Applied Biosystems, Foster City, CA, USA). Reactions were carried out at least in triplicate repeats in two independent experiments. The geometric mean of the glyceraldehyde-3-phosphate dehydrogenase (*GAPDH*) housekeeping gene was used as an internal control to normalize the variability in expression levels. All procedures were carried out according to the instructions. The forward primer for *ABCG2* was 5' CAC CTT ATT GGC CTC AGG AA-3', the reverse primer was 5' CCT GCT TGG AAG GCT CTA TG-3'. The forward primer for *GAPDH* was 5' GAG TCA ACG GAT TTG GTC GT-3', the reverse primer was 5' GAT CTC GCT CCT GGA AGA TG-3'.

Statistical Analysis

All experiments were repeated at least thrice and the differences were determined by using the Student *t* test. The statistical significance was determined at $P < 0.05$.

All supplementary materials are available online at www.molmed.org.

RESULTS

Axitinib Particularly Reversed *ABCG2*-Mediated MDR *In Vitro*

To examine the reversal activity of axitinib in *ABCB1*-, *ABCG2*-, *ABCC1*-, *ABCC4*- and LRP-mediated MDR in cancer cells, we first examined the cytotoxic activity of axitinib in different cell lines via MTT assay. As shown in Supplementary Figure S1, more than 90% of cells survived when treated with axitinib alone up to 1.0 $\mu\text{mol/L}$ in S1, S1-M1-80, KB, KBv200, SW1573, SW1573/2R120, NIH3T3 and NIH3T3/MRP4-2 cells, and up to 0.4 $\mu\text{mol/L}$ in HL60 and HL60/ADR cells. Therefore, we used 1.0 or 0.4 $\mu\text{mol/L}$ (in HL60 and HL60/ADR cells) axitinib to perform an MDR reversal assay *in vitro*. As shown in Table 1, the S1-M1-80 cells were significantly resistant to topotecan and mitoxantrone: the IC_{50} values were 12.790 ± 0.241 and $14.668 \pm$

$0.636 \mu\text{mol/L}$ in S1-M1-80 cells, respectively, which were 51- and 76-fold higher than that in S1 cells (0.250 ± 0.014 and $0.193 \pm 0.021 \mu\text{mol/L}$, respectively). Thus, overexpression of *ABCG2* resulted in a significant resistance to topotecan and mitoxantrone. Axitinib strongly enhanced the cytotoxicity of topotecan and mitoxantrone in S1-M1-80 cells in a dose-dependent manner, but had no such effect on drug-sensitive parent S1 cells. At the presence of 1.0 $\mu\text{mol/L}$ axitinib, the IC_{50} values of topotecan and mitoxantrone in S1-M1-80 cells reduced from 12.79 ± 0.241 to $3.020 \pm 0.242 \mu\text{mol/L}$ and from 14.668 ± 0.636 to $2.904 \pm 0.482 \mu\text{mol/L}$, respectively, representing a 4.11- and 5.05-fold drug sensitization. In addition, axitinib almost entirely reversed the resistance to mitoxantrone in *ABCG2*-482-G2 and *ABCG2*-482-T7 cells, which were transfected with *ABCG2* (Table 2). However, axitinib did not alter the cytotoxicity of non-*ABCG2* substrate (cisplatin) in MDR cells or their parental sensitive cells. All cell drug-resistance models with overexpression of *ABCG2* showed no detectable expression of other MDR proteins (Supplementary Figure S2). These results suggest that axitinib is an effective modulator in restoring the sensitivity of resistant cells to anticancer drugs *in vitro*.

Moreover, to investigate whether axitinib had the potential ability to reverse MDR mediated by other ABC transporters, we examined the modulator activity of axitinib on *ABCB1*/P-gp-, *ABCC1*/MRP1-, *ABCC4*/MRP4-, and LRP-mediated drug resistance in cancer cells. As illustrated in Table 1, axitinib displayed no modulating activity in KBv200, HL60/ADR, NIH3T3/MRP4-2 and SW1573/2R120 cell lines. Taken together, our data suggest that axitinib probably specifically reverses *ABCG2*-mediated MDR.

Axitinib Reversed *ABCG2*-Mediated MDR *In Vivo*

An established S1-M1-80 cell xenograft model in nude mice was used to evaluate the efficacy of axitinib to reverse the resistance to topotecan *in vivo*.

Topotecan and axitinib had minimal inhibitory activity toward S1-M1-80 tumors when administered alone. Surprisingly, the antitumor activity of topotecan was significantly enhanced when it was administered in combination with axitinib (Figure 1, $P < 0.05$ comparing topotecan to topotecan plus axitinib). The weight of tumors excised from mice were 0.704 ± 0.418 , 0.651 ± 0.280 , 0.555 ± 0.344 and 0.224 ± 0.097 g for saline, topotecan, axitinib and combination groups, respectively. The inhibition rate in the combination group was 68.2%. No significant body weight loss or treatment-related deaths occurred during the study, indicating that axitinib effectually enhanced the antitumor activity of topotecan without causing additional toxicity. The S1 cell xenograft model in nude mice was established to examine the effect of axitinib on the parental sensitive cells. As shown in Supplementary Figure S3, after treatment of the S1 cell xenograft model in the same way as the S1-M1-80 tumor model, compared with animals treated with saline or axitinib alone, both topotecan and the combination of axitinib with topotecan produced significant inhibition of tumor growth. S1 cells remained sensitive to topotecan and there was no significant difference in tumor size between topotecan and the combination group.

Axitinib Targeted to SP Cells and Enhanced the Efficacy of Chemotherapeutic Drugs in SP Cells

We examined the existence of SP cells in A549 cells by Hoechst 33342 staining to generate a Hoechst blue-red profile. The SP gate was defined as the diminished region in the presence of FTC, which blocked the activity of Hoechst 33342 dye transporter. A549 cells contained about 5.06% SP cells, which decreased significantly in the presence of FTC (Figure 2A).

To test whether SP cells isolated in our study were enriched for tumorigenic cells, we examined the tumor formation rate of the SP and non-SP cells in a xenograft model. Our results showed that

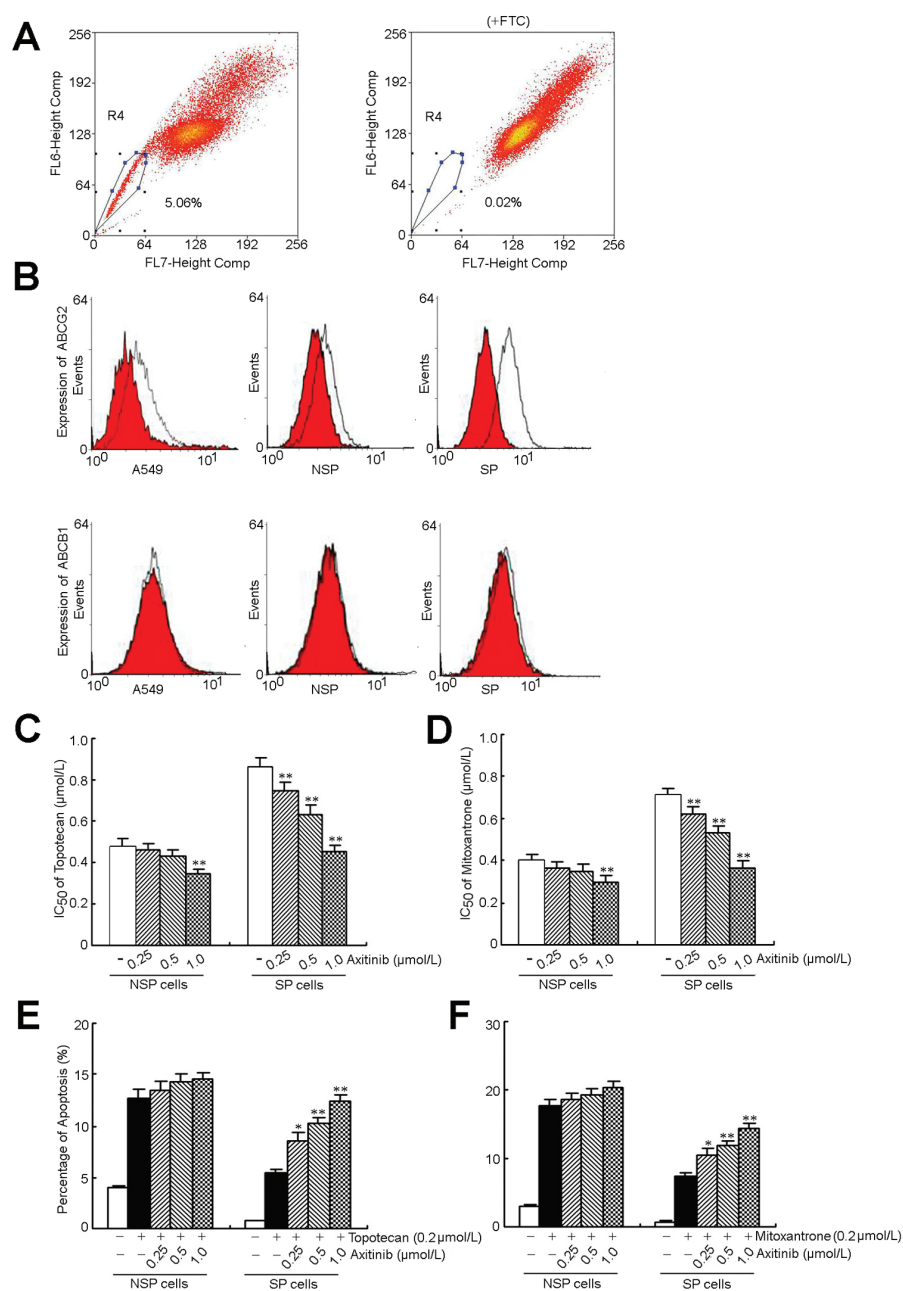


Figure 2. Axitinib targeted to SP cells and enhanced the efficacy of topotecan and mitoxantrone in the inhibition of proliferation and induction of apoptosis. (A) The A549 cells were stained with Hoechst 33342 as described in Materials and Methods. Gated on forward and side scatter to exclude debris, Hoechst red versus Hoechst blue (R2) was used to sort SP cells. (B) The cell surface expression of *ABCG2* and *ABCB1*. (C, D) Induction of 50% cell death in SP and non-SP cells by topotecan, mitoxantrone and axitinib. Growth inhibition was determined by the MTT assay according to the protocol described in Materials and Methods. (E, F) Sorted SP and non-SP cells treated with topotecan, mitoxantrone and axitinib in the indicated concentrations for 48 h, respectively. Apoptosis was analyzed by flow cytometry as the percentage of cells labeled by annexin V and propidium iodide. All of these experiments were repeated at least thrice, and a representative experiment is shown. Columns, means of triplicate determinations; * $P < 0.05$; ** $P < 0.01$, compared with topotecan or mitoxantrone treatment.

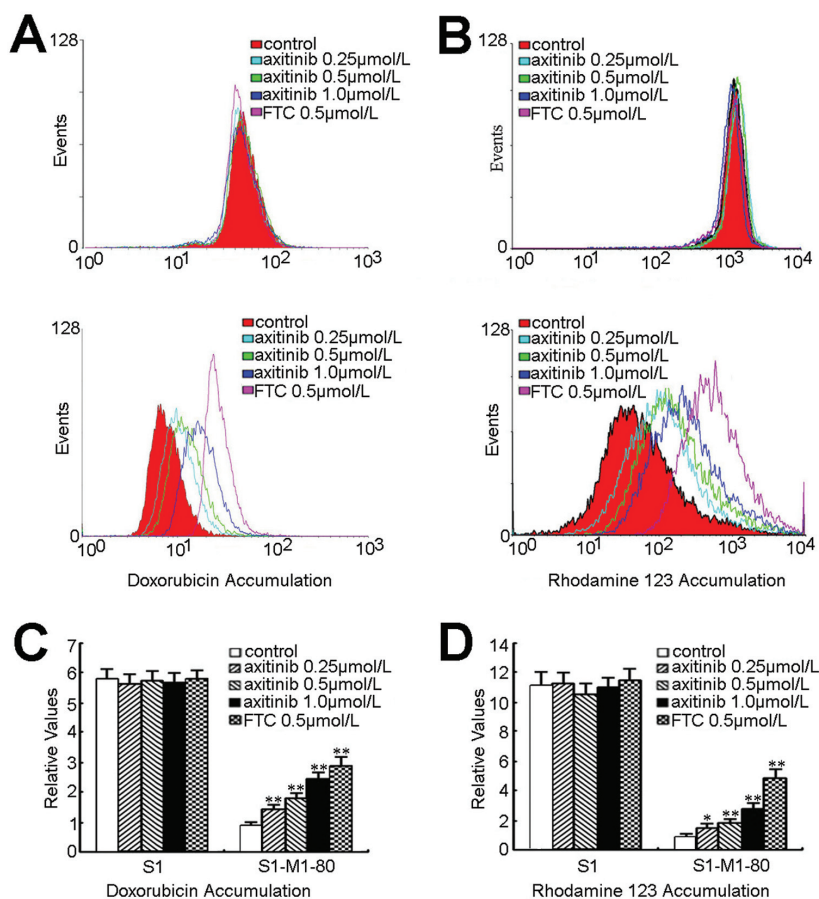


Figure 3. Effect of axitinib on the intracellular accumulation of Dox and Rho123 in S1 and S1-M1-80 cells. The accumulation of Dox (A) and Rho 123 (B) was measured by flow cytometric analysis after cells were preincubated with or without axitinib and FTC for 3 h at 37°C and then incubated with 10 $\mu\text{mol/L}$ Dox or 5 $\mu\text{mol/L}$ Rho123 for another 3 h or 0.5 h at 37°C, respectively, as described in Materials and Methods. (A) Intracellular accumulation of Dox and Rho123 was not significantly affected by axitinib in S1 cells. (B) The accumulation of Dox and Rho123 was increased in a dose-dependent manner in S1-M1-80 cells. (C, D) Dox and Rho123 levels were expressed as fold change in fluorescence intensity relative to control MDR cells. They are calculated by dividing the fluorescence intensity of each sample with that of MDR cells treated with Dox and Rho123. Columns, means of triplicate determinations; bars, standard deviation. ** $P < 0.01$ versus the MDR control group. Independent experiments were performed at least three times, and a representative experiment is shown.

the SP cells gave rise to tumors with 1×10^4 cells (one of six animals), whereas at least 1×10^6 non-SP cells were needed to form a tumor. At the same injection dose (1×10^6 cells), the tumor generated by the SP cells (389 mm^3) is 3.6-fold larger in volume than that of the non-SP cells (107 mm^3) (Supplementary Figure S4).

We next analyzed the cell surface expression of *ABCG2* and *ABCB1*. The SP cells showed much higher expression of

ABCG2 than the non-SP cells. The A549 cells also showed a low expression of *ABCG2*. All of the A549 cell subsets showed no expression of *ABCB1* (Figure 2B). Then we tested whether axitinib could enhance the cytotoxic effect of chemotherapeutics. As shown in Figure 2C, the SP cells exhibited higher resistance to chemotherapeutic drugs than non-SP cells. It is interesting to note that axitinib significantly enhanced the sensi-

tivity of SP cells to topotecan and mitoxantrone in a dose-dependent manner, but had no such effect on non-SP cells (Figures 2C, D). Axitinib had no effect on the apoptosis induced by topotecan and mitoxantrone in non-SP cells, but it drastically enhanced the apoptosis of SP cells (Figures 2E, F). Quantitative analysis of the apoptotic SP cells showed that axitinib induced an increase of the percentage of apoptotic cells in a dose-dependent manner: topotecan from $5.6\% \pm 0.9\%$ to $12.2\% \pm 1.1\%$ and mitoxantrone from $7.5\% \pm 1.1\%$ to $14.4\% \pm 1.3\%$ (Supplementary Figure S5). The results of the apoptosis assay reveal that axitinib can target to SP cells and increase the cell apoptosis induced by topotecan and mitoxantrone.

Axitinib Inhibited the Function of *ABCG2*-Mediated Transport

The results above indicated that axitinib could enhance the sensitivity of MDR cancer cells to certain *ABCG2* substrate anticancer drugs. To ascertain the potential mechanisms, we examined the effect of axitinib on the accumulation of Dox and Rho 123 in cells overexpressing *ABCG2*. In the absence of axitinib, the intracellular levels of Dox and Rho 123 were very low in MDR cells, whereas axitinib significantly increased the intracellular accumulation of Dox and Rho 123 in a dose-dependent manner (Figure 3). The fluorescence index of Dox in the presence of 0.25, 0.5 and 1.0 $\mu\text{mol/L}$ of axitinib was increased by 1.28-, 1.67- and 2.16-fold in S1-M1-80 cells, respectively (Figures 3A, C). As shown in Figures 3B, D, axitinib at 0.25, 0.5 and 1.0 $\mu\text{mol/L}$ increased the intracellular accumulation of Rho 123 by 1.38-, 1.81- and 2.91-fold in S1-M1-80 cells, respectively. However, axitinib did not alter the intracellular accumulation of Dox and Rho 123 in the parental sensitive S1 cells. Taken together, these results suggest that axitinib significantly inhibits *ABCG2*-mediated transport function.

Drug efflux function of *ABCG2* is associated with ATP hydrolysis that is stimulated in the presence of its substrates. To assess the effect of axitinib on the

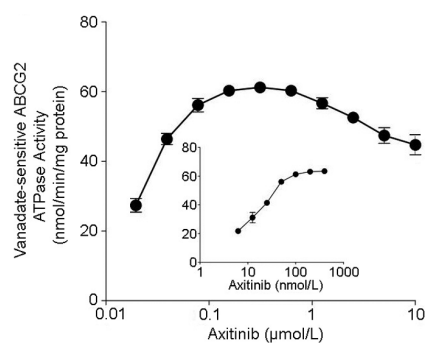


Figure 4. Effect of axitinib on the ATPase activity of *ABCG2*. The vanadate-sensitive ATPase activity of *ABCG2* was determined at different concentrations of axitinib. The inset shows the effect of lower concentrations of axitinib on *ABCG2* ATPase activity. ATP hydrolysis was monitored by measuring the amount of inorganic phosphate released using a colorimetric assay.

ATPase activity of *ABCG2*, we measured *ABCG2*-mediated ATP hydrolysis using a range of concentrations of axitinib under conditions in which the activity of other major membrane ATPases was suppressed by sodium vanadate. As shown in Figure 4, axitinib stimulated the ATPase activity of *ABCG2* in a concentration-dependent manner. A maximum *ABCG2* ATPase activity of 63.4 ± 2.8 nmol Pi/min per mg protein was attained in the presence of a low concentration of axitinib (~ 400 nmol/L) (inset of Figure 4). At a higher concentration of axitinib, a drop in the stimulated *ABCG2* ATPase activity was observed. The data suggested that axitinib may be a substrate of *ABCG2*.

Axitinib Did Not Alter the Expression Level of *ABCG2* at the mRNA or Protein Level

The reversal of *ABCG2*-mediated MDR can be achieved by either inhibiting *ABCG2* function or lowering *ABCG2* expression. Therefore, we determined the effect of axitinib on the expression of *ABCG2* at the mRNA and protein levels. S1-M1-80 cells were incubated with axitinib at $1.0 \mu\text{mol/L}$ for 24, 48 and 72 h, or at 0.25, 0.5, 1.0 and $2.0 \mu\text{mol/L}$ for 48 h.

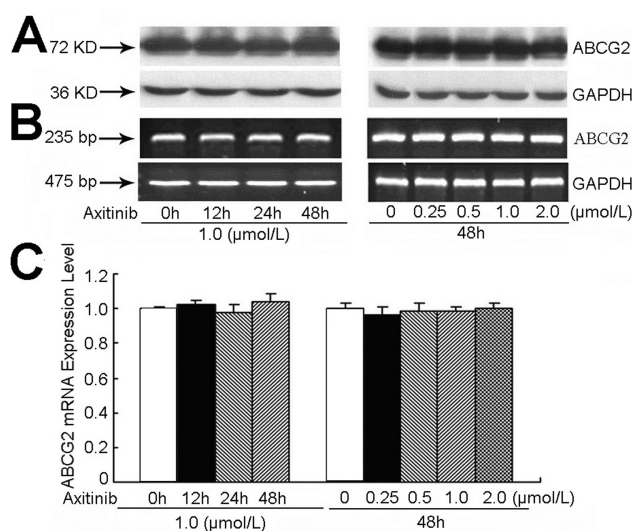


Figure 5. Effect of axitinib on the expression of *ABCG2* in MDR cells at the protein (A) and mRNA (B, C) levels. S1-M1-80 cells were treated with axitinib at $1.0 \mu\text{mol/L}$ for 12, 24 and 48 h, or at 0.25, 0.5, 1.0 and $2.0 \mu\text{mol/L}$ for 48 h, respectively. The change at mRNA level was determined by reverse transcription-PCR (B) and quantitative real-time PCR (C) as described in Materials and Methods. Independent experiments were performed at least three times and results from a representative experiment are shown.

Our results indicated that axitinib did not significantly alter the protein or mRNA expression level of *ABCG2* in S1-M1-80 cells (Figure 5). These data suggest that axitinib most likely exerts its MDR-reversal activity via direct inhibition of *ABCG2*-mediated efflux, rather than downregulation of its expression.

Axitinib Did Not Block the Phosphorylation of AKT and ERK1/2 at MDR Reversal Concentrations

Accumulating studies have shown that the inhibition of AKT and ERK1/2 pathways may decrease the resistance to anti-neoplastic drugs in cancer cells (43,44). To determine whether the concentration of axitinib used in our experiments attenuated cell survival signaling pathways, we measured the change of total and phosphorylated forms of AKT and ERK1/2 in S1 and S1-M1-80 cells. As shown in Figure 6, axitinib did not alter the total or phosphorylated forms of AKT and ERK1/2 in S1 (Figure 6A) and S1-M1-80 cells (Figure 6B). This suggests that the MDR reversal effect of axitinib in S1-M1-80 cells is independent of the

blockade of AKT and ERK1/2 signal transduction pathway.

DISCUSSION

The cancer stem cell hypothesis suggests that the formation and growth of tumors are driven by rare cancer stem cells, and increasing evidence also indicates that cancer stem cells play an important role in tumor initiation, progression and metastasis, as well as chemoresistance (45–47). Isolation and observation of CSCs have been achieved through selecting the SP cells, the subset of cells capable of effluxing the DNA-intercalating dye Hoechst 33342. SP cells have been identified in both human primary tumors and human cancer cell lines of several tissue origins, including thyroid, ovary, breast, glial cells and hepatic oval cells, and in all these cases the SP cells exhibit features of CSCs. Recent strong evidence has shown that cancer stemlike phenotypes are often correlated with expression and function of *ABCG2*, which may be responsible for their drug resistance phenotype (48–50). Elevated expression of *ABCG2* has been observed in a number of cancer stem cells

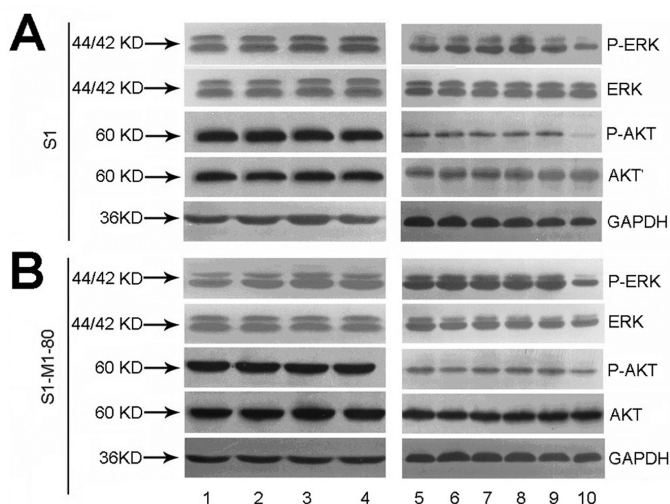


Figure 6. Effect of axitinib on the blockade of AKT and ERK1/2 phosphorylation. S1 and S1-M1-80 were treated with axitinib at 1.0 $\mu\text{mol/L}$ for 0, 3, 6 and 12 h (lanes 1–4, respectively), or at 0, 0.25, 0.5, 1.0 and 2.0 $\mu\text{mol/L}$ for 24 h (lanes 5–9, respectively), and lane 10 was the positive control (cells were treated with lapatinib at 10 $\mu\text{mol/L}$). An equal amount of protein was loaded for Western blot analysis as described in Materials and Methods. The experiments were repeated at least three times independently, and a representative experiment is shown.

isolated from retinoblastoma, pancreas, liver and lung. In addition, *ABCG2* and CD133, a widely identified CSC marker, are coexpressed in melanoma and pancreatic carcinoma. These data suggest that *ABCG2* is a promising molecular marker for identification of CSCs in tumors. New therapeutic strategies targeting *ABCG2*-positive CSCs may effectively eliminate CSCs and overcome current chemotherapeutic limitations.

Axitinib is an oral small-molecule inhibitor of VEGFR-1, -2 and -3; PDGFR and c-KIT TKs. Further studies demonstrated that axitinib alone produced remarkable antitumor efficacy associated with antiangiogenesis effects across pre-clinical models regardless of the RTK expression profile in tumor cells. Clinical trials with axitinib are showing promising antitumor activity against advanced renal cell carcinoma (7), thyroid cancer (8) and non-small cell lung cancer (6). In combination studies, additive or synergistic enhancement of TKIs and response to chemotherapeutic agents alone was observed when axitinib was combined with docetaxel, carboplatin and gemcitabine.

Importantly, combining axitinib with docetaxel generated marked suppression of disease progression compared with docetaxel alone in a docetaxel-resistant Lewis lung carcinoma model (51–55). More studies are underway to provide deeper insight into how axitinib and chemotherapeutic agents can be best used for maximal activity in animal models.

In the present study, we examined the effect of axitinib on enhancing chemotherapeutic efficacy in SP cells and the ability of axitinib to reverse MDR in drug-resistant cell lines. Our data showed that axitinib enhanced the chemotherapeutic sensitivity of topotecan and mitoxantrone and increased apoptosis induced by the two drugs in SP cells. In addition, nontoxic concentrations of axitinib produced a 4.11-fold topotecan sensitization and a 5.05-fold mitoxantrone sensitization in S1-M1-80 cells, but had no such effect in the drug-sensitive parent S1 cells, indicating that the sensitization of the resistant cells by axitinib was attributable to its specific effect on *ABCG2*.

To determine whether the favorable effects of axitinib *in vitro* can be extended

to an *in vivo* paradigm, we have examined the effect of axitinib on enhancing the antitumor activity of topotecan in S1-M1-80 cell xenograft model in mice. Consistent with the *in vitro* results, our data indicated that axitinib in combination with topotecan resulted in markedly enhanced antitumor activity of topotecan in this *ABCG2*-overexpressing tumor xenograft model and did not increase the toxic side effects (Figure 1 and Supplementary Figure S3).

To investigate the mechanisms of reversal of *ABCG2*-mediated MDR by axitinib, *ABCG2* expression and transport activity were examined. Consistent with the overexpression and therefore higher transport function of *ABCG2*, S1-M1-80 cells had lower intracellular accumulation of Dox and rhodamine 123 than S1 cells (Figure 3 and Supplementary Figure S2). Axitinib treatment significantly increased the accumulation of Dox and rhodamine 123 in a dose-dependent manner but had no effect in the parent S1 cells. We also found that axitinib stimulated the ATPase activity of *ABCG2* in a concentration-dependent manner (Figure 4), indicating that axitinib may directly interact with the drug-substrate binding site on *ABCG2*.

As shown in Supplementary Figure S4, SP cells that are isolated by their ability to efflux Hoechst 33342 dye were enriched in tumor-initiating capability compared with non-SP cells. We also found that axitinib enhanced the cytotoxicity of topotecan and mitoxantrone in SP cells *in vitro*. Kataoka *et al.* have reported that treatment of SP cells with dofequidar reversed the drug resistance of xenografted SP cells *in vivo* just as it did *in vitro* (56). Because the SP cells isolated in our study did overexpress *ABCG2* (Figure 2B), we can conclude that the *in vitro* effects of axitinib on SP cells can be extended to an *in vivo* paradigm as efficient as dofequidar. Thus it could be used in conjunction with other conventional anticancer drugs to eradicate the cancer stem cells.

Taken together, these data strongly indicated that axitinib can inhibit the transport function of *ABCG2*, thereby increas-

ing the intracellular concentration of its substrate chemotherapeutic drugs. It is possible that the downregulation of *ABCG2* expression may potentiate the reversal effect of axitinib on *ABCG2*-mediated MDR. However, axitinib treatment did not change the expression of *ABCG2* at both mRNA and protein levels (Figure 5). We thus proposed that the MDR reversal effect of axitinib was due to the inhibition of efflux function of *ABCG2* as revealed in the drug accumulation assay.

Receptor TKs such as VEGFR, PDGFR and c-Kit play a key role in modulating cell proliferation, differentiation and survival by activating downstream signal molecules such as signal transducers and activators, PI3K/AKT and ERK1/2 (57). Aberrant activation of receptor TKs is believed to be associated with cancer growth, angiogenesis and metastasis. Moreover, several studies have revealed that activation of the PI3K/AKT and/or ERK pathways is associated with resistance to conventional chemotherapeutic drugs (58,59). Our data revealed that total and phosphorylation forms of AKT and ERK1/2 remained unchanged in S1 and S1-M1-80 cells after treatment with different concentrations of axitinib (Figure 6), indicating that blockade of AKT and ERK1/2 activation was not involved in the reversal of *ABCG2*-mediated MDR by axitinib. Compared with other *ABCG2* inhibitors, axitinib is more potent and specific, which is ideal for future clinical studies. Nonetheless, as with other modulators it will be essential to evaluate the effect of the axitinib on the pharmacokinetic disposition of other antineoplastic drugs.

CONCLUSION

In conclusion, axitinib can enhance the efficacy of conventional chemotherapeutic drugs in SP cells and *ABCG2*-overexpressing MDR cells via directly inhibiting the drug transport function of *ABCG2*. Our results suggest that axitinib may be used in combination with conventional *ABCG2* substrate chemotherapeutic drugs to overcome multidrug

resistance in the clinic. It should be discussed that axitinib would be used both as an antineoplastic drug and as an MDR reversal agent in the future.

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DISCLOSURE

The authors declare that they have no competing interests as defined by *Molecular Medicine*, or other interests that might be perceived to influence the results and discussion reported in this paper.

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