

Activation of phosphatidylinositol 3-kinase/Akt signaling mediates sorafenib-induced invasion and metastasis in hepatocellular carcinoma

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Abstract. Sorafenib, an antiangiogenic agent, can promote tumor invasion and metastasis. The phosphatidylinositol 3-kinase (PI3K)/Akt/Snail-dependent pathway plays an important role in tumor invasion and metastasis. Yet, little is known concerning the role of the PI3K/Akt/Snail-dependent pathway in sorafenib-induced invasion and metastasis of hepatic carcinoma (HCC). A human HCC orthotopic xenograft model was established, and sorafenib (30 mg/kg/day) was administered orally. Tumor growth and intrahepatic metastasis were assessed, and immunohistochemistry was applied to analyze the activation of the PI3K/Akt/Snail-dependent pathway. HCC cell lines were treated with sorafenib (1, 5 and 10 μ M), and proliferation, migration and invasion were assessed. Western blotting and real-time polymerase chain reaction (RT-PCR) were used to examine the related gene expression of epithelial-mesenchymal transition (EMT) markers and the PI3K/Akt/Snail-dependent pathway. Sorafenib inhibited tumor growth and promoted intrahepatic invasion and metastasis of the orthotopic tumors grown from SMMC7721-GFP cells *in vivo*. Additionally, sorafenib promoted EMT and invasion and metastasis of HCC cells *in vitro*. Importantly, sorafenib enhanced PI3K and Akt activation and upregulation of the expression of transcription factor Snail, a critical EMT mediator. The upregulation of transcription factor Snail expression by sorafenib may be related to activation of the PI3K/AKT signaling pathway. The PI3K/Akt/Snail-dependent pathway may mediate the pro-invasive and pro-metastatic effects of sorafenib on HCC by inducing EMT.

Introduction

Hepatic carcinoma (HCC) is the fifth most common malignancy worldwide and the second leading cause of cancer-related death in Asia generally and in China in particular (1). Currently, surgical resection and liver transplantation offer the best potential for treating HCC (2-4), but most HCC patients are diagnosed in advanced stages. At present, sorafenib, a multikinase inhibitor with antiangiogenic and antiproliferative effects, currently sets the new standard for advanced HCC (5,6). However, the survival benefit is only 2.8 months.

Antiangiogenic therapy has been thought to hold significant potential for the treatment of cancer (7). However, clinical and preclinical observations indicate that these therapies may have limited efficacy. Although these agents typically produce inhibition of primary tumor growth, lasting responses are rare, with only a moderate increase in progression-free survival and little benefit in overall survival (8). In addition, recent reports describe that treatment of tumor-bearing mice with antiangiogenic drugs leads to increased local tumor cell invasion and enhanced distant metastasis after prolonged treatment or after only short-term treatment (9,10). Notably, sorafenib, the only approved molecular-targeted drug for HCC, was found to promote invasion and metastasis of HCC by increased intrahepatic metastasis, lung metastasis, and circulating tumor cells in tumors with higher expression of HTATIP2 in xenograft models (11). Therefore, it is important to clarify the molecular mechanisms of invasion and metastasis caused by sorafenib from all aspects in HCC.

Epithelial-mesenchymal transition (EMT) plays a key role in tumor invasion and metastasis. During this process, epithelial cells lose their epithelial signatures while acquiring the characteristics of mesenchymal cells including morphology, cellular structure and biological function (12). Transcription factor Snail has also been shown to confer survival properties either concomitantly with induction of EMT or independent of EMT (13-15). Snail, Slug and Twist transcription factors can act as E-box repressors and block E-cadherin transcription (16). In addition, Snail transcription factor can mediate an increase in expression of mesenchymal markers such as

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vimentin, fibronectin, matrix metalloproteinases (MMPs) and RhoA (17-20). The overall effect of Snail is increased migration and invasion (18,19).

Numerous signaling pathways are involved in the regulation of EMT. PI3K/Akt signaling is an important survival/proliferative pathway involving various growth factors, cytokines and activation of receptors (21). In addition, the PI3K/Akt signaling pathway plays a key role in the control of cell invasion and metastasis and the activation of PI3K/AKT is a central feature of EMT in the development of cancer (22-27). On the one hand, the PI3K/AKT signaling pathway can increase the expression of matrix metalloproteinases to induce EMT (28,29). On the other hand, the PI3K/AKT signaling pathway can upregulate the expression of transcription factor Snail to induce EMT (30-32). Notably, activation of the PI3K/Akt signaling pathway plays a key role in mediating resistance to sorafenib. The combination of MK-2206, an Akt inhibitor, and sorafenib overcomes such resistance (33). Yet, little is known concerning the role of the PI3K/Akt signaling pathway on the invasion and metastasis induced by sorafenib in HCC.

In the present study, we tested and verified that sorafenib promotes invasion and metastasis of HCC by inducing EMT. More importantly, we showed that activation of the PI3K/Akt/Snail-dependent pathway may play a key role in this process.

Materials and methods

Reagents and antibodies. Sorafenib was purchased from Bayer Corporation (West Haven, CT, USA). Antibodies against E-cadherin, N-cadherin, vimentin, Snail and GAPDH were purchased from Epitomics (Burlingame, CA, USA); antibodies against p-PI3K and p-AKT were purchased from Bioworld Technology (Minneapolis, MN, USA).

Cell culture. The human HCC cell lines SMMC7721 and HCCLM3 originated from the American Type Culture Collection (ATCC) and were cultured in RPMI-1640 containing 10% fetal bovine serum (FBS; Biochrom, Berlin, Germany) in 5% CO₂ at 37°C. SMMC7721-GFP cells were SMMC7721 cells transfected with green fluorescence protein (GFP) and were labeled as SMMC7721-GFP cells.

Cell proliferation, migration and invasion assays. Cell proliferation analysis was performed as previously described by us (34). Briefly, cells were plated at 5,000/well in 96-well microtiter plates and incubated overnight at 37°C in a humidified incubator containing 5% CO₂. On the following day, various concentrations of sorafenib were added to the wells, and cultures were incubated for an additional 24, 48 and 72 h. Cell viability was determined using a Cell Counting Kit-8 (Dojindo, Gaithersburg, MD, USA) according to the manufacturer's instructions. For cell migration assay, cell migration was assessed using the Transwell assay (Boyden chambers; Corning, Cambridge, MA, USA). Cells (5x10⁴) were seeded in serum-free medium in the upper chamber and allowed to migrate toward the lower chamber that contained 10% FBS. After 48 h, cells that had traveled through and adhered to the underside of the membrane were counted at x200 magnification. The cell invasion assay was carried out similarly, except

that 50 µl Matrigel (BD Biosciences, Franklin Lakes, NJ, USA) diluted 1:6 with serum-free medium was added to each well overnight before cells (2x10⁵) were seeded onto the membrane.

Animal models and treatments. Six-week-old BALBc nu/nu female mice were obtained from the Shanghai Institute of Material Medica, Chinese Academy of Science. All mice were bred in laminar flow cabinets under specific pathogen-free conditions. We followed internationally recognized guidelines on animal welfare. The study design was approved by the Animal Ethics Committee, and the experiments were undertaken in accordance with the Ethical Principles of Animal Experimentation of Fudan University. SMMC7721-GFP cells [5x10⁶/0.2 ml phosphate-buffered saline (PBS)] were subcutaneously inoculated into the right flanks of 6-week-old BALBc nu/nu female mice. After 4 weeks, non-necrotic tumor tissue was cut into 1 mm³ pieces and orthotopically implanted into the liver. Treatment was started 2 weeks after orthotopic implantation of the tumors. Mice were randomly separated into two groups with 6 mice in each group. Mice in the experimental group received 30 mg/kg/day sorafenib, whereas the control mice received vehicle alone. Animal weight was measured twice a week for 4 weeks. At the end of the experiment, mice were sacrificed, tumors were excised from each mouse, weighed and snap-frozen for further analysis.

Detection of metastasis. Tumors were excised and their largest (a) and smallest (b) diameters were measured to calculate tumor volume ($V = ab^2/2$). The livers were also excised, and green fluorescent protein-positive metastatic foci were imaged by Lumazone imaging system (Mag Biosystems, Tucson, AZ, USA). Hematoxylin and eosin staining (H&E) was further applied to detect liver metastasis.

Western blot analysis. Cells were washed with cold PBS and lysed in culture dishes using PhosphoSafe™ Extraction Reagent (Merck, Darmstadt, Germany) containing 1% protease inhibitor cocktail (EDTA-Free; Thermo, San Jose, CA, USA). Protein concentrations were then determined using Bio-Rad detergent compatible protein assays (Bio-Rad, Hercules, CA, USA). A total of 30 µg protein from each of the control and treated cell lysates was loaded on 8-12% gradient NuPAGE gels (Novex, San Diego, CA, USA), electrophoresed under reducing conditions, and transferred onto polyvinylidene difluoride membranes (0.22 Å; Millipore). Western blot analysis was carried out as previously described (34).

Immunohistochemistry. Procedures for the immunohistochemistry were previously described (35). Briefly, the tumor sections were stained with rabbit anti-p-Akt, and rabbit anti-p-PI3K at 4°C overnight. Goat anti-rabbit IgG/horseradish peroxidase was applied as the secondary antibody according to the standard protocols provided by the manufacturer. For negative controls, primary antibodies were replaced with PBS. The procedures were performed by two independent investigators, both of whom were blinded to the model/treatment type for the series of experiments.

Real-time polymerase chain reaction. RT-PCR analysis was performed as previously described by us (36). The following

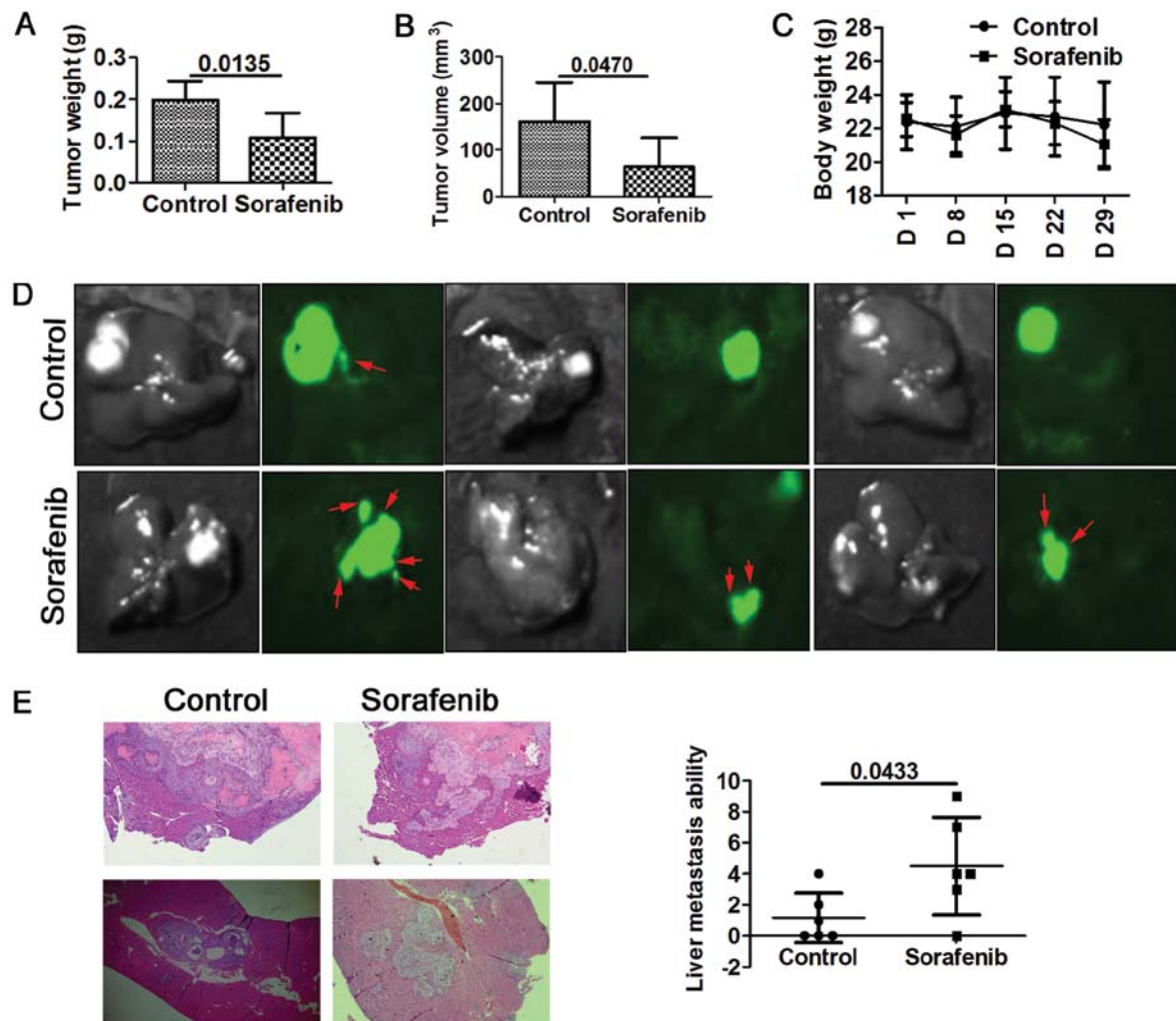


Figure 1. Sorafenib suppresses primary tumor growth and promotes intrahepatic metastasis of hepatic tumors *in vivo*. Mice were orthotopically implanted with SMMC7721-GFP cells and treated with sorafenib (30 mg/kg/day, i.g.) for four weeks. (A and B) Weight and volume of the hepatic tumors. At the end of the experiment, primary tumors from the control and sorafenib-treated mice were carefully excised. Weight and volume were measured and are presented as bar graphs. $P < 0.05$. (C) Body weights of the control and sorafenib-treated mice are presented as a line graph. Values are means \pm SD. (D) At the end of the experiment, animals from the control and sorafenib-treated groups were imaged to visualize intrahepatic metastasis by Lumazine imaging system, and representative images are presented (red arrows represent micrometastases). (E) Liver tumors and liver tissues were analyzed using H&E staining (magnification, $\times 40$). Liver metastases were quantified by counting the number of metastatic colonies in one histological section of the mid-portion of each liver sample from each mouse. Representative images and dot plots are shown. Data are expressed as the means \pm SD ($P < 0.05$). GFP, green fluorescence protein; H&E, hematoxylin and eosin staining.

primers for amplification of human genes were used: E-cadherin forward, 5'-AGCCCCGCCTTATGATTCTCTG-3' and reverse, 5'-TGCCCCATTTCGTTCAAGTAGTCAT-3'; N-cadherin forward, 5'-CCACGCCGAGCCCCAGTAT-3' and reverse, 5'-GGCCCCCAGTCGTTTCAGGTAAT-3'; vimentin forward, 5'-CCTTGACATTGAGATTGCCACCTA-3' and reverse, 5'-TCATCGTGATGCTGAGAAGTTTCG-3'; Snail forward, 5'-CAGCCTGGGTGCCCTCAAGAT-3' and reverse, 5'-GCACACGCCTGGCACTGGTA-3'.

Statistical analysis. All analyses of the results were performed using the GraphPad Prism software version 5.0 (GraphPad Software, San Diego, CA, USA) and the SPSS 19.0 software package (SPSS, Inc., Chicago, IL, USA). Statistical analyses were performed using the Student's t-test and analysis of variance (ANOVA) models. Differences were considered statistically significant at $P < 0.05$.

Results

Sorafenib promotes invasion and migration *in vivo*. To elucidate the effects of sorafenib on HCC invasion and migration, mice were orthotopically implanted with SMMC7721-GFP cells and treated with 30 mg/kg/day sorafenib. Tumor growth and metastasis were monitored. Our results showed that sorafenib substantially reduced the primary tumor growth compared with the control tumors. Tumor weight and volume were reduced in the sorafenib-treated mice (Fig. 1A and B). Additionally, sorafenib was well tolerated by the mice as no apparent weight loss was noted (Fig. 1C). Unfortunately, sorafenib-treated mice developed more intrahepatic metastatic lesions and exhibited irregular tumor margins as detected by green fluorescence imaging (Fig. 1D). To further explore the effect of sorafenib on the invasion and metastasis of HCC, liver metastatic nodules were evaluated by H&E staining as observed under a micro-

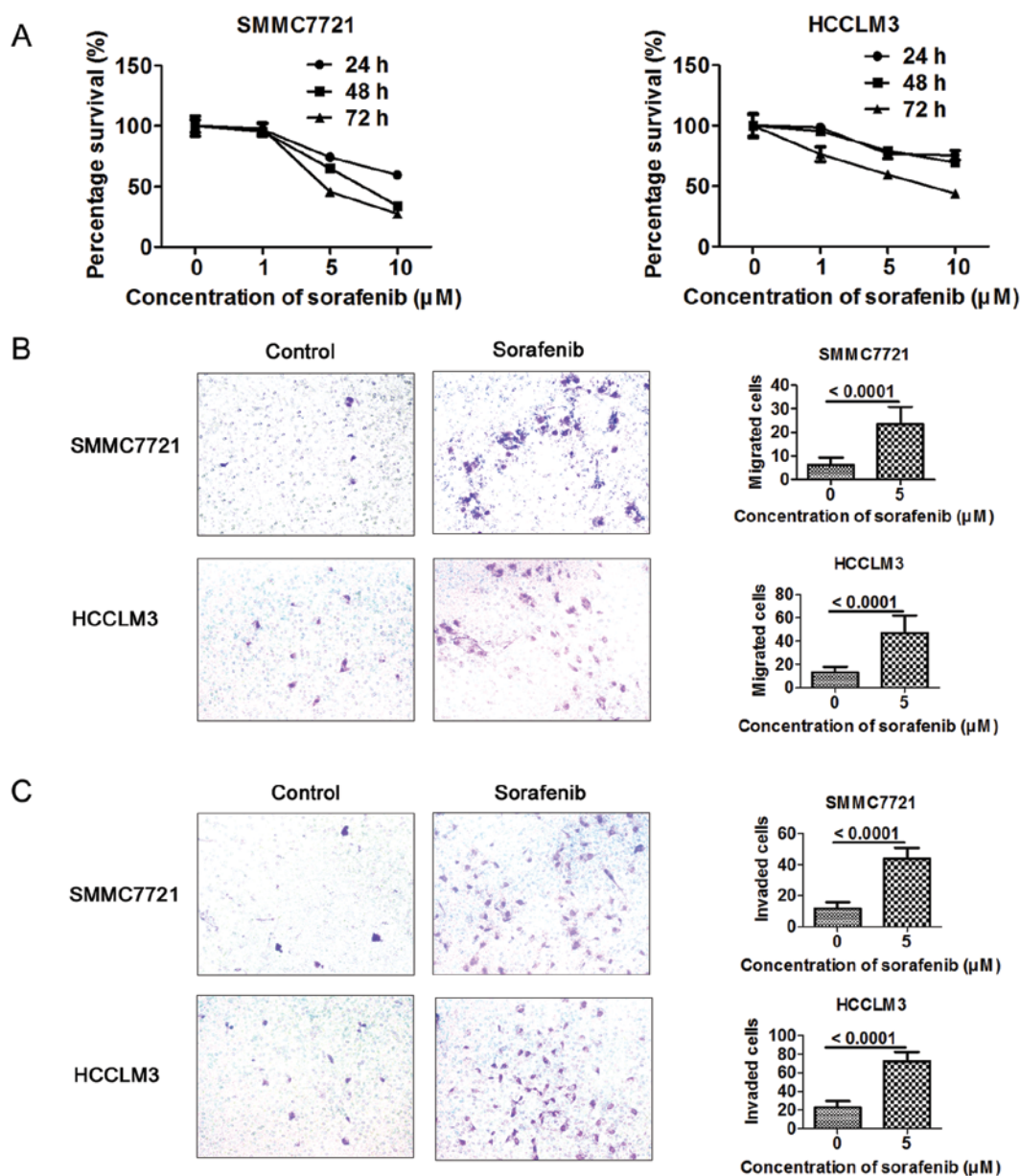


Figure 2. Sorafenib inhibits proliferation and promotes migration and invasion of HCC cells. (A) SMMC7721 and HCCLM3 cells were plated in a 96-well plate, treated with various concentrations of sorafenib and cell viability was measured by Cell Counting Kit-8. (B) Migration of HCC cells was measured by Transwell assay according to the manufacturer's instructions. Values are means \pm SD ($P < 0.05$). (C) Invasion of HCC cells was also measured by Transwell assay according to the manufacturer's instructions. Values are means \pm SD ($P < 0.05$).

scope. The number of metastatic nodules was then statistically analyzed. A higher number of intrahepatic metastatic nodules was detected in the sorafenib-treated mice (Fig. 1E).

Sorafenib promotes invasion and migration of HCC cells. As sorafenib promoted invasion and migration *in vivo*, we wanted to further validate whether sorafenib could promote the invasion and migration *in vitro*. Cell proliferation assay was applied to assess the proliferative effect on hepatoma cells after sorafenib treatment. The antiproliferative effect of sorafenib on HCC cells was dose- and time-dependent at a concentration of 1-10 μ M in the SMMC7721 and HCCLM3 cells (Fig. 2A). Sorafenib at a concentration of 5 μ M, with little effect on cell proliferation, was applied to assess the effect of sorafenib on the migration and invasion of HCC cells.

Cells (5×10^4 or 2×10^5) were seeded in the upper chamber. A higher number of metastatic and invasive cells were detected in the sorafenib-treated HCC cells as assessed by Transwell assay (Fig. 2B and C).

Sorafenib promotes EMT in HCC cells. EMT is well known to closely correlate with cancer metastasis. To test and verify whether 5 μ M sorafenib promotes the EMT process, we evaluated the expression of EMT markers in the sorafenib-treated and the control cells. As expected, SMMC7721 and HCCLM3 cells treated with 5 μ M sorafenib underwent significant morphological changes and displayed the mesenchymal phenotype (Fig. 3A). Importantly, epithelial marker E-cadherin was downregulated and mesenchymal markers N-cadherin and vimentin were upregulated in the sorafenib-

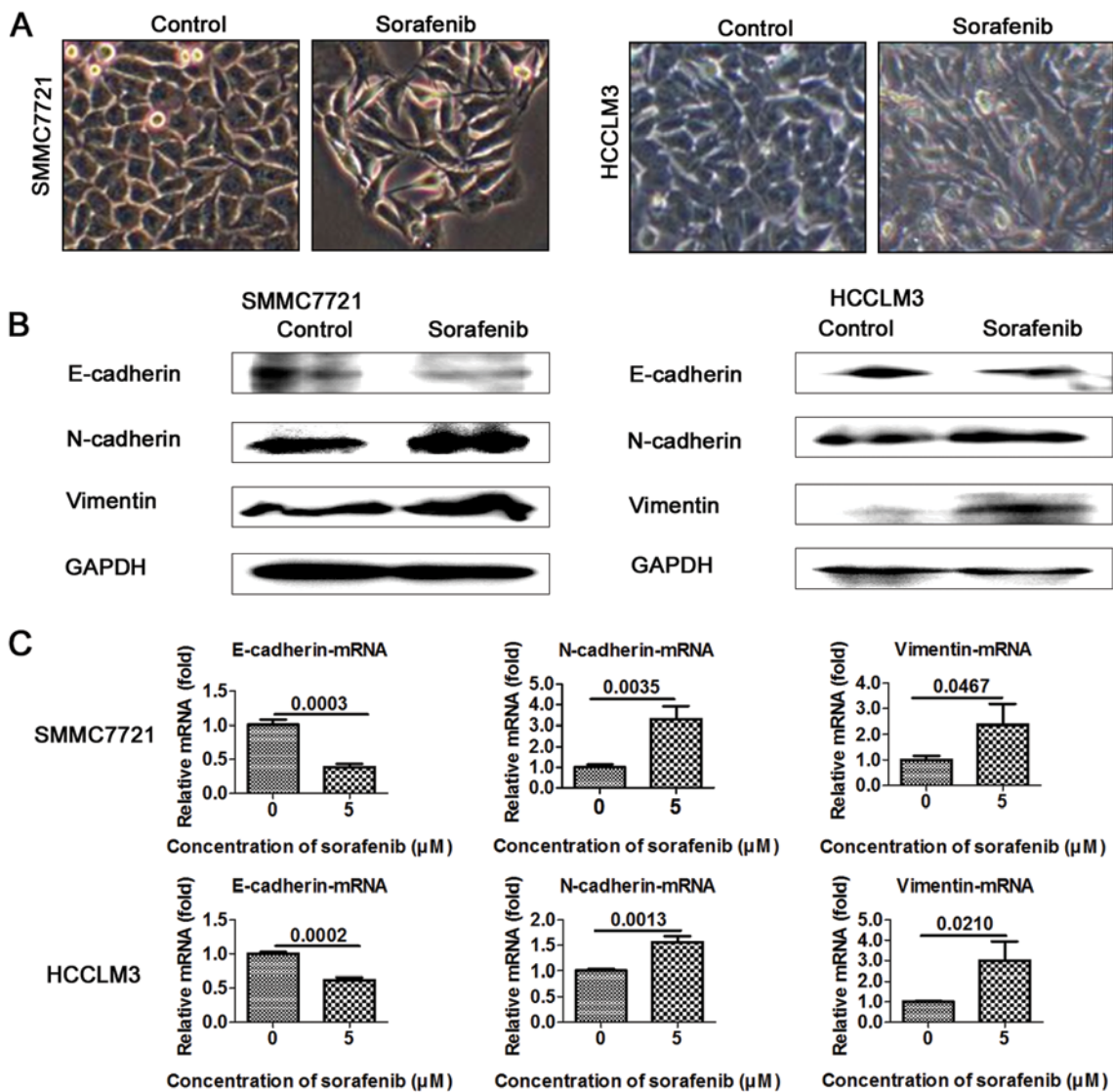


Figure 3. Sorafenib promotes EMT *in vitro*. Cells were treated with 5 μ M sorafenib, and EMT markers were evaluated by western blotting and RT-PCR. (A) After 72 h of treatment, morphological changes were evaluated in the SMMC7721 and HCCLM3 cells. (B and C) EMT markers, including E-cadherin, N-cadherin and vimentin, were assessed by western blotting and RT-PCR in the SMMC7721 and HCCLM3 cells. GAPDH was used as a loading control in the western blot analysis. EMT, epithelial-mesenchymal transition.

treated cells (Fig. 3B). RT-PCR assay further confirmed the decreased levels of epithelial marker E-cadherin and the increased levels of mesenchymal markers N-cadherin and vimentin in the SMMC7721 and HCCLM3 cells (Fig. 3C).

Sorafenib upregulates the expression of Snail in vitro. As zinc-finger transcriptional repressor Snail plays a key role in EMT-mediated tumor invasion and metastasis, we ascertained whether Snail is involved in sorafenib-mediated EMT. HCC cells were treated with 5 μ M sorafenib and western blot analysis and RT-PCR were carried out to measure Snail expression. As anticipated, transcription factor Snail was upregulated in the SMMC7721 and HCCLM3 cells, when compared to the controls (Fig. 4A and B).

Sorafenib activates the PI3K/AKT signaling pathway in vivo and vitro. As a highly conserved cellular program, EMT has been documented to involve several important pathways.

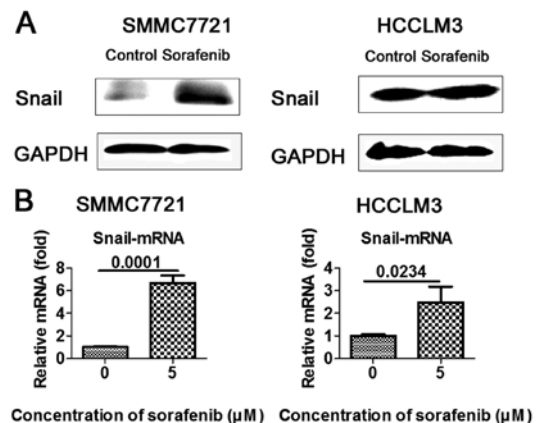


Figure 4. Sorafenib promotes the expression of transcription factor Snail *in vitro*. (A and B) After 72 h of treatment with 5 μ M sorafenib, the expression of transcription factor Snail was assessed by western blotting and RT-PCR in SMMC7721 and HCCLM3 cells. GAPDH was used as a loading control in the western blot analysis.

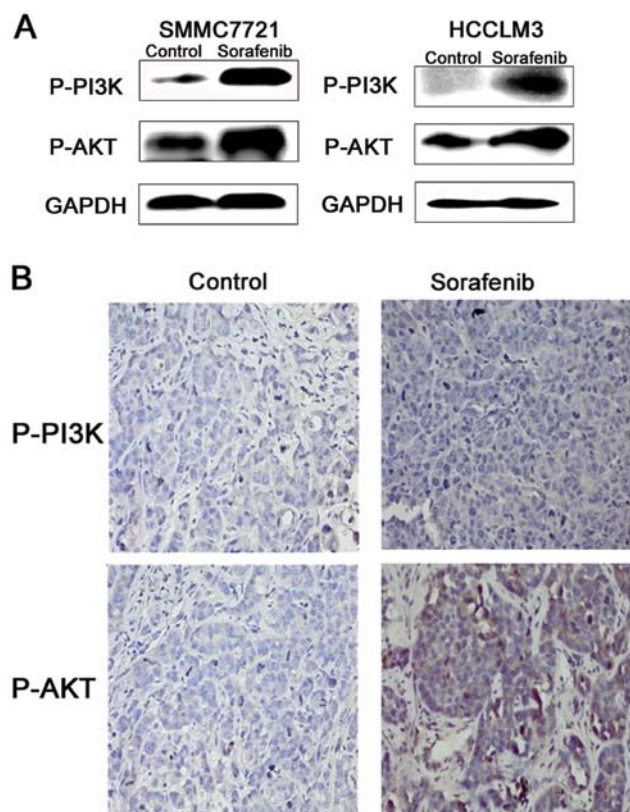


Figure 5. Sorafenib activates the PI3K/AKT signaling pathway *in vitro* and *in vivo*. (A) After 72 h of treatment with 5 μ M sorafenib, the phosphorylation of PI3K/AKT was analyzed by western blotting in SMMC7721 and HCCLM3 cells. GAPDH was used as a loading control in the western blot analysis. (B) At the end of the experiment, the control and sorafenib-treated tumors were excised for immunohistochemistry for P-PI3K and P-AKT.

Accumulating research suggests that PI3K/Akt activation plays a pivotal role in tumor progression via induction of EMT. The PI3K/Akt/GSK-3 β /Snail-dependent signaling pathway is involved in HCC. Thus, we detected the activity of PI3K/AKT in the sorafenib-induced invasion and metastasis of HCC. The results showed that the PI3K/AKT signaling pathway was activated in the HCC cells treated with 5 μ M sorafenib (Fig. 5A). In addition, the marginal tissues of the xenografts were analyzed by immunohistochemical staining as described earlier. PI3K/AKT phosphorylation levels were also upregulated (Fig. 5B).

Discussion

As a result of the SHARP and ORIENTAL trials, sorafenib has become the new standard therapy for patients with advanced hepatic carcinoma (HCC) (5,6). However, the survival benefit is only a few months. Furthermore, tumors may progress during sorafenib treatment (9-11). In the present study, we demonstrated that sorafenib exerted an antitumor effect and inhibited tumor growth in mouse models of cancer. However, sorafenib also promoted invasion and metastasis of HCC in this tumor model by inducing EMT. Similar observations were reported by other authors. Importantly, we found that sorafenib upregulated the expression of transcription factor Snail and activated the PI3K/Akt signaling pathway.

In a previous study, increased local invasion and distant metastasis during or after treatment with sorafenib were observed (11). EMT plays a key role in tumor invasion and metastasis. EMT is also reported to be involved in the progression of HCC and is correlated with the prognosis of patients (37). In the present study, more metastatic lesions in the livers of nude mice were detected in the sorafenib treatment group. In addition, HCC cell lines, including SMMC7721 and HCCLM3, were treated with 5 μ M sorafenib, with little effect on cell proliferation as confirmed by Cell Counting Kit-8. Surprisingly, morphology of the cells underwent significant changes and presented a mesenchymal phenotype after treatment for 72 h. Then EMT-related markers were analyzed. As anticipated, mesenchymal markers were significantly upregulated and epithelial markers were markedly decreased in the sorafenib-treated cells. Transwell assay was also used to analyze the ability of hepatoma cell invasion and migration. Invasion and migration capacity of the HCC cell lines was enhanced. Therefore, these data indicate that sorafenib may promote HCC invasion and metastasis by the induction of EMT, consistent with other reports.

The Snail transcription factor, a member of the Snail superfamily, is a zinc finger protein that can mediate EMT through downregulation of cell adhesion molecules such as E-cadherin by binding several E-boxes located in the promoter region (16). Snail has also been shown to confer survival properties either concomitantly with induction of EMT or independent of EMT. Snail plays an important role in inducing EMT in HCC cells (38). In cancer patients, an EMT-phenotype transcriptome profile, with increased Snail expression correlates with invasive tumors. Phosphorylation and subsequent degradation of Snail is controlled by GSK-3 β , which is predominantly regulated by PI3K/Akt (39). The PI3K/Akt/GSK-3 β /Snail-dependent signaling pathway can mediate invasion and metastasis of HCC (40,41). Increasing evidence also demonstrates that activation of the PI3K/Akt pathway plays a central role in the EMT process and correlates with an aggressive phenotype in several types of malignancies (22-27). Several signaling pathways that induce EMT and metastasis often converge at or activate PI3K/Akt, which itself is frequently activated during tumor progression. Hyperactivation of Akt is closely associated with elevated invasion and metastasis, resulting in a poor prognosis and a greater probability of relapse in many different cancer types (42-46). The PI3K/Akt signaling pathway plays a key role in invasion and metastasis of HCC. It was therefore of significance to investigate whether the PI3K/Akt/Snail-dependent signaling pathway participates in sorafenib-induced EMT. The PI3K/Akt signaling pathway was analyzed in the human HCC SMMC7721 and HCCLM3 cells. Notably, we found that 5 μ M sorafenib activated the PI3K/Akt signaling pathway and upregulated the expression of transcription factor Snail. Immunohistochemical staining was then applied to the xenograft marginal tissues. As anticipated, these results were further confirmed *in vivo*.

In conclusion, the present study showed that sorafenib upregulated the expression of transcription factor Snail and activated the PI3K/AKT signaling pathway. Importantly, these may be associated with sorafenib-induced invasion and metastasis of HCC. Therefore, inhibition of the expres-

sion of transcription factor Snail or combined with PI3K/AKT signaling pathway inhibitors may enhance the effectiveness of sorafenib treatment of HCC. Currently, relevant studies are being carried out. The present study may provide the theoretical basis for the combined treatment of sorafenib and PI3K/AKT signaling pathway inhibitors to treat HCC.

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