Inhibition of autophagy enhances the targeted therapeutic effect of sorafenib in thyroid cancer

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Abstract. The multi-target kinase inhibitor sorafenib has been approved for the treatment of patients with advanced differentiated thyroid cancer. However, different sensitivities to sorafenib have been observed, and few patients have benefited from sorafenib treatment in the long term. In the event of acquired resistance to sorafenib it is not beneficial to continue treatment in most patients. Autophagy can be induced in a variety of cancer treatments and plays an important role in cancer treatment. The role of autophagy in sorafenib treatment of thyroid cancer has not been fully demonstrated. The present study investigated whether autophagy is activated by sorafenib during the treatment of thyroid cancer, examined the underlying mechanisms, and explored potential strategies to enhance the therapeutic sensitivity of sorafenib. Chloroquine (CQ) is an autophagy inhibitor that has been reported to increase sensitivity to various cancer treatments. Thyroid cancer xenograft model mice were treated with sorafenib, CQ, or a combination of sorafenib and CQ. We observed that CQ or sorafenib treatment suppressed tumor growth, while mice treated with the combination of sorafenib and CQ displayed significantly reduced tumor growth compared with those treated with sorafenib or CQ alone. Western blotting results indicated that sorafenib concurrently inhibited the activities of the MAPK and AKT/mTOR pathways in thyroid cancer. Autophagy was activated by sorafenib in thyroid cancer, both in vitro and in vivo, which was at least in part due to suppression of the AKT/mTOR pathway. Combination treatment including CQ could inhibit the autophagic flux induced by sorafenib. Silencing the key autophagy gene ATG5 using small interfering RNA also increased the anticancer effect of sorafenib. In summary, the present study revealed that inhibition of autophagy enhances the anticancer effect of sorafenib, and the combination of CQ with sorafenib treatment represents a potential therapeutic strategy for treating advanced differentiated thyroid cancer.

Introduction

A marked increase in the number of cases of thyroid cancer has been consistently observed in several countries, including China. It is estimated that approximately 90,000 new thyroid cancer cases were diagnosed in China in 2015 (1). More than 90% of thyroid cancer cases are differentiated thyroid cancers (DTC). Some DTC patients with distant metastasis are resistant to 131Iiodine. Traditional cancer treatments are usually not effective for 131Iiodine-resistant DTC patients, who have shorter overall survival times than 131Iiodine-responsive DTC patients (2,3).

The biological behaviors of tumors are driven by oncopgenes (4). The MAPK and PI3K/AKT pathways are two important pathways driving thyroid tumor development and progression (5). The activities of BRAF and RAS, two key components of the MAPK pathway, affect cancer cell proliferation and resistance to anticancer treatment (6). Multiple genes in the MAPK pathway are targeted by sorafenib, a kinase inhibitor (7,8). Sorafenib was recently approved as the first targeted drug for advanced metastatic DTC by the Chinese Food and Drug Administration. However, sorafenib only improves progression-free survival for ~5 months compared with patients on a placebo (9), and the majority of patients acquire resistance to sorafenib after 1 or 2 years of administration, which is an obstacle in clinical practice (10).

Autophagy is an evolutionarily conserved process in which cytoplasmic materials are delivered to lysosomes for degradation (11). The relationship between autophagy and cancer is complex due to the dual roles of autophagy in tumor progression and cancer therapy (11,12). Several studies have reported that sorafenib induced autophagy in cancer cells (13), and it has been reported that autophagy can be induced by sorafenib in a dose- and time-dependent in several hepatocellular carcinoma (HCC) cell lines (14,15). It has also been observed that sorafenib modulated the expression of LC3, Beclin-1, Atg5 and Atg12, and decreased the expression of p62 in HCC cells in vitro (16,17). Several studies have reported that sorafenib inhibited AKT phosphorylation in renal cancer cells and
HCC, as reviewed by Prieto-Dominguez et al (13). In addition, inhibition of AKT enhanced sorafenib-induced autophagy, and switched protective autophagy to autophagic cell death in HCC (17). Although the role of autophagy in cancer has been extensively investigated, its role in the treatment of thyroid cancer by sorafenib has not been clearly determined.

We previously observed that sorafenib exerted a therapeutic effect by targeting the MAPK and AKT/mTOR pathways. However, the AKT/mTOR pathway is a negative regulator of autophagy and the role of autophagy in cancer treatment is complex (17,18). In this study, we aimed to investigate whether autophagy was induced by sorafenib treatment in thyroid cancer cells and to elucidate whether autophagy inhibition increases or reduces sorafenib-induced thyroid cancer cell death.

Materials and methods

Cell culture and materials. The human thyroid cancer cell lines 8505C and FTC133 were cultured at 37°C in Dulbecco's modified Eagle's medium (Gibco; Thermo Fisher, Shanghai, China) supplemented with 10% fetal bovine serum (Gibco; Thermo Fisher), 100 U/ml penicillin and 100 g/ml streptomycin in a humidified incubator with 5% CO₂. Antibodies against Beclin-1 (mouse, 1:1,000, SC-10086), GAPDH (mouse, 1:1,000, SC-51907), p-ERK (mouse, 1:200, SC-7383), ERK (rabbit, 1:1,000, SC153) and AKT (mouse, 1:1,000, SC-5298) were purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA), and the anti-p-AKT (Ser473) antibody was purchased from Abcam (rabbit, 1:200, Cambridge, Massachusetts, USA). Antibodies against cleaved-caspase-3 (rabbit, 1:1,000, #9661), p62 (rabbit, 1:1,000, #88588), LC3 (rabbit, 1:1,000, #12741), p70s6 (rabbit, 1:1,000, #2708) and p-p70s6 (rabbit, 1:500, #2211) were purchased from Cell Signaling Technology, Inc. (Danvers, MA, USA). BEZ-234 was purchased from Selleck (Shanghai, China), and sorafenib was generously donated by the Bayer Company (Shanghai, China). Cells were transfected with scrambled (negative control) small interfering (si)RNA (5'-UUCUCCGAACGUGUCACGUTT-3' and 5'-ACGUGACACUUCGGAGATT-3'), AKTsiRNA (5'-GACGGGCAUAAGAGUCUUGGTT-3' and 5'-UAGGAUUGUGCCGUTT-3'), or ATG5 siRNA (5'-GUCCAUCAAGGAUGCAUUTT-3' and 5'-AUUCAAGGUAGGACTT-3') and purchased from GenePharma (Shanghai, China). Transfection was performed using Lipofectamine™ 2000 (Invitrogen, Shanghai, China).

Mouse xenograft. Nude BALB/c mice (5-6 weeks old; male; Animal Core Facility of Nanjing Medical University, Nanjing, China) were used to generate xenograft tumor models. All animal studies complied with the management rules of the National Health and Family Planning Commission of China and were approved by the Ethics Committee of Zhejiang Cancer Hospital. A suspension of 5x10⁶ 8505C or FTC133 cells in 100 µl PBS was inoculated subcutaneously into the right flanks of the mice. When the tumor sizes averaged approximately 5x5 mm, the mice were randomly divided into four groups that were matched for tumor volume, and treatment was initiated (6 mice per treatment group). Treatment groups were as follows: the vehicle control group (DMSO, ≤0.1%), the chloroquine group (CQ; Sigma-Aldrich, Shanghai, China), the sorafenib group, and the sorafenib + CQ group. Sorafenib was administered at doses of 30 mg/kg every day for 21 days. CQ was administered at doses of 60 mg/kg every day for 21 days. In the sorafenib + CQ group, sorafenib was administered 20 min after CQ administration. All drugs were administered via intraperitoneal injection. A caliper was used to assess the tumors every 4 days. After 21 days of treatment, the mice were sacrificed and the tumor tissues were harvested for study. The mice were sacrificed by anesthetizing with an intraperitoneal injection of 0.8% pentobarbital sodium (60 mg/kg), followed by cervical dislocation. All efforts were made to reduce pain experienced by the mice.

Western blot analysis. Tumor tissues or cell pellets were lysed in RIPA lysis buffer and a protease inhibitor cocktail (Sigma-Aldrich). Total protein samples were separated by 10% or 12% sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and transferred onto nitrocellulose membranes. The membranes were blocked with 5% non-fat dry milk and incubated overnight with the various primary antibodies at 4°C. Goat anti-mouse (sc-2005) or anti-rabbit (sc-2004) antibodies were purchased from Santa Cruz Biotechnology, Inc. and diluted to 1:1,000 in 5% skim milk, Tris-HCI (pH 7.5) and 0.1% Tween-20. The immunoblots were subsequently washed and incubated with the goat anti-mouse or anti-rabbit antibody for 1 h at room temperature. The results were visualized using chemiluminescence reagent (sc-2048, Santa Cruz Biotechnology, Inc.).

RNA extraction and reverse transcription-quantitative PCR (RT-qPCR). Total RNA was isolated using an RNA Prep Pure kit (Tiangen, Beijing, China), according to the manufacturer's instructions. The synthesis of cDNA was performed with HiScript II Q RT Super Mix for qPCR (+gDNA wiper) (Vazyme Biotech Co., Nanjing, China). qPCR was performed with an AceQ qPCR SYBR Green Master Mix (without ROX) (Vazyme Biotech Co.). GAPDH was used as an internal control. Primer sequences used in the qPCR analysis were purchased from GenePharma (Shanghai, China), and the sequences were as follows: AKT, 5'-CCAACACCTTTCACTAACC-3' and 5'-CTCTCTTTTCTTCTTCT-3'; ATG5, 5'-GCAGATGGACAGTTGCACACA-3' and 5'-TTTTCCCATCTTCAGATCAA-3'; and GAPDH, 5'-CGGAGTCAACGGGATAATTTGCTGTAT-3' and 5'-AGCCCTTCTCCATGGTGGTGAGAC-3'. Melt curve was made to determine the optimal condition (95°C 15 sec, 60°C 30 sec, 95°C 15 sec). The PCR protocol is as follows: denaturation 95°C for 5 min, then 40 amplification cycles (95°C for 5 sec and 60°C for 31 sec, at a ramp-rate of 1.6°C/sec).

Apoptosis assay. 8505C and FTC133 cells (2x10⁵/well) were seeded into a 6-well plate. After treatment with ATG5Ri mimic, sorafenib (10 µM), BEZ-235 (0.8 µM) or combined sorafenib + BEZ-235, the cells were harvested and analyzed for apoptosis by Annexin V and propidium iodide staining, using an FITC Annexin V Apoptosis Detection kit (Life Technologies, Waltham, MA USA), according to the manufacturer's instructions, and a flow cytometer (FACS, Beckman Coulter, Miami, FL, USA).
Immunohistochemistry (IHC). Paraffin-embedded mouse tumor samples were sliced into 5-µm sections and mounted on microscope slides. Slides were incubated at 37°C overnight for deparaffinization. The slides were dipped in dimethylbenzene three times (10 min/dip). The slides were then immediately dipped in 100, 95, 90, 80, 70 and 50% alcohol continuously for 2 min at each percentage (%) of alcohol. Antigen retrieval was performed by heating for 15 min in 0.01 M sodium citrate buffer (pH 6.0). The slides were washed in TBS three times (5 min/wash), treated with 3% H2O2 and blocked with 5% BSA for 20 min, then incubated with 50 µl cleaved-caspase-3 (1:200 dilution) at 4°C overnight. Subsequently, the slides were washed, and incubated with a mouse anti-rabbit antibody for 1 h at room temperature. A DAB horseradish peroxidase color development kit (ZSGB-BIO, Beijing, China) was used to detect positive staining. The slides were also counterstained with hematoxylin for 25 sec, washed under running water for 3 min, then washed with distilled water for 1 min followed by dipping in 50, 70, 80, 90 and 100% alcohol continuously for 1 min each. Finally, the slides were dipped in dimethylbenzene for 3 min, air-dried and mounted with Permount™ Mounting Medium (eBioscience, San Diego, CA, USA).

Cell proliferation and viability assays. 8505C and FTC133 cells were plated at a density of 1x10^4 cells/well in 96-well microtiter plates, and each plate was incubated for 24 h at 37°C in 5% CO₂. Following treatment, the absorbance of the contents of each well was assessed at 470 nm following the use of a CCK-8 kit (Beyotime, Guangzhou, China).

Statistical analysis. Data are presented as the mean ± standard error of the mean. The results were analyzed using One-way ANOVA, and multiple comparison using Student-Newman-Keuls (SNK) and least significant difference (LSD) tests. P<0.05 was considered to indicate a statistically significant difference.

Results

CQ increases the therapeutic effect of sorafenib and suppresses tumor growth in xenograft models. To investigate whether autophagy influences the therapeutic efficacy of sorafenib in vivo, 5-6-week-old male BALB/c nude mice were used as an animal model. CQ is a potential autophagy inhibitor that has been used to regulate autophagy in vivo and in vitro (19,20). Mice were randomly divided into control, sorafenib (30 mg/kg/day), CQ (60 mg/kg/day), or sorafenib + CQ groups (n=6/group) at day 7 after subcutaneous injection of 5x10^6 cancer cells. Tumor volumes were monitored every 4 days, and the mice were sacrificed after 21 days of treatment for isolation of tumor tissues (Fig. 1A and B). Compared with the tumor sizes in the control group (1,280.93±245.27 mm³ in 8505C xenograft mice; 2,176.57±343.82 mm³ in FTC133 xenograft mice), the tumor volumes significantly decreased after administration of CQ or sorafenib alone (602.36±101.18 and 698.61±54.53 mm³ in 8505C xenograft mice; 600.42±107.33 and 697.07±171.34 mm³ in FTC133 xenograft mice), the tumor volumes significantly decreased after administration of CQ or sorafenib alone (602.36±101.18 and 698.61±54.53 mm³ in 8505C xenograft mice, respectively; 600.42±107.33 and 697.07±171.34 mm³ in FTC133 xenograft mice, respectively) (Fig. 1C and D), while the combined treatment of CQ + sorafenib significantly suppressed tumor volumes compared with CQ or sorafenib treatments alone in the 8505C and FTC133 xenograft mice (358.99±86.54 and 344.36±83.36 mm³, respectively) (Fig. 1C and D). There were no significant differences between CQ and sorafenib treatment groups in 8505C or FTC133 xenograft mice.
Sorafenib induces autophagy and apoptosis, and combined treatment with CQ increases autophagy and apoptosis. 8505C and FTC133 xenograft mice were sacrificed and the tumor tissues were isolated for further studies after 21 days of treatment. The proteins LC3 and p62 were used as autophagy markers for monitoring autophagic flux and were examined by western blotting of lysed tumor tissues. The expression level of LC3II/GAPDH increased significantly in the sorafenib treatment group compared with the control group, while the expression level of p62 was reduced in the sorafenib treatment group compared with the control group (Fig. 2A-C). To confirm the effect of sorafenib on autophagy in thyroid cancer, we treated 8505C and FTC133 cell lines with various concentrations of sorafenib in vitro for 24 h. With
an increase in sorafenib concentration, LC3II/GAPDH levels increased significantly compared with the control. When treated with 10 µM sorafenib, p62 protein levels were obviously reduced compared with the control (Fig. 2D-F). These results indicated that sorafenib increased autophagic flux in 8505C and FTC133 xenograft mouse tumors and in vitro experiments. In CQ-treated tumor tissues, the expression level of LC3II/GAPDH increased significantly compared with the control. However, the expression level of p62 in the CQ treatment group increased significantly compared with the control group, which appeared to be contrary to the sorafenib group (Fig. 2A-C). These results indicated that autophagic flux was inhibited by CQ.

Caspase-3 is activated in apoptotic cells by both extrinsic and intrinsic pathways (21,22). We used caspase-3 activation to assess apoptosis since caspase-3 is an early marker for apoptosis (23). To assess the effect of sorafenib and CQ on tumor apoptosis, we used IHC analysis of cleaved caspase-3 in different treatment groups of mice. As expected, in the sorafenib- and CQ-treated groups, the level of cleaved caspase-3 increased compared with the control, suggesting CQ or sorafenib induced apoptosis in 8505C and FTC133 xenograft mouse tumors. In the CQ + sorafenib combined treatment group, a further increase of cleaved caspase-3 was observed compared with CQ- or sorafenib-treated groups in both 8505C and FTC133 xenograft mice (Fig. 2G). Our results demonstrated an activation of caspase-3 in 8505C and FTC133 xenograft mouse tumors, suggesting that caspase-mediated apoptosis occurs following sorafenib and CQ treatment.

Sorafenib concurrently inhibits activity of the MAPK and AKT/mTOR pathways in thyroid cancer. Tumor tissues were used to analyze the target-suppressing effects of sorafenib. As shown in Fig. 3A-D, compared with the control groups, the
expression of the ERK protein was not significantly different in the sorafenib treatment groups of 8505C and FTC133 xenograft mice. However, phosphorylation levels of ERK were significantly inhibited in both 8505C and FTC133 xenograft mice that were treated with sorafenib. Furthermore, we observed that the phosphorylation levels of AKT and p70s6k were significantly reduced in the sorafenib treatment groups compared with control groups. The results indicated that CQ exerted similar effects to sorafenib. The phosphorylation levels of ERK, AKT, and p70s6k were significantly reduced after CQ treatment. Furthermore, in 8505C and FTC133 xenograft mice treated with combined CQ + sorafenib, the dephosphorylation effect was enhanced compared with that achieved by either drug alone (Fig. 3A-D).

Beclin-1 is a key autophagy-related gene that takes part in the initial process of autophagy (24). Beclin-1 protein expression was increased in xenograft mice treated with CQ or sorafenib, while combined CQ + sorafenib treatment increased the expression of Beclin-1 protein more than CQ or sorafenib treatment alone (Fig. 3A-D).

**Inhibition of the AKT/mTOR pathway activates autophagy and apoptosis, and enhances sorafenib-induced autophagy and apoptosis induction.** The MAPK pathway is a well-known target of sorafenib. However, the AKT/mTOR pathway was inhibited by sorafenib in the present study. The AKT/mTOR pathway is a major pathway that regulates the process of autophagy (25,26). Next, we demonstrated the role of the AKT/mTOR pathway in sorafenib-treated thyroid cell lines. To construct an AKT gene-silencing cell model, we transfected siRNAs into 8505C and FTC133 cell lines. The silencing effect was examined by western blotting. The protein levels of p-AKT were reduced significantly in AKT siRNA-transfected cell line models (27). LC3II and p62 proteins were examined again for the evaluation of autophagy levels. Following silencing of AKT by siRNA, LC3II increased and p62 decreased compared with the control group (Fig. 4A-C). AKT silencing in combination with sorafenib treatment significantly enhanced LC3II expression levels and reduced p62 compared with AKT silencing or sorafenib treatment alone (Fig. 4A-C). BEZ-235, a dual, ATP-competitive PI3K and mTOR inhibitor, was selected to demonstrate the role of the AKT/mTOR pathway in sorafenib-induced autophagy. In a previous study, 0.8 µM BEZ-235 inhibited AKT/mTOR pathway activity and inhibited 8505C and FTC133 cell line proliferation (27). Thus, 8505C and FTC133 cell lines were treated with 0.8 µM BEZ-235 for 24 h, and western blotting was used to evaluate the expression levels of LC3II and p62. LC3II expression increased and p62 expression decreased compared with the control group in both 8505C and FTC133 cell lines (data not shown). Combined treatment with sorafenib and BEZ-235 significantly enhanced LC3II expression and
reduced p62 expression (data not shown). The AKT/mTOR pathway negatively regulates autophagy and apoptosis (17). In the present study, 8505C and FTC133 cell lines treated with AKT siRNA exhibited increased cleaved capase-3 expression, and combined treatment of AKT siRNA with sorafenib enhanced cleaved capase-3 expression compared with either treatment alone (Fig. 4D and E).

Effect of autophagy on sorafenib treated thyroid cancer cell lines. In our results, sorafenib treatment or AKT/mTOR pathway suppression induced autophagy in thyroid cancer cell lines. As autophagy plays a dual role in cancer treatment, we examined the role of autophagy in sorafenib treatment of thyroid cancer (11,12). ATG5 is a key gene in the process of autophagy that can be silenced to demonstrate the role of autophagy in different treatments (6). Three different siRNAs (ATG5Ri) were transfected into 8505C and FTC133 cell lines, and RT-qPCR was used to evaluate their silencing effect. ATG5 siRNA2 appeared to have strong silencing effects and was selected for use in further experiments (data not shown).

Treatment with sorafenib or ATG5Ri increased the apoptosis of 8505C and FTC133 cell lines compared with the control group (apoptosis rate, 4.9 and 5.6%; the apoptosis rates after sorafenib or ATG5Ri treatment were 24.7 and 25.2%, respectively, in 8505C cells (Fig. 5A and B), and 53.2 and 27.3%, respectively, in FTC133 cells (Fig. 5C and D). Combined treatment with sorafenib and ATG5Ri increased apoptosis rates more than either treatment alone in both cell lines (42.8% in 8505C cells; 69.7% in FTC133 cells; Fig. 5).

Silencing of ATG5 significantly increased the apoptosis rate after combined treatment of sorafenib and BEZ-235 in FTC133 cells (72.6 vs. 68.1%; Fig. 5). Silencing of ATG5 increased the anti-proliferative effect of sorafenib and combined treatment of sorafenib and BEZ-235 in both 8505C and FTC133 cell lines (Fig. 6).
Resistance to $^{131}$iodine therapy is an important factor leading to poor prognosis for DTC patients (2,3). Several targeted kinase inhibitors have been used in clinical trials and administered to patients who have advanced/progressive $^{131}$iodine-refractory DTC disease (28). Sorafenib targets the RAF serine/threonine kinases, and also potently inhibits tyrosine kinase receptors, such as vascular endothelial growth factor receptor and platelet-derived growth factor receptor-$\beta$, which promote angiogenesis (7,8). It has been reported that sorafenib has therapeutic effects in several types of cancer, such as HCC, acute myeloid leukemia, advanced renal cell carcinoma or prostate cancer (29-32). Sorafenib is the first tyrosine kinase inhibitor (TKI) approved for advanced/progressive $^{131}$iodine-refractory DTC patients in China. After sorafenib treatment, the majority of advanced/progressive $^{131}$iodine-refractory DTC patients exhibit a favorable prognosis. However, most patients acquire resistance to sorafenib after 1 or 2 years of administration, and the mechanism of this acquired resistance remains unclear (10). In the treatment of various cancers, including liver, kidney and prostate cancer, sorafenib can achieve a few months of progression-free survival, but drug resistance frequently emerges (31-33). Mechanisms of sorafenib resistance may be related to EGFR activation, downstream signaling of EGFR (particularly Ras/Raf/MEK/ERK), the PI3K/AKT pathway, autophagy or epithelial-mesenchymal transition, as reviewed by Beardsley et al (32), and Chen et al (33).

The process of autophagy could be activated during cancer treatment by radiation (34), chemotherapy (35) or TKI drugs (12,15). The role of autophagy in cancer treatment is complex since it is a ‘double-edged sword’: autophagy may promote or inhibit cell death (12,36,37). The varying roles of autophagy in cancer treatment primarily depend on different contexts and treatments (13,37,38). In different HCC cell lines, the role of autophagy induced by sorafenib was reversed, as detailed by Prieto-Dominguez et al (13). It has been reported that inhibition of AKT reverses acquired resistance to sorafenib by switching protective autophagy to autophagic cell death (17).

It has been demonstrated that sorafenib triggers autophagy in different cancer contexts (12,15). In the present study, sorafenib induced autophagy in thyroid cancer cell lines and mice xenograft models. Inhibition of autophagy by either a pharmacological inhibitor (CQ) or siRNA knockdown of essential autophagy genes increased the effect of sorafenib treatment. CQ has been safely used for decades in patients for malaria prophylaxis and for the treatment of rheumatoid arthritis (39). The anticancer role of CQ has primarily been attributed to its autophagy regulation role. Autophagy is an evolutionarily conserved cellular degradation process that engulfs cytoplasmic materials in double-membrane structures (autophagosomes), which are subsequently transported to lysosomes for degradation and recycling (11,40). CQ is an autophagy inhibitor that prevents autophagosomes from fusing with lysosomes, thereby disrupting the process of autophagy (20). The primary role of CQ in cancer treatment is to enhance the efficacy of a drug or other treatment measure (41). However, CQ does not markedly alter anticancer efficiency in certain types of cancer; it may depend on the tumor type and context (42). To the best of our knowledge, no previous studies have demonstrated the role of CQ in sorafenib treatment of thyroid cancer. In the present study, thyroid cancer xenograft mice were administered CQ. The results indicated that the processes of autophagy were inhibited by CQ, and that the anticancer efficiency of sorafenib was significantly increased. Recent studies have indicated that the synergistic effects of CQ in cancer treatment are independent of autophagy in several types of cancer (43-45). The mechanism of the synergistic effects of CQ is consistent with lysosomal cell death (43), vessel normalization (44) and other mechanisms (45). In order to confirm the role of autophagy in sorafenib treatment of thyroid cancer, further regulatory pathways and autophagy genes were evaluated.
Activation of the PI3K/AKT signaling pathway mediates acquired resistance to sorafenib in HCC (46). Sorafenib has been reported to inhibit both the MAPK and AKT/mTOR pathways in lymphoma xenografts (47). Similarly, in the present study, sorafenib inhibited both the MAPK and AKT/mTOR pathways. mTOR is a negative regulator of autophagy and promotes the proliferation of cancer cells (18). In the present study, the AKT/mTOR pathway was selected to demonstrate the roles of autophagy in thyroid cancer treatment by sorafenib. We silenced the AKT/mTOR pathway by siRNA technology or with an inhibitor (BEZ-235 is a dual inhibitor of PI3K and AKT). Subsequent results indicated that the downstream gene activity of the AKT/mTOR pathway was suppressed, cell proliferation was significantly reduced, and apoptosis and autophagy were enhanced. These results indicated that sorafenib induced autophagy, at least in part due to inhibition of the AKT/mTOR pathway.

Beclin-1 is a key gene participating in the initial process of autophagy (48). The expression of Beclin-1 was promoted after sorafenib treatment in the present study. This finding indicates that a complex mechanism participates in sorafenib-induced autophagy. However, due to the dual role of autophagy in cancer treatment, it is not clear whether sorafenib-induced autophagy or AKT/mTOR pathway inhibition promote or inhibit cell death (37).

ATG5 is an E3 ubiquitin ligase that is critical for autophagy due to its role in autophagosome elongation (6). Silencing of ATG5 demonstrates the role of autophagy in a cancer treatment process (6). Treatment with ATG5 siRNA in the present study induced apoptosis in 8505C and FTC133 cell lines. Combined treatment with an ATG5 siRNA increased the tumor-suppressive effect of sorafenib in 8505C and FTC133 cell lines. The apoptosis rate of the ATG5 silencing + sorafenib + BEZ235 treatment group increased more than that of the sorafenib + BEZ235 treatment group in FTC133 cells. We did not observe a significantly different apoptosis rate between the ATG5 silencing + sorafenib + BEZ-235 treatment group and the sorafenib + BEZ-235 treatment group in 8505C cells. In our data, silenced ATG5 suppressed the proliferation of 8505C and FTC133 cells, and ATG5 silencing combined with sorafenib treatment significantly suppressed cell proliferation in 8505C and FTC133 cells. Our previous study indicated that silencing the AKT/mTOR pathway induced apoptosis, suppressed proliferation and induced autophagy in both 8505C and FTC133 cells (27). In the present study, sorafenib inhibited both the MAPK and AKT/mTOR pathways in 8505C and FTC133 cells. Thus, AKT/mTOR inhibition by sorafenib represents a positive effect of sorafenib treatment in thyroid cancer. However, autophagy induced by sorafenib involves a complex mechanism and can only be partially attributed to AKT/mTOR suppression. Comprehensive analysis of our results revealed that autophagy is a protective mechanism in sorafenib treatment of thyroid cancer. Our study had limitations. The additive or synergistic effect of sorafenib and CQ in mouse xenograft models can not be easily established and can be examined by future studies.

In summary, the present study demonstrated that sorafenib induces autophagy in thyroid cancer, and that this effect is, at least in part, due to AKT/mTOR pathway suppression. We demonstrated that CQ increased the efficacy of sorafenib in treating thyroid cancer in xenograft mice, resulting in decreased tumor growth. Treatment with CQ or ATG5 silencers blocked the processes of autophagy and increased apoptosis in sorafenib-treated tumors. Autophagy has a cytoprotective role in sorafenib treatment of thyroid cancer. Combined treatment of sorafenib with CQ or other autophagy inhibitors is a novel and potentially useful clinical strategy to improve the efficacy of sorafenib-targeted thyroid cancer therapies. Our findings provide a basis for investigating the use of sorafenib with autophagy inhibitors to treat iodine-resistant, advanced DTC.

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References


