Chloroquine affects autophagy to achieve an anticancer effect in EC109 esophageal carcinoma cells in vitro

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Abstract. Esophageal carcinoma is a malignancy that severely threatens human health, with a high incidence rate and a low 5-year survival rate. Resistance to chemotherapy frequently emerges during its treatment, partly due to the induction of autophagy. Therefore, targeting autophagy may be a promising therapeutic approach for the treatment of esophageal carcinoma. In the present study, it was investigated how chloroquine (CQ) can influence the growth ability and biological behaviors of EC109 esophageal squamous carcinoma cells in vitro, as well as the potential molecular mechanisms behind its activity. It was demonstrated that CQ could suppress the growth and proliferation of EC109 cells in a time- and dose-dependent manner; migration and colony formation abilities were also inhibited by CQ. Furthermore, subsequent to the exposure to CQ, the number of autophagosomes was clearly increased in EC109 cells over-expressing green fluorescent protein tagged-light chain (LC)3 when observed by fluorescence microscopy. Protein expression of the endogenous autophagy markers LC3-II and p62 was elevated subsequent to CQ treatment, whereas the expression of proteins from the protein kinase B/mechanistic target of rapamycin target of rapamycin pathway was inhibited. This suggested that CQ could induce the formation of autophagosomes in the initiation of autophagy, but inhibit the degradation of autophagosomes in a later stage of autophagy. The overall effect was that autophagic cell death was activated by CQ, as confirmed by flow cytometry. Overall, the anticancer effect of chloroquine on EC109 was revealed to be mediated through modulating autophagy, and this may produce promising therapeutic benefits for esophageal carcinoma.

Introduction

Esophageal cancer is the eighth most common cancer worldwide, with 456,000 new cases and ~0.4 million mortalities reported in 2012. Due to the geographical variation in the occurrence of esophageal cancer, 49% of new cases occurred in China (1), particularly in two counties renowned for the high occurrence rates of esophageal cancer: Linxian, Henan and Yanting, Sichuan (2). This is due to the tumors being asymptomatic in early stages, meaning >50% of patients present at an advanced stage that is too late for treatment by esophagectomy.

Chemotherapy is the main treatment for advanced esophageal cancer (3). However, chemotherapy is associated with drug resistance and side effects. Therefore, the identification of novel and safe treatments for esophageal cancer is necessary.

Autophagy is an intricate and conserved process. Autophagic flux refers to the process by which damaged organelles and unfolded proteins are sequestered into autophagosomes within the cytoplasm, which fuse with lysosomes to form autolysosomes, to induce the degradation of intracellular components (4). The induction of autophagy has been identified as a drug resistance mechanism that promotes cancer cell survival via self-digestion (5,6). Therefore, targeting autophagy may be a promising approach for cancer therapy.

Chloroquine (CQ) has been safely and widely used as an anti-malarial for >60 years (7). A number of studies have suggested that CQ can inhibit autophagy through the prevention of lysosome acidification, and subsequently, the inhibition of autophagosome-lysosome fusion, to block the degradation of autolysosomes at the last step of autophagy (8,9). Previous studies have indicated that CQ exhibited an antitumor effect on several types of cancer, including glioblastoma (10,11), hepatocellular carcinoma (12), breast cancer (13), prostate cancer (14) and pancreatic cancer (15), which prompts us to hypothesize that CQ may be able to influence the growth of esophageal carcinoma through the modulation of autophagy. Therefore, the aim of the present study is to explore the antitumor effect of CQ on the esophageal squamous carcinoma cell line EC109, and the potential mechanism for this effect.

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Materials and methods

Cells and reagents. EC109 human esophageal squamous cell carcinoma cells were obtained from the Institute of Rheumatism Immunity, Affiliated Hospital of North Sichuan Medical College (Nanchong, China). EC109 cells were grown in RPMI-1640 supplemented with 10% fetal bovine serum (both from Hyclone; GE Healthcare Life Sciences, Logan, UT, USA), penicillin (100 U/ml) and streptomycin (100 µg/ml; Sigma-Aldrich; Merck KGaA, Darmstadt, Germany). Cells were incubated in a humidified incubator in 5% CO₂ at 37°C.

CQ was kindly provided by Dr Yibin Deng (Hormel Institute, University of Minnesota, Austin, MN, USA). RPMI-1640 was purchased from Hyclone; GE Healthcare Life Sciences. All primary antibodies, including light chain (LC3) (cat. no. 4108), p62/SQSTM1 (cat. no. 5114), phosphorylated (p)-mechanistic target of rapamycin (mTOR; cat. no. 2971), p-protein kinase B (Akt; cat. no. 9271), p-70S6K (cat. no. 9205), p-4E binding protein 1 (4EBP1; cat. no. 9451) and GAPDH (cat. no. 2118) were obtained from CST Biological Reagents Co., Ltd. (Shanghai, China).

*MTT viability assay.* Cells were seeded at a density of 2x10⁴ per well in a 96-well plate and incubated for 24 h. CQ was added to each well in a concentration series of 50, 100, 150 and 200 µmol/l and incubated for a further 12, 24 and 36 h. A total of 20 µl MTT (5 mg/ml; Sigma-Aldrich) was added to each well and incubated at 37°C for 4 h, then 100 µl DMSO was added to each well to dissolve crystals and the plate was agitated for 10 min. Absorbance values at 490 nm were detected using a multi-mode detection platform (Molecular Devices Austria GmbH, Wals, Austria). Inhibition rate was calculated as 
\[
\frac{(A_{\text{control}} - A_{\text{blank}}) - (A_{\text{treated}} - A_{\text{blank}})}{(A_{\text{control}} - A_{\text{blank}})} \times 100
\]
Each experiment was assayed in triplicate.

*Scratch assay.* Cells were seeded in a 6-well plate at a density of 5x10⁴ cells per well and cultured until the cells formed a confluent monolayer. A 10 µl pipette tip was used to scratch the cells and create a wound. The medium was removed and the cultures were washed twice with PBS to remove non-adherent cells. The cells were then incubated with complete RPMI medium, with and without 200 µmol/l CQ. The migratory ability of the cells was determined according to the extent of closure of the scratch, observed at 0 and 24 h after incubation with CQ. Images were acquired using an inverted microscope (DMI 3000B; Leica Microsystems GmbH, Wetzlar, Germany).

*Transwell migration assay.* Cells (6x10⁴) in RPMI 1640 with 5% FBS were seeded into the upper chamber of polycarbonate membrane Transwell inserts (Corning Incorporated, Corning, NY, USA) with or without 200 µmol/l CQ. RPMI-1640 supplemented with 20% FBS was added to the lower chamber as a chemo-attractant solution. Following a 48-h incubation, the remaining cells on the upper chamber surface were removed, and the cells that had migrated to the underside of the chamber were fixed in methanol and stained with 10% Giemsa for 10 min at room temperature. The cell migration ability was determined by the number of cells that had migrated to the lower side of the filter. These experiments were repeated three times, and the filters were photographed under an inverted microscope.

*Green fluorescent protein (GFP)-LC3 transfection.* EC109 cells were transfected with a GFP-LC3 plasmid using Lipofectamine™ 3000 (Invitrogen; Thermo Fisher Scientific, Inc., Waltham, MA, USA) and incubated for 24 h and treated with and without CQ. The images were obtained by fluorescence microscope.

*Western blot analysis.* Cells treated with or without 200 µmol/l CQ were lysed in radioimmunoprecipitation assay buffer (Beyotime Institute of Biotechnology, Haimen, China) containing proteinase inhibitors, and lysates were collected. The protein samples were detected using SDS-PAGE (8-15% gel). A total of 40 µg of protein was loaded into each lane of the SDS-PAGE gel and transferred onto a polyvinyl difluoride membrane. Protein was quantified using a BCA assay. Membranes were blocked in 5% fat-free milk for 1 h at room temperature, and probed with the previously specified primary antibodies (dilution 1:1,000) overnight at 4°C, followed by horseradish peroxidase-conjugated secondary antibodies (1:5,000; Boster Biological Technology, Pleasanton, CA, USA) and incubated for 1 h at room temperature. Bands were visualized using an enhanced chemiluminescence kit (Merck KGaA).

*Cell death assessment by flow cytometry.* Cells were seeded at 5x10⁴ cells per well in a 6-well plate and incubated with or without 200 µmol/l CQ for 24 h. Cells were harvested by trypsinization and washed with PBS. Subsequent to staining with Annexin V-fluorescein isothiocyanate for 15 min at 4°C and propidium iodide for 2 min at room temperature (Nanjing KeyGen BioTech, Co., Ltd., Nanjing, China), cells were detected using flow cytometry using FACSComp (version 4.1; Beckman Coulter, Inc., Brea, CA, USA).

*Statistical analysis.* The results are expressed as the mean ± standard deviation and t-tests or one-way analysis of variance followed by a least significant difference t-test were performed using SPSS (version 13.0; SPSS, Inc., Chicago, IL, USA) statistical software. P<0.05 was considered to indicate a statistically significant difference.

**Results**

Proliferation, migration and colony formation abilities of EC109 are inhibited by CQ. An increasing number of studies have described the anti-tumor effect of CQ. Thus, the first aim of the present study was to explore whether CQ affected the growth, migration or colony-forming ability of EC109 cells. According to the results of the MTT assay, CQ suppressed the viability of EC109 cells in a time and dose-dependent manner (Fig. 1A). The scratch assay indicated that the migration
ability of EC109 cells was inhibited after a 24 h CQ treatment (Fig. 1B). A Transwell migration assay was performed for the same purpose, and a similar result was obtained (Fig. 1C). The colony formation ability of EC109 cells was evaluated using a soft agar assay; it was revealed that the number and size of colonies were reduced in the CQ group compared with the control group (Fig. 1D).

Cell death is induced by CQ in EC109 cells. The results of flow cytometry demonstrated that the cell death ratio in the CQ group was significantly increased compared with the control group (P<0.05; Fig. 2).

CQ modulates autophagy through inducing the formation of autophagosomes and preventing the degradation of autophagosomes in EC109 cells. A number of studies have indicated that the induction of autophagy promoted tumor cell survival, whereas others have achieved the opposite result (16). Therefore, the present study aimed to explore whether CQ could modulate autophagy and influence the survival of EC109 cells.

The GFP-LC3 fused protein translocates rapidly to nascent autophagosomes in a punctate distribution during the initiation of autophagy. Subsequent to the transfection of a GFP-LC3 plasmid into EC109 cells to monitor autophagosome formation by fluorescence microscopy, it was observed that the number of autophagosomes was evidently increased following CQ treatment (Fig. 3A). LC3 has two isoforms, LC3-I and LC3-II; LC3-I is cleaved by autophagy associated protein ATG4 and lipidated to form LC3-II during the initiation of autophagy. Thus, the expression of LC3-II is often used as a marker to evaluate the progression of autophagy (17). Western blot analysis revealed that CQ treatment markedly enhanced LC3-II expression (Fig. 3B), which indicated that CQ could induce the formation of autophagosomes in EC109 cells; this result corroborated the conclusion drawn from the GFP-LC3 transfection experiment.

Another autophagy marker, p62, is sequestered within autophagosomes and then degraded by lysosomes during the last stage of autophagy. The present study demonstrates that
the expression of p62 was upregulated following CQ treat-
ment compared with the control (Fig. 3B). Collectively, these
results demonstrate that CQ inhibited autophagy by preventing
autophagosome degradation at the last stage of autophagy,
despite inducing autophagosome formation during the initia-
tion of autophagy.

Based on the results outlined in the previous section, it
is hypothesized that CQ could suppress the growth and
proliferation of EC109 cells in a time- and dose-dependent
manner and reduce their migration and colony formation
ability by preventing autophagosome degradation. In order
to elucidate the mechanism for this process as induced by CQ,
the expression of key proteins in the Akt/mTOR signaling
pathway, which serves an important role in promoting cell
growth and proliferation and is associated with the regulation
of autophagy, was assessed. Western blotting demonstrated
that the expression of autophagic negative regulatory factors,
p-Akt and p-mTOR, decreased on CQ treatment. This
suggests that autophagosome formation was initiated. The
phosphorylation levels of the downstream proteins, includ-
ing the cell translation regulation factors, p70S6K and 4EBP1,
also decreased with CQ treatment (Fig. 3C), and therefore,
CQ may initiate the formation of autophagosomes, but inhibit
the degradation of the autophagosome in the final stage of
autophagy.

Discussion

Esophageal carcinoma is one of the most common types of
malignant neoplasm and a leading cause for cancer-associated
mortality worldwide (1). Radiotherapy and chemotherapy
are the most common treatments for patients with advanced
esophageal carcinoma, with cisplatin, fluorouracil, oxali-
platin and capcitabine being the most commonly used
chemotherapy agents (18). These drugs gradually induce
chemoresistance and side effects due to the higher and longer
dosages required. Developing an effective drug may cost $800
million and 15 years (19), therefore, a more effective way to
solve the problem may be to develop new chemotherapeutics
from existing drugs.

Over the past ten years, autophagy inhibition has been
a popular topic in the field of cancer therapy. Autophagy is
an important regulatory mechanism involved in cell growth,
maturity and death, functions which are associated
with a number of diseases, including cancer. Autophagic
activity in the context of cancer is a double-edged sword:
It degrades damaged organelles and recycles macromol-
ecules to maintain a stable cellular environment, preventing
tumor formation (20), but autophagy may also contribute
to tumorigenesis by helping cancer cells to survive in
response to growth-limiting conditions, including nutrient
deficiency, absence of growth factors, hypoxia and exposure
to cytotoxic drugs (6). A recent study revealed that in an
increasing proportion of cases, chemotherapy failure is
due to drug-initiated autophagy, which eventually induces
chemoresistance (21).

As an inhibitor of autophagy, a number of studies have
suggested CQ as a promising approach for cancer therapy
as it is relatively safe and inexpensive (7,22,23). However,
the specific anti-tumor mechanisms of CQ remain unchar-
acterized. In order to explore whether CQ could be used
as a potential chemotherapeutic agent for the treatment of
esophageal carcinoma, it was used to treat EC109 esophageal
squamous cell carcinoma cells. The MTT assay results indi-
cated that CQ was able to suppress the proliferation of EC109
cells in a time and dose-dependent manner, and the migra-
tion ability of the cells was also inhibited, as demonstrated
using scratch and Transwell chamber tests. Furthermore, CQ
suppressed colony formation on soft agar, which indicated
that the transformation ability of EC109 cells was impaired.
Collectively, these findings demonstrated that CQ may have an antitumor effect on EC109 cells.

In order to determine whether autophagy was involved in the antitumor activity of CQ, autophagosome formation was monitored by transecting EC109 cells with a GFP-LC3 plasmid, and observing the expression of LC3 by fluorescence, and LC3-II and p62 by western blot analysis. As the substrates of autophagic flux, LC3 and p62 are involved in the process of autophagosome-lysosome fusion, which can be used to evaluate the formation and degradation of the autophagosome (24). Results demonstrate that CQ treatment increased the expression of LC3 and p62, indicating that autophagosome formation was increased, and autophagosome degradation was markedly inhibited, subsequent to CQ treatment.

To investigate the signaling mechanism of CQ-mediated autophagy in EC109 cells, the Akt/mTOR pathway, which serves a central role in regulating cell growth, was studied. The over-activation of this pathway contributes to cancer development and progression; Akt positively regulates the phosphorylation and activation of mTOR, which is a sensor of nutrient levels to promote cell growth (25). Furthermore, the Akt/mTOR pathway has been demonstrated to be involved in the regulation of autophagy; the inhibition of mTOR induces autophagy via the phosphorylation and inactivation of Unc-51-like autophagy activating kinase 1, a serine/threonine kinase that initiates autophagy (26). mTOR exists in two cellular complexes, mTORC1 and mTORC2 (27). mTORC1 phosphorylates S6K1 and 4E-BP1, which contribute to translation, protein synthesis and cell growth. mTORC2 regulates the phosphorylation of Akt. In the present study, the expression of p-mTOR, p-Akt, p-p70S6K1 and p-4EBP1 were markedly downregulated following CQ treatment.

The increased LC3-II expression and the inhibition of the Akt/mTOR pathway indicated that CQ induced autophagic flux and facilitated the formation of autophagosomes. However, the upregulation of p62 indicated that CQ also inhibited the degradation of the autophagosomal contents. Taken together, these results demonstrate that, as an inhibitor of autophagy, CQ induced the formation of the autophagosome, but inhibited the degradation of autophagosome in the last stage of autophagy. The overall effect was autophagic cell death activation by CQ, as confirmed by flow cytometry. Liu et al came to a similar conclusion regarding cervical cancer by incubating SiHa cells with hydroxychloroquine, the hydroxylated analog of CQ (28); another study demonstrated that autophagy of SiHa cells via inhibition of CXCR4 and hedgehog signaling. Mol Cancer Ther 35: 322-332, 2016.

Based on the data of the present study, CQ exerted an antitumor effect on EC109 through the modulation of autophagy, suggesting that CQ may exhibit promising therapeutic benefits for the treatment of esophageal squamous cell carcinoma.

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References