EHHM, a novel phenolic natural product from Livistona chinensis, induces autophagy-related apoptosis in hepatocellular carcinoma cells

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Abstract. Hepatocellular carcinoma (HCC) ranks the second cause of cancer-associated mortality worldwide. In the present study, the effects and mechanisms of a new phenolic natural product E-[6'-(5'-hydroxypentyl) tricosyl]-4-hydroxy-3-methoxycinnamate (EHHM) isolated from Livistona chinensis on the growth of HCC cells were investigated. It was observed that EHHM treatment significantly suppressed cell proliferation and colony formation, and induced cell apoptosis via a mitochondria-dependent caspase pathway in HepG2 cells in a time- and dose-dependent manner. Meanwhile, EHHM treatment also led to upregulated expression of autophagy protein 5 (Atg5), Beclin 1 and light chain 3 (LC3)-II proteins, and accumulation of green fluorescent protein-LC3 punctate florescent foci in HCC cells, suggesting that EHHM-induced apoptosis is accompanied by autophagy induction. Western blotting revealed that EHHM-induced autophagy is related to the inhibition of the Akt/mechanistic target of rapamycin/p70 ribosomal protein S6 kinase signaling pathway. Furthermore, treatment with Atg5 small interfering RNA or autophagy inhibitors significantly enhanced EHHM-mediated growth inhibition and apoptotic cell death, indicating that autophagy serves as a self-protective mechanism in EHHM-treated HCC cells, and that combined

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treatment with EHHM and autophagy inhibitors may be an effective therapeutic strategy for HCC.

Introduction

Hepatocellular carcinoma (HCC) is the most frequently diagnosed cancer and the second major cause of cancer mortality worldwide (1-3). Infection with hepatitis B or C virus is the leading cause of HCC (4). Surgical resection is mostly selected in HCC treatment, followed by radiation therapy and chemotherapy; however, the 5-year survival rates following resection are very low, and the recurrence rates, mainly due to primary tumor metastasis and formation of new tumors, exceed 70% (5,6). Therefore, it is necessary and urgent to develop new effective chemotherapeutic agents for the treatment of HCC.

Livistona chinensis, a species of subtropical palm tree in eastern Asia, is widely distributed and cultivated in tropical and temperate regions, including several islands in the South China Sea, Southern Japan and Taiwan (7). The extract from the fruit of L. chinensis has been shown to exhibit antibacterial activity (8), and was traditionally used for folk remedy for various tumors, including chorionic carcinoma, esophageal carcinoma, leukemia and nasopharyngeal carcinoma (9,10). E-[6'-(5'-hydroxypentyl) tricosyl]-4-hydroxy-3-methoxycinnamate (EHHM), a new phenolic compound isolated from the L. chinensis fruit, has been shown to have antioxidant activity and a potent anticancer activity by inhibiting cancer cell proliferation (9,11). However, the antitumor effects and possible mechanisms of EHHM against human HCC have not yet been investigated. In the present study, it was demonstrated that EHHM can inhibit the proliferation and induce the apoptosis of HCC cells. In addition, EHHM induced autophagy in HCC cells.

The induction of cancer cell apoptosis is the main consideration in anticancer compound research (12-15). Cell morphology shrinkage, nuclear fragmentation, chromatin condensation and chromosomal DNA fragmentation are typically observed during apoptosis (16). However, to escape from apoptosis, cancer cells may also trigger various other pathways

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such as autophagy in order to overcome the adverse environment (17-19).

Autophagy is an intracellular degradation process, through which, a portion of the cytoplasmic components is delivered to lysosomes to be degraded (20). The cell fates are different when autophagy is induced in response to multiple environmental stresses (20). Autophagy can induce cell death (21) or serve as a survival mechanism that protects tumor cells from apoptotic death (22,23). Meanwhile, several studies have suggested that, in certain circumstances, inhibition of protective autophagy can promote cell apoptosis (24). Therefore, autophagy may be a novel approach for cancer treatment. In the present study, it was observed that EHHM induced cytoprotective autophagy through the Akt/mechanistic target of rapamycin (mTOR) signaling pathway in HCC cells, and that inhibition of autophagy by small interfering RNA (siRNA) targeting autophagy-related genes or autophagy inhibitors could lead to the enhancement of EHHM-mediated cytotoxicity through increasing cell apoptosis in vitro and in vivo.

Materials and methods

Reagents. EHHM was isolated and identified as described in a previous study (9). EHHM was dissolved in dimethyl sulfoxide (DMSO) at a stock solution of 50 mM, stored at -20°C and diluted with culture medium prior to each experiment. The human HCC cell line HepG2, which was obtained from the American Type Culture Collection (Manassas, VA, USA), was routinely maintained in Dulbecco's modified Eagle medium (DMEM; Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) with 10% fetal calf serum (Gibco; Thermo Fisher Scientific, Inc.) and incubated in a humidified atmosphere with 5% CO₂ at 37°C. 3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) was purchased from Roche Diagnostics (Indianapolis, IN, USA). Trypan blue was purchased from Sangon Biotech Co., Ltd. (Shanghai, China). The pEGFPc1-LC3 plasmid was a gift from Professor Marja Jäättelä (Institute of Cancer Biology, Danish Cancer Society, Copenhagen, Denmark) (25). Benzyloxycarbonyl-Val-Ala-Asp (OMe) fluoromethylketone (Z-VAD-FMK) was purchased from Sigma-Aldrich (Merck Millipore, Darmstadt, Germany). Lipofectamine 2000 and FITC Annexin V/Dead Cell Apoptosis kit were purchased from Invitrogen (Thermo Fisher Scientific, Inc.). JC-1 Mitochondrial Membrane Potential Assay kit was purchased from Nanjing KeyGen Biotech Co., Ltd. (Nanjing, China). 4',6-Diamidino-2-phenylindole and Hoechst 33258 were obtained from Sangon Biotech Co., Ltd. Bafilomycin A1 (Baf A1) and 3-methyl adenine (3-MA) were purchased from Sigma-Aldrich (Merck Millipore). ECL Western Blotting Substrate kit was purchased from Abcam (Cambridge, MA, USA). Antibodies were purchased from the following suppliers: Anti-\beta-actin antibody (cat. no. AF0003) from Beyotime Institute of Biotechnology (Haimen, China); anti-caspase 3 antibodies (cat. no. ab13585) from Abcam; and anti-light chain 3 (LC3; cat. no. 3868), anti-autophagy protein 5 (Atg5; cat. no. 12994) and anti-Beclin 1 (cat. no. 3495) antibodies from Cell Signaling Technology, Inc. (Danvers, MA, USA).

Cell proliferation and viability assays. Cell proliferation was determined by MTT assay as described previously (26).

Briefly, 90 μ l cells (10⁵ cells) were seeded into a 96-well plate and precultured for 24 h, prior to be treated with EHHM for 24 or 48 h. Next, 10 μ l of MTT (5 mg/ml) was added to each well and incubated for additional 4 h at 37°C. The medium was removed carefully, and 150 μ l of DMSO was added into the plate, and then the 96-well plate was agitated for 10 min. The absorbance (A) was measured at 570 nm by an automated microplate reader (BioTek Instruments, Inc., Winooski, VT, USA). The cell death rate was calculated as follows: Cell death (%) = (mean A₅₇₀ of the control group - mean A₅₇₀ of the experimental group)/(mean A₅₇₀ of the control group - mean A₅₇₀ of the blank group) x 100%. Cell viability was estimated by trypan blue dye exclusion (27).

Soft agar colony formation assay. Colony formation in soft agar was tested by suspending 1x10³ cells in 1 ml of DMEM with 0.3% low melting point agarose (Merck Millipore) in the presence of different concentrations of EHHM, and then plating them on a bottom layer containing 0.6% agarose and 10% fetal bovine serum (Gibco; Thermo Fisher Scientific, Inc.) in a 6-well plate in triplicate. After 15 days, the plates were stained with 0.1% gentian violet, and the colonies were counted under a light microscope (28,29).

Nuclei morphology observation by Hoechst staining. Cell morphological changes in the nucleus were determined by Hoechst 33258 staining as described previously (30). Briefly, cells treated with different concentrations of EHHM were fixed with 4% polyoxymethylene for 10 min and then stained with Hoechst 33258 (1 mg/ml) at room temperature in the dark for 5 min. Morphological changes of nuclei were observed and imaged using a fluorescence microscope with an excitation wavelength of 330-380 nm. Cells with round nuclear morphology were considered normal, while those with condensed/fragmented and bright nuclei were regarded as apoptotic.

Analysis of cell proliferation and apoptosis. Cells were treated with EHHM, 3-MA or Baf A1 for the indicated concentrations and time points. Apoptotic cells were analyzed using annexin V-fluorescein isothiocyanate (FITC) /propidium iodide (PI) double staining methods as described previously (31). Briefly, HepG2 cells were treated with the indicated concentrations of EHHM in the presence or absence of 3-MA (5 mM) or Baf A1 (10 nM) for 24 h. Cells were digested into a single-cell suspension with ehylenediaminetetraacetic acid-free trypsin and stained according to the manufucturer's protocol of the FITC Annexin V/Dead Cell Apoptosis kit. The stained cells were immediately analyzed by flow cytometry. Cells undergoing early-stage apoptosis are stained with annexin V-FITC only (green fluorescence), while cells at late-stage apoptosis and necrotic cells are stained with both annexin V-FITC and PI (both green and red fluorescence) (32). In addition, to determine whether EHHM-induced apoptosis is caspase-dependent, HepG2 cells were pretreated with 20 mM Z-VAD-FMK for 1 h followed by treatment with 15 μ M EHHM for 24 h, and the inhibition rate was determined by MTT assay, as described previously.

Transfection of cells with siRNA. siRNA targeting human Atg5 (5'-CCUUUGGCCUAAGAAGAAATTdTdT-3') was

synthesized by Shanghai GenePharma Co., Ltd. (Shanghai, China), and a nonspecific siRNA served as a negative control. Cells seeded in 60-mm dishes were transfected with siRNAs for 36 h prior to treatment with EHHM.

JC-1 mitochondrial membrane potential detection. Cell mitochondrial membrane potential changes were analyzed by JC-1 staining as described previously (33). Cells were incubated with or without EHHM for 24 h, followed by incubation with JC-1 working solution, and then the cells were incubated in 5% CO₂ for 20 min. Cells was washed twice with JC-1 incubation buffer, and then the cells labeled with JC-1 were detected with a fluorescence microscope. JC-1 can selectively enter the intact mitochondria with high membrane potential and form polymers to emit red fluorescence, whereas in mitochondria with low membrane potential, it remains in the monomeric form in the cytosol and emits a green fluorescence (16).

Western blotting. Cells were treated with different concentrations of EHHM in the presence or absence of 3-MA (5 mM) or Baf A1 (10 nM) for 24 h, and then lysed in lysis buffer [50 mM Tris-HCl (pH 6.8), 2% w/v sodium dodecyl sulfate (SDS), 10% glycerol, 10 mM dithiothreitol, 1 mM phenylmethane sulfonyl fluoride and 1X protease inhibitor cocktail (Roche, Basel, Switzerland)]. Equal amounts of protein samples were quantified using the RC DC TM Protein Assay Kit I (Bio-Rad Laboratories, Hercules, CA, USA) and electrophoresed by 8-12% SDS-polyacrylamide gel electrophoresis and transferred to a polyvinylidene fluoride membrane. The membranes were blocked with a 5% non-fat milk powder solution in Tris-buffered saline with Tween 20 (TBST) buffer [50 mM Tris-HCl (pH 7.6), 150 mM NaCl and 0.05% Tween 20] for 1 h at room temperature and then incubated with the corresponding primary antibodies (1:1,000) overnight at 4°C. Upon washing with TBST, membranes were incubated with horseradish peroxidase-conjugated anti-mouse (cat. no. 31430; 1:5,000) and anti-rabbit (cat. no. 31460; 1:5,000) IgG secondary antibodies (Thermo Fisher Scientific, Inc.) for 2 h at room temperature and detected with ECL Western Blotting Substrate kit (Thermo Fisher Scientific, Inc.) (34).

Visualization of cells with green fluorescent protein (*GFP*)-*LC3 vesicles*. Cells plated in 6-well plates were transfected with the pEGFPc1-LC3 plasmid using Lipo-fectamine 2000. After 24 h of transfection, cells were treated with different concentrations of EHHM in the presence or absence of autophagy inhibitors, 3-MA (5 mM) or Baf A1 (10 nM), for 12 h. The patterns of GFP-labeled LC3 in transfected cells were examined and imaged by fluorescence microscopy.

HCC xenograft experiments. A total of 24 female 6-7 week-old nude immunodeficient mice (nu/nu) (15-20 g) were purchased from the Guangdong Province Medical Animal Center (Guangzhou, China) and bred in the laboratory animal facility of Tsinghua University (Shenzhen, China). The mice were housed at 25°C and exposed to 12 h light/dark cycles with free access to food and water. All animal studies were approved by the Institutional Animal Care and Use

Committee of Tsinghua University. The mice were subcutaneously injected with HepG2 cells ($2.0x10^6$ cells) in 100 μ l of DMEM without serum. The mice were randomly divided into four groups (6 mice/group) when the tumors reached a palpable size and then treated with vehicle control, 3-MA (24 mg/kg) and EHHM (50 mg/kg), alone or in combination with 3-MA, every 2 days. Vernier caliper measurements of the longest perpendicular tumor diameters were performed every 2 days to estimate the tumor volume with the following formula: $4\pi/3 \times (width/2)^2 \times (length/2)$, which represents the three-dimensional volume of an ellipsoidal tumor tissue (35).

Statistical analysis. Data from three independent experiments were expressed as the mean \pm standard deviation, unless otherwise noted. The Student's *t* test was used to evaluate the variables in the present study. P<0.05 was considered to indicate a statistically significant difference.

Results

Proliferation-inhibition and apoptosis-induction effect of EHHM on HepG2 cells. EHHM is a novel plant metabolite isolated from the fruit of L. chinensis (Fig. 1A). In the present study, the cytotoxicity of EHHM towards human HepG2 cells was examined. By MTT assay, it was demonstrated that EHHM had a moderate cytotoxicity towards HepG2 cells, with a half maximal inhibitory concentration of $41.62 \pm 4.57 \,\mu\text{M}$ (Fig. 1B). Meanwhile, using trypan blue dye exclusion analysis, it was demonstrated that EHHM inhibited the proliferation and growth of HepG2 cells in a time- and dose-dependent manner (Fig. 1C). Next, the effect of EHHM on colony formation was tested using soft agar colony formation assays. The results revealed that the clonogenic activity of HepG2 cells was significantly suppressed (Fig. 1D and E). These results suggested that EHHM can significantly inhibit the proliferation and growth of HCC cells in a time- and dose-dependent manner.

EHHM induces apoptosis of HCC cells via a mitochondriadependent caspase activation cascade. It was tested whether EHHM can induce the apoptosis of HCC cells. By Hoechst 33258 staining, HepG2 cells treated with the indicated concentrations of EHHM exhibited chromatin condensation and fragmentation, which are typical apoptotic nuclear morphological changes (Fig. 2A). Meanwhile, flow cytometry assay using annexin V/PI staining also confirmed that treatment with EHHM induced apoptosis in HepG2 cells in a time- and dose-dependent manner (Fig. 2B). Caspase 3, a critical executor of apoptosis, is either partially or totally responsible for the proteolytic cleavage of numerous key proteins such as the nuclear enzyme poly (ADP-ribose) polymerase (PARP) (36). The inactive full-length caspase 3 is processed during apoptosis, generating an activated 17-kDa fragment termed cleaved caspase 3 (37). In the present study, both cleaved caspase 3 and cleaved PARP were detected subsequent to EHHM treatment at different concentrations (15, 45, 90 and 150 μ M), and a reduction in pro-caspase 3 (through cleavage) was observed (Fig. 2C). To further confirm that EHHM-induced cell apoptosis is caspase-dependent, HepG2 cells were pretreated with Z-VAD-FMK (20 mM), a pan-caspase inhibitor, for 1 h and then treated with 15 μ M EHHM for additional 24 h. As



Figure 1. Inhibitory effects of EHHM on HepG2 cells. (A) Chemical structure of EHHM. (B) The inhibitory effects of EHHM on HepG2 cells were analyzed by $3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide assay. (C) The inhibitory effects of EHHM on the viability of HepG2 cells were analyzed by trypan blue exclusion assay. (D) HepG2 cells were treated with EHHM at the indicated concentrations, and colony formation was observed. (E) The colonies were counted, and the data were expressed as a percentage of the control group (<math>0-\mu$ M EHHM treatment). The experiment was conducted in triplicate, and the mean \pm standard deviation was calculated (**P<0.01). EHHM, E-[6-(5'-hydroxypentyl)tricosyl]-4-hydroxy-3-methoxycinnamate.

shown in Fig. 2D, EHHM-induced apoptosis was significantly suppressed. Since the activation of caspase 3/caspase 9 involves mitochondria permeability changes and the release of cytochrome c, the present study next examined changes in the mitochondrial transmembrane potential following EHHM treatment. As shown in Fig. 2E, in EHHM-treated HepG2 cells, JC-1 remained in the cytoplasm in its monomeric form and emitted green fluorescence, whereas in the cells without EHHM treatment, JC-1 accumulates and aggregates in the mitochondria, emitting red fluorescence. This result suggested that EHHM treatment damaged the mitochondrial transmembrane of HepG2 cells. Meanwhile, cytochrome c release from mitochondria was also investigated. As shown in Fig. 2C, a significant increase in released cytochrome c was detected at 24 h after 15- μ M EHHM treatment.

Overall, these results suggested that EHHM treatment significantly suppressed HepG2 proliferation and induced apoptosis in a time- and dose-dependent manner, and that EHHM-induced apoptosis in HepG2 cells is mediated via mitochondria-dependent caspase cascade activation.

EHHM induces Akt/mTOR signaling-related autophagy in HepG2 cells. Notably, several vacuoles were observed in EHHM-treated HepG2 cells, but not in untreated cells, by light microscopy, indicating that EHHM treatment may be associated with cell autophagy.

Autophagy, a cellular pathway involved in protein and organelle degradation, plays complicated roles in cancer

development (38). During the autophagy process, LC3-I is converted into LC3-II via a ubiquitin-like system involving Atg7 and Atg3, which promotes LC3-II relocalization to autophagic vesicles (39). The conversion of LC3-I into LC3-II and the presence of LC3-II in autophagosomes have been used as indicators of autophagy activation (40). To test whether EHHM induces autophagy formation, HepG2 cells transfected with GFP-LC3 were treated with EHHM, and autophagic vacuoles were detected. As shown in Fig. 3A, the number of autophagic vacuoles with a punctate staining pattern of GFP-LC3-II was significantly increased in EHHM-treated cells compared with untreated cells. Accordingly, the protein levels of LC3-II were also significantly increased in HepG2 cells treated with EHHM in a dose-dependent manner (Fig. 3B). In addition, several studies have demonstrated that Beclin 1 participates in the initiation of the autophagosome formation, and that Atg5, an E3 ubiquitin ligase, is important in autophagosome elongation (41,42). Therefore, the expression levels of these two proteins were also investigated in EHHM-treated cells. As shown in Fig. 3B, the expression levels of Atg5 and Beclin 1 proteins in cells exposed to EHHM were also increased in a dose-dependent manner.

The phosphatidylinositol 3-phosphate kinase (PI3K)/Akt/ p70 ribosomal protein S6 kinase (p70S6K) signaling pathway is known to regulate autophagy, growth, proliferation and survival in response to a variety of stresses in mammalian cells (43). To investigate whether the Akt/mTOR/p70S6K signaling pathway was involved in EHHM-induced apoptosis



Figure 2. EHHM induces the apoptosis of HepG2 cells via a mitochondria-dependent caspase activation cascade. (A) HepG2 cells were treated with EHHM for 24 h, and evaluated by Hoechst 33258 assay. Scale bar, 50 μ m. (B) HepG2 cells were treated with EHHM for 24 h, and the apoptotic cell death rate was analyzed by annexin V/propidium iodide staining and flow cytometry. (C) HepG2 cells were treated with EHHM (15 μ M), and western blotting was performed using the indicated antibodies. (D) HepG2 cells were pretreated with Z-VAD-FMK (20 mM) for 1 h and then treated with EHHM at 15 μ M for 24 h, and the inhibition rate was determined by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide assay (*P<0.05). (E) The mitochondrial membrane potential of HepG2 cells treated with EHHM at the indicated concentration for 24 h was determined by JC-1 staining. Scale bar, 20 μ m. Data represents the mean ± standard deviation of three independent experiments. EHHM, E-[6'-(5'-hydroxypentyl)tricosyl]-4-hydroxy-3-methoxycinnamate; FITC, fluorescein isothiocyanate; PARP, poly (ADP-ribose) polymerase; Cy, cyanine.



Figure 3. Autophagy is triggered by EHHM in HepG2 cells. (A) HepG2 cells were transfected with a GFP-LC-3-expressing plasmid for 24 h, and then treated with EHHM at the indicated concentrations for additional 24 h. The cells were then observed under a confocal microscope. Scale bar, $10 \mu m$. (B) HepG2 cells were treated with EHHM at the indicated concentrations for 24 h, and western blotting was performed with antibodies against autophagy marker proteins. (C and D) EHHM treatment inhibited the AKT/mTOR/p7086K signaling pathway in a (C) time- and (D) dose-dependent manner. EHHM, E-[6'-(5'-hydroxypentyl)tricosyl]-4-hydroxy-3-methoxycinnamate; GFP, green fluorescent protein; LC3, light chain 3; Atg5, autophagy protein 5; p-, phosphorylated; mTOR, mechanistic target of rapamycin; p7086K, p70 ribosomal protein S6 kinase.



Figure 4. Knockdown of Atg5 inhibits autophagy but enhances EHHM-induced apoptosis in HepG2 cells. (A) Western blot analysis of Atg5 expression in HepG2 cells with the indicated treatment. (B) HepG2 cells were treated with control siRNA or Atg5 siRNA, with or without EHHM (15 μ M) for 24 h. The inhibition rate was determined by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide assay. (C) Western blot analysis of LC3-I/II and caspase 3 activation of HepG2 cells with the indicated treatments. (D) The apoptosis of HepG2 cells subjected to the indicated treatments was analyzed by flow cytometry. EHHM, E-[6-(5'-hydroxypentyl)tricosyl]-4-hydroxy-3-methoxycinnamate; LC3, light chain 3; Atg5, autophagy protein 5; siRNA, small interfering RNA; FITC, fluorescein isothiocyanate; PI, propidium iodide.

and autophagy, western blotting was performed to detect the expression levels of the associated proteins. As shown in Fig. 3C and D, EHHM inhibited the phosphorylation of Akt, mTOR and p70S6K in a time- and dose-dependent manner.

Taken together, these data indicated that EHHM was able to induce autophagy in HepG2 cells, which involves the Akt/mTOR/p70S6K signaling pathway.

Autophagy inhibition enhances the cell growth-inhibition and apoptotic effect of EHHM. As demonstrated above, EHHM treatment induces cell apoptosis and autophagy in HepG2 cells. As previously reported, autophagy may serve different roles in response to different cellular stresses, protecting or killing the cells (44). To identify which type of autophagy is induced by EHHM, HepG2 cells were transfected with Atg5 siRNA (Fig. 4A), and the proliferation effect of EHHM on HepG2 cells was evaluated using an MTT assay. As illustrated in Fig. 4B, knockdown of Atg5 expression significantly enhanced the inhibition effect of EHHM on HepG2 cells compared with control siRNA. Simultaneously, knockdown of Atg5 resulted in the decrease of LC3-I/II conversion and the increase in cleaved caspase 3 expression (Fig. 4C). A flow cytometry assay further confirmed the above results. As shown in Fig. 4D, knockdown of Atg5 significantly enhanced EHHM-induced apoptosis of HepG2 cells.

Furthermore, the effects of autophagy specific inhibitors (3-MA and Baf A1) on EHHM-induced growth inhibition and apoptosis were investigated. 3-MA inhibits autophagy via blocking the autophagosome formation in the early stage of autophagosome formation (45), whereas Baf A1, a macro-lide antibiotic derived from *Streptomyces griseus*, inhibits autophagy by blocking the fusion of autophagosomes with lysosome in the late stage of autophagy (46). As shown in Fig. 5A, the EHHM-induced upregulation of Atg5 was inhibited by 3-MA and Baf A1 treatment. As expected, the accumulation of GFP-LC3-II on autophagic vacuoles induced by EHHM



Figure 5. Autophagy inhibition enhances the growth inhibitory and pro-apoptotic effects of EHHM. (A) HepG2 cells were treated with 3-MA or Baf A1, followed by EHHM (15 μ M) treatment for additional 24 h. Western blot analysis was performed to detect the expression levels of Atg5 and Beclin 1. (B) HepG2 cells transfected with a GFP-LC-3-expressing plasmid were treated with 3-MA or Baf A1, followed by treatment with EHHM (15 μ M) for additional 24 h. The cells were then observed under a confocal microscope, and (C) the ratios of cells with autophagy formation were calculated. Scale bar, 10 μ m. (D) HepG2 cells were treated with 3-MA or Baf A1, followed by treatment with EHHM (15 μ M) for additional 24 h, and the cell inhibition rate was then analyzed by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide assay. (E) HepG2 cells were treated with 3-MA, followed by EHHM (15 μ M) treatment for additional 24 h, and the apoptotic cell death rate was analyzed by annexin V/PI staining and flow cytometry. Data represent the mean \pm standard deviation of three independent experiments (*P<0.05). EHHM, E-[6'-(5'-hydroxypentyl)tricosyl]-4-hydroxy-3-methoxycinnamate; GFP, green fluorescent protein; LC3, light chain 3; Atg5, autophagy protein 5; p-, phosphorylated; mTOR, mechanistic target of rapamycin; p70S6K, p70 ribosomal protein S6 kinase; siRNA, small interfering RNA; DMSO, dimethyl sulfoxide; Baf A1, bafilomycin A1; 3-MA, 3-methyl adenine; PI, propidium iodide.



Figure 6. EHHM exhibits antitumor effects on HCC xenograft tumor *in vivo*. (A) Tumor-bearing mice were treated with 3-MA or EHHM, with or without 3-MA, and the tumor volumes were calculated every 2 days. (B) EHHM with or without 3-MA treatment did not affect the body weight in the murine model. EHHM, E-[6'-(5'-hydroxypentyl)tricosyl]-4-hydroxy-3-methoxycinnamat; 3-MA, 3-methyl adenine.

was also decreased following 3-MA treatment (Fig. 5B and C). Accordingly, 3-MA pretreatment significantly promoted EHHM-induced growth inhibition and apoptotic cell death in HepG2 cells (Fig. 5D and E). Baf A1 treatment markedly promoted the accumulation of GFP-LC3-II autophagic vacuoles in EHHM-treated cells, indicating that Baf A1 inhibited the progression of autophagy by blocking the fusion of autophagosomes with lysosome (Fig. 5B). Similar to 3-MA, Baf A1 pretreatment also significantly promoted EHHM-induced growth inhibition of HepG2 cells (Fig. 5D). These results suggested that EHHM-induced autophagy is a protective response to cell death, and that inhibition of this type of autophagy can enhance the anti-HCC effect of EHHM.

EHHM exhibits antitumor effects on HCC xenograft tumors in vivo. To further evaluate the anti-tumor effect of EHHM

in vivo, 2.0×10^6 HepG2 cells suspended in 100 μ l of serum-free DMEM were subcutaneously inoculated into nude mice to generate xenografted murine models. When the tumors could be measured, the tumor-bearing mice were randomly divided into four groups (n=6 mice/group) and administered vehicle control, 3-MA (24 mg/kg) or EHHM (50 mg/kg), alone or in combination with 3-MA, every 2 days. As shown in Fig. 6A, the tumor growth was markedly inhibited by EHHM compared with the vehicle control and 3-MA groups, and 3-MA treatment significantly promoted the inhibitory activity of EHHM. By contrast, the above treatments did not significantly reduce the mice body weight, which suggested that EHHM plus 3-MA had no apparent side effects (Fig. 6B). These results preliminarily suggested that EHHM has a potent antitumor activity both in vitro and in vivo, and that the combination of EHHM and autophagy inhibitors provides synergistic anti-proliferative and pro-apoptotic effects in human HCC cells.

Discussion

A previous study reported that EHHM purified from Livistona chinensis is a key active agent, which exhibited potential anticancer activities towards several cancers, including human myeloid leukemia, HCC and human nasopharyngeal carcinoma (9). HCC is the most common primary cancer in East Asia, particularly in China (47-49), and viral infection, aflatoxin and cirrhosis are the main causes (50). In the present study, the anti-HCC effect and possible mechanisms of EHHM were investigated. The results revealed that EHHM efficiently inhibited the proliferation and induced the apoptosis of HepG2 cells in a time- and dose-dependent manner. In addition, EHHM treatment caused changes in the mitochondrial transmembrane potential, release of cytochrome c from the mitochondria to the cytosol, and activation of the caspase cascade, which indicated that EHHM-induced apoptosis may occur mainly through the activation of the mitochondrial pathway.

Unexpectedly, it was also observed that EHHM treatment induced autophagy formation in HCC cells. EHHM treatment increased the accumulation of autophagic vacuoles, promoted LC3-I/II conversion, and upregulated the expression of several important autophagosome-regulatory genes such as Atg5 and Beclin 1 in HepG2 cells. In fact, accumulating evidence indicated that autophagy could be induced in response to various anticancer treatments in different types of cancer cells, including HCC (51). The PI3K/mTOR/p70S6K signaling pathway has been implicated in regulating autophagy and cell death in response to anticancer treatments (43). In the present study, it was also demonstrated that EHHM treatment significantly inhibited the phosphorylation of Akt, mTOR and p70S6K in a time- and dose-dependent manner, which indicated that EHHM-induced autophagy is involved in the inactivation of the Akt/mTOR/p70S6K signaling pathway.

However, the role of autophagy induced by anticancer treatment is not yet fully understood. Currently, there are different perspectives for the role of autophagy in cancer development and chemotherapy (20). Autophagy has been mainly divided into two types depending on its roles for cell fate: One is protective autophagy, which protects cancer cells from death; the other one is autophagic programmed cell death, which can initiate a cell death process in response to anticancer treatments (20). The different roles of autophagy may be related to different cell types and different anticancer therapies (44). Accumulating evidence indicated that protective autophagy may be an important factor that confers cellular resistance to chemotherapy (52). In the present study, the role of EHHM-induced autophagy in cell proliferation and apoptosis in HepG2 cells was investigated. Pretreatment with Atg5 siRNA or autophagy inhibitors (siRNA against Atg5, 3-MA or Baf A1) inhibited EHHM-induced autophagy, and significantly enhanced the EHHM-induced growth inhibitory activity and pro-apoptotic ability, indicating that EHHM-induced autophagy is a protective autophagy. Similarly, several earlier studies have also reported that the use of autophagy inhibitors or siRNAs targeting autophagy-associated genes significantly promoted the apoptosis induced by anticancer treatments in various cancer cells (23,53-55). Therefore, our results, together with previous findings, raise the possibility that there is an unknown crosstalk or feedback network between autophagy and apoptosis in response to anticancer treatments. Furthermore, our study also suggests an attractive strategy for anti-HCC therapy: EHHM in combination with an autophagy inhibitor may have a synergistic effect in HCC chemotherapy.

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