

# Galangin suppresses human laryngeal carcinoma via modulation of caspase-3 and AKT signaling pathways

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**Abstract.** Laryngeal cancers are mostly squamous cell carcinomas. Although targeting radio-resistant cancer cells is important for improving the treatmental efficiency, the signaling pathway- and therapeutic strategy-related to laryngeal carcinoma still require further study. Galangin is an active pharmacological ingredient, isolated from propolis and *Alpinia officinarum Hance*, and has been reported to have anticancer and anti-oxidative properties through regulation of cell cycle, resulting in angiogenesis, apoptosis, invasion and migration without triggering any toxicity in normal cells. PI3K/AKT and p38 are important signaling pathways to modulate cancer cell apoptosis and proliferation through caspase-3, NF- $\kappa$ B and mTOR signal pathways. Autophagy is also enhanced by activating LC3s and Beclin 1. In the present study, galangin was found to suppress laryngeal cancer cell proliferation. Also, flow cytometry, immunohistochemical and western blot analysis indicated that cell apoptosis was induced for galangin administration, promoting caspase-3 expression through regulating PI3K/AKT/NF- $\kappa$ B. Furthermore, galangin inhibited laryngeal cancer cell proliferation, related to p38 inactivation by galangin treatment. Additionally, mTOR activation regulated by PI3K/AKT was reduced by galangin, suppressing cancer cell transcription and proliferation. Our data also indicated that the tumor volume and weight in nude mice were reduced for galangin use *in vivo* accompanied by Ki-67 decrease and TUNEL increase in tumor tissues. Together, our data indicated that galangin has a potential role in suppressing human laryngeal cancer via inhibiting tumor cell proliferation, activating apoptosis and autophagy, which were regulated by p38 and AKT/NF- $\kappa$ B/mTOR pathways, providing a therapeutic strategy for human laryngeal cancer treatment.

## Introduction

Laryngeal cancer consists mostly of squamous cell carcinomas (also known as the larynx or laryngeal carcinoma), indicating their origin from human skin of larynx (1,2). According to GLOBOCAN 2012, there were an estimated 156,877 new cases and 83,376 deaths in the world, and the adjusted incidence and mortality rates were 2.1/100,000 and 1.1/100,000, respectively (3). Cancer could develop in any part of the larynx according to a previous study. However, the cure rate presently is influenced by the tumor location (4). The origin of laryngeal cancer is a specialised area, which needs the coordinated expertise of ear, nose and throat surgeons and oncologists (5,6). The laryngeal cancer symptoms rely on the tumor size and location. Specific treatment has a close relationship with the type, location, as well as the stage of laryngeal cancer (7,8). Treatment includes radiotherapy, surgery and chemotherapy, alone or in combination. However, a patient affected severely may need a laryngectomy, the total or the partial remove of the vocal cords. In 2013, data indicated that laryngeal cancer has led to a large number of deaths since 1990 (9). Furthermore, the survival rate of patients with laryngeal cancer has decreased compared with the survival rate of patients with all other types of cancers as a whole (4). Hence, finding effective and safe therapeutic strategy is urgently needed to improve cancer treatments. It has been indicated that the treatment of a variety of drugs targeting different signaling pathways can provide effective strategy for various cancer cell mutations, and postpone cancer adaptation procedure subsequently.

Galangin (Fig. 1A) is a member of the flavonol class of flavonoids, which occurs at high concentrations in the rhizome of *Alpinia officinarum Hance*, as well as in propolis, applied in China for centuries as a spice and a traditional Chinese medicine for various diseases (10,11). As a potential scavenger of free radicals, including singlet oxygen and superoxide anion, galangin has various bioactivities and influences many cellular processes (12). In addition to its anti-mutagenic, anti-oxidant, as well as anti-inflammatory functions, galangin has been reported to possess antitumor role in a number of *in vitro* and *in vivo* systems, including melanoma, hepatoma and leukaemia (13,14). Furthermore, a previous study suggested that galangin possesses therapeutic potential as an antitumor agent for liver cancer (15). Galangin induced apoptosis by suppressing tumor cell migration, promoting caspase-3 and increasing ROS

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production (16). However, the effect of galangin on human laryngeal cancer progression has not been investigated yet. In the present study, we investigated the anticancer effects of galangin on two types of human laryngeal cancer cells.

Apoptosis is a distinct genetic and biochemical pathway of cell death necessary for cell growth, development and maintenance of homeostasis in organisms. Caspase-9 was activated with the elevated level of cleaved caspase-9, which in turn cleaved caspase-3 and ultimately induced apoptosis (17,18). Autophagy is a cellular process of catabolic degradation in which damaged, dysfunctional, or superfluous organelles and proteins are sequestered, engulfed, and recycled to maintain cellular metabolism, viability and homeostasis (19,20). The mTOR kinase, which is activated by signaling pathway originating from growth factors, plays a critical role in regulating autophagy progression (21). There are diverse signaling pathways implicated in the regulation of mTOR signaling, including positive regulation of mTOR (PI3K/Akt and p38 MAPK signaling) suppressing autophagy (22,23). The PI3K/Akt signaling pathway is implicated in cell migration and invasion, which induces the expression of NF- $\kappa$ B transcription factor, resulting in cancer cell proliferation (24).

This study is the first time that galangin was used to treat human laryngeal cancer TU212 and HEP-2 cells, to prove if galangin had a potential effect on inhibiting laryngeal cancer progression in the future. In the present study, TU212 and HEP-2 cells, exposed to galangin, suppressed cancer cell proliferation, migration and invasion. Caspase-3, PI3K/AKT and p38 signaling pathways were investigated. These results indicated that galangin performed its anticancer effect on human laryngeal cancer possibly via inactivating PI3K/AKT- and p38-signaling pathways.

## Materials and methods

**Cells and treatment.** Human laryngeal cancer cell lines, TU212 and M4e, human normal larynx epithelial HBE cells, and human normal liver HHL-5 cells, were purchased from the American Type Culture Collection (ATCC; Manassas, VA USA). The mouse normal larynx epithelial RTE cells, and human laryngeal cancer HEP-2 cells were purchased from the Nanjing KeyGen Biotech, Co., Ltd. (Nanjing, China). TU212, HEP-2 and RTE cells were routinely cultured in RPMI-1640 medium (Gibco, Carlsbad, CA, USA), containing 10% fetal bovine serum (FBS; Gibco), 1% penicillin/streptomycin. The cell lines M4e, HBE and HHL-5 were cultured in Dulbecco's modified Eagle's medium (DMEM; Gibco) supplemented with 10% FBS, 100 U/ml penicillin and 100  $\mu$ g/ml streptomycin. All cells were kept in a humidified atmosphere with 5% CO<sub>2</sub> and 95% humidity at 37°C in an incubator. Galangin (>98% purity), purchased from the Hangzhou DayangChem, Co., Ltd., (Hangzhou, China) was used for the treatment of human laryngeal cancer dissolved in dimethyl sulfoxide (DMSO) and then stored at -20°C for experimental treatment use. The final DMSO concentration in cells is <0.1% (v/v) in each treatment.

**Cell viability analysis.** In order to calculate the growth inhibitory role of galangin in different cell lines,  $\sim 1 \times 10^3$  cells/well were seeded in plates (Corning Inc., Corning, NY, USA) with the respectively complete growth media. The following day,

the cells were treated with different concentrations of galangin for different time as shown in the figures and incubated at 37°C. Then, cell viability was calculated by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl-2H-tetrazolium bromide (MTT) analysis at 570 nm.

**Wound healing analysis.** TU212 and HEP-2 cells were seeded in 60-mM dishes and incubated until confluence. After a 3-h cell pre-treatment with 50  $\mu$ M mytomicin C, wounds were created by scratching cell sheets with a sterile 200- $\mu$ l pipette tip. The culture medium was replaced with fresh medium containing either DMSO or galangin. The images of a specific position on the scratched areas were taken by an inverted microscope (Leica Microsystems, Wetzlar, Germany) using a x10 objective every 24 h. The wound widths were assessed and the relative wound widths were evaluated.

**Cancer cell migration and invasion.** TU212 and HEP-2 cells were seeded into the upper chamber of a Transwell insert pre-coated with 5  $\mu$ g/ml fibronectin for migration or a BD™ Matrigel invasion chamber for invasion. Medium with 10% serum was put in the lower chamber as a chemo-attractant, and cells were then incubated for 4 h of migration invasion. Non-migratory and non-invasive cancer cells were removed from the upper chamber using a cotton bud. The cells on the lower insert surface were stained with Diff-Quick. Cells were finally calculated as the number of cells observed in five different microscope fields of three independent inserts.

**Apoptosis analysis.** Apoptosis assay of tissue samples was determined by TUNEL used an In Situ Cell Death Detection kit, fluorescein (Roche Applied Science, Indianapolis, IN, USA) following the manufacturer's protocol. Tumor tissue sections were counterstained with hematoxylin. Then, the number of TUNEL-positive cells was evaluated under a microscope. The number of apoptosis cells were counted by the ratio of apoptotic cells to the total cells. The experiment was conducted three times independently for each cell line.

TU212 and HEP-2 cell apoptosis was determined by flow cytometric (FCM) analysis. The tumor cells were harvested, and then washed three times with chilled phosphate-buffered saline (PBS), stained with Annexin V-FITC and propidium iodide (PI) diluted in the binding buffer, and tested by FACSCalibur FCM (BD Biosciences, San Jose, CA, USA) for 15-min incubation at the room temperature in the dark. Fluorescence was then detected at an excitation wavelength of 480 nm through 530 nm FL-1 and 585 nm FL-2 filters. The apoptotic cells were then quantified.

**Western blot analysis.** For western blot analysis, the cancer cells of TU212 and HEP-2 were washed with chilled PBS and lysed on ice in modified RIPA buffer, containing 50 mM Tris-HCl pH 7.4, 1% NP-40, 0.25% Na-deoxycholate, 150 mM NaCl, 1 mM Na<sub>3</sub>VO<sub>4</sub>, and 1 mM NaF, with protease inhibitors (100  $\mu$ M phenylmethylsulfonyl fluoride, 10  $\mu$ g/ml leupeptin, 10  $\mu$ g/ml pepstatin and 2 mM EDTA). The lysates were centrifuged at 12,000 x g for 20 min at 4°C and the supernatant fractions were collected. The proteins were separated by SDS-PAGE electrophoresis and transferred to Immobilon-P membranes. The specific proteins were

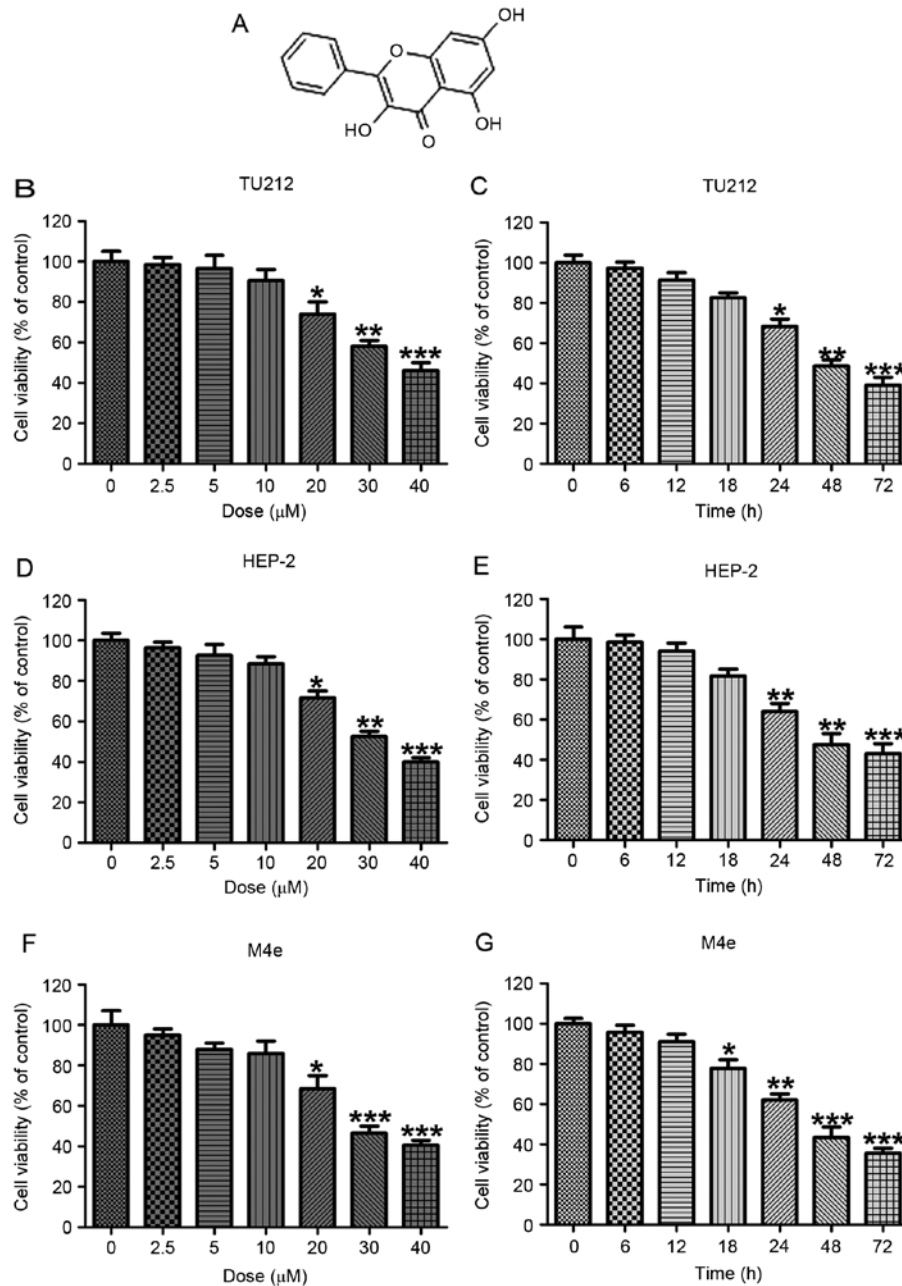


Figure 1. The human laryngeal carcinoma cell viability was calculated by MTT analysis. (A) The chemical structure of galangin. (B) The human laryngeal carcinoma TU212 cells were treated with different concentrations of galangin (0-40  $\mu\text{M}$ ) for 24 h, and the cell viability was calculated by MTT. (C) TU212 cells were treated with 10  $\mu\text{M}$  galangin for the indicated time. The cell viability was evaluated by MTT assays. (D) The human laryngeal carcinoma HEP-2 cells were treated with different concentrations of galangin (0-40  $\mu\text{M}$ ) for 24 h, and then the cell viability was calculated by MTT. (E) HEP-2 cells were treated with 10  $\mu\text{M}$  galangin for the indicated time. The cell viability was evaluated by MTT assays. (F) The human laryngeal carcinoma M4e cells were treated with different concentrations of galangin (0-40  $\mu\text{M}$ ) for 24 h, and the cell viability was calculated by MTT. (G) M4e cells were treated with 10  $\mu\text{M}$  galangin for the indicated time. The cell viability was evaluated by MTT assays. All analysis were conducted in triplicate, and the results are the mean  $\pm$  SEM of three independent experiments. \* $P < 0.05$ , \*\* $P < 0.01$  and \*\*\* $P < 0.001$  (compared to the control/Con).

detected using an enhanced chemiluminescence (ECL) western blotting kit according to the manufacturer's instructions. The primary antibodies including PI3K, Bcl-2, Bax, caspase-9, caspase-3, PARP, LC3, Beclin 1, Ras, Raf, p38, AKT, p-AKT, NF- $\kappa\text{B}$ , p-NF- $\kappa\text{B}$  and GAPDH. GAPDH was used as the loading control. The gray value of the western blotting band was analyzed by ImageJ software (Version 1.4.2b, Mac OS X; National Institutes of Health, Bethesda, MD, USA USA), representing the specific protein expression levels.

*Nude mouse xenograft tumor assay.* The 6-week-old SPF male BALB/c nude mice, weighed 20-25 g, and were obtained from Vital River Laboratory Animals Co., Ltd. (Beijing, China). Before the experiments, all mice were required to adapt to the environment for a week. They were housed in a specific pathogen-free, temperature and humidity-controlled environment ( $25 \pm 2^\circ\text{C}$ ,  $50 \pm 2\%$  humidity) with a standard 12-h light/12-h dark cycle with food and water in their cages. All procedures were in accordance with the Regulations of Experimental Animal Administration issued by the Ministry of Science and

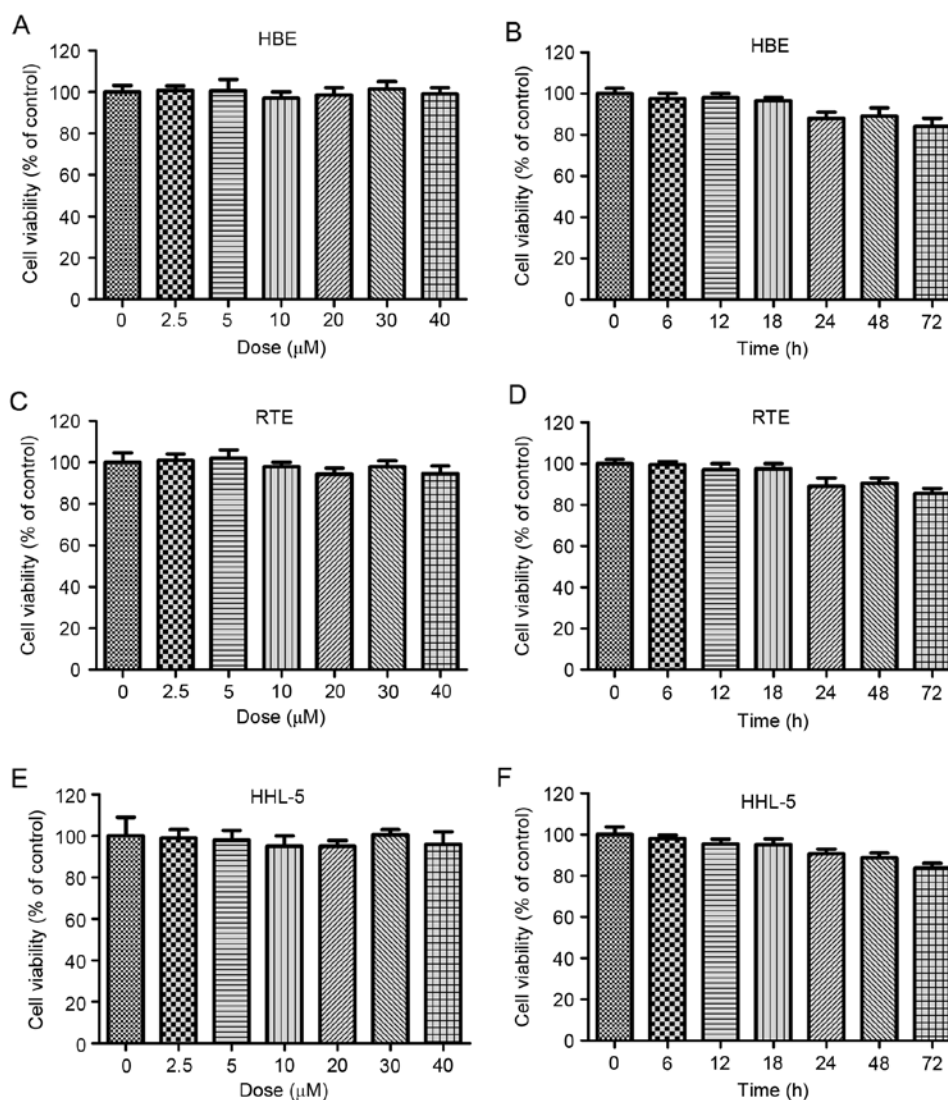


Figure 2. Galangin shows no toxicity on normal cells. (A) The human larynx epithelial HBE cells were treated with different concentrations of galangin, ranged from 0 to 40  $\mu\text{M}$ , for 24 h, and then the cell viability was calculated by MTT. (B) HBE cells were treated with 40  $\mu\text{M}$  galangin for the indicated time. The cell viability was evaluated by MTT assays. (C) The mouse larynx epithelial RTE cells were treated with different concentrations of galangin, ranged from 0 to 40  $\mu\text{M}$ , for 24 h, and the cell viability was calculated by MTT. (D) RTE cells were treated with 40  $\mu\text{M}$  galangin for the indicated time. The cell viability was evaluated by MTT assays. (E) The human normal liver HHL-5 cells were treated with different concentrations of galangin, ranged from 0 to 40  $\mu\text{M}$ , for 24 h, and then the cell viability was calculated by MTT. (F) HHL-5 cells were treated with 40  $\mu\text{M}$  galangin for the indicated time. The cell viability was evaluated by MTT assays. All analysis were conducted in triplicate, and the results are the mean  $\pm$  SEM of three independent experiments.

Technology of China. The mice were randomly divided into 4 groups: the control (Con) group, the galangin group (10 mg/kg), the galangin group (20 mg/kg) and the galangin group (30 mg/kg). Subsequently, 150  $\mu\text{l}$  of the TU212 cell suspension (containing  $2 \times 10^7$  cells) was injected subcutaneously into the right axilla area of the mice. Ten days before the cancer cell inoculation, the treatment groups were gavaged with galangin every day and observed for the growth of tumors at 42 days. The control mice were only injected with the equal volume of TU212 cells. The growth of tumors were determined after the mice were sacrificed, and stored for the following experiments.

**Histopathological examination of tissues.** Histopathological evaluation was performed on mice that were collected. Samples were fixed with 10% buffered formalin, imbedded in paraffin and sliced. Samples were subjected to immunohistochemical staining (Ki-67) according to CST Technology Co. Introduction

and performed by Shanghai Zhenda Biotechnology, Co., Ltd. (Shanghai, China).

**Immunofluorescence assay.** TU212 and HEP-2 cells were cultured on sterilized glass coverslips overnight and treated with galangin for 24 h. After being fixed with 4% paraformaldehyde solution and blocked with 4% BSA in PBS, cells on coverslips were incubated with TSC1, LC3II and Bax primary antibody and anti-rabbit secondary antibody conjugated with Alex Flour 555. Images were captured with a fluorescence microscope (Olympus, Tokyo, Japan).

**Statistical analysis.** Data were expressed as mean  $\pm$  SEM from three or more experiments. Treated cells and the corresponding controls were compared using GraphPad Prism (version 6.0; GraphPad Software, Inc., La Jolla, CA, USA) by a one-way ANOVA with Dunn's least significant difference

tests. Differences between groups were considered significant at  $P < 0.05$ .

## Results

*Galangin inhibits human laryngeal carcinoma cell viability without toxicity on normal cells.* The human laryngeal carcinoma cell viability treated by galangin was calculated. As shown in Fig. 1B, TU212 cells were treated with various concentrations of galangin for 24 h, then the MTT analysis was used to calculate cell viability. With the increasing of galangin concentration, we found that TU212 cell viability was reduced, especially at  $>10 \mu\text{M}$ , showing significant difference compared to the Con ones. Additionally, after the TU212 cells were treated with  $10 \mu\text{M}$  galangin for different time as indicated, we found that the cell viability was reduced in a time-dependent manner (Fig. 2C). Furthermore, in human laryngeal carcinoma HEP-2 cells, the cell viability was decreased with the upregulation of galangin concentrations, especially at  $>10 \mu\text{M}$  (Fig. 1D). Also, HEP-2 cell viability was decreased by  $10 \mu\text{M}$  galangin administration for different times. In addition, significant difference was observed for over 24 h (Fig. 1E). Finally, M4e cells were also included to confirm the suppressive role of galangin in human laryngeal carcinoma. From Fig. 1F, shows that the M4e cell viability was downregulated for galangin treatment, which was comparable to the Con ones at  $>20 \mu\text{M}$ . Similarly, the cell viability was lower by  $10 \mu\text{M}$  galangin treatment for different times (Fig. 1G). The data indicated that galangin indeed has a potential value in reducing human laryngeal carcinoma cell proliferation in a dose- and time-dependent manner.

Galangin has been proved to be toxic for human laryngeal carcinoma cells, and further study was needed to calculate its effects on normal cells. As shown in Fig. 2A and B, no significant difference was observed in human normal larynx epithelial HBE cells, between the various groups treated under different concentrations of galangin for different times. In line with the results above, in the mouse larynx epithelial cells, no significant difference was observed after galangin treatment under different conditions (Fig. 2C and D). The human normal liver HHL-5 cells were included to measure the toxicity of galangin on normal cells. As shown in Fig. 2E and F, galangin treatment under different conditions did not alter the cell viability of HHL-5. The results indicated that galangin shows no toxicity to normal cells, providing its effectivity in human laryngeal carcinoma treatment in future.

*Galangin promotes laryngeal carcinoma cell morphology alteration and suppresses migration, invasion as well as proliferation.* Exposure of TU212 and HEP-2 cells to galangin at different concentrations resulted in necrotic morphological changes and a downregulation in the percentage of viable cells, which was dose-dependent (Fig. 3A). Pretreatment with galangin considerably increased the inhibition of migrated cancer cells (Fig. 3B and C). In addition, we determined the invasion of TU212 and HEP-2 cells treated with galangin at the indicated dose. Fig. 4A shows that the invasion of TU212 and HEP-2 cells was markedly downregulated for galangin treatment compared to the control ones. In addition, images were taken at 0 and 24 h after galangin treatment as shown

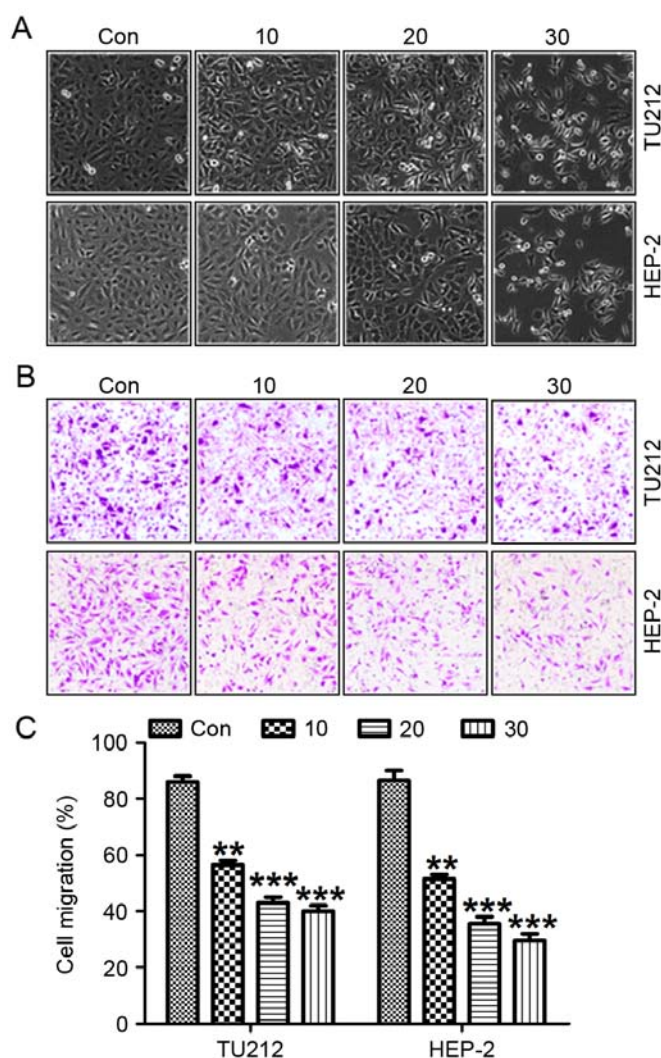


Figure 3. Galangin promotes laryngeal carcinoma cell morphology alteration and migration suppression. (A) Galangin induced morphological changes in human laryngeal carcinoma cells of TU212 and HEP-2. (B) The representative images of migrated cancer cells are shown. (C) Galangin reduced the number of migrated TU212 and HEP-2 cells under different conditions. All analysis was conducted in triplicate, and the results are mean  $\pm$  SEM of three independent experiments. \*\* $P < 0.01$  and \*\*\* $P < 0.001$  (compared to the control/Con).

in Fig 4B and C. TU212 and HEP-2 cells treated with galangin for 24 h indicated that galangin suppressed the cancer cell migration, which was comparable to the control group in the absence of galangin. These results suggested that galangin suppressed TU212 and HEP-2 cell proliferation.

*Galangin induces apoptosis in human laryngeal carcinoma cells of TU212 and HEP-2.* In this regard, we attempted to explore the mechanism of galangin on TU212 and HEP-2 cell growth and progression. The data from flow cytometric assay clearly showed that galangin induced apoptosis in TU212 cancer cells (Fig. 5A and B) and HEP-2 (Fig. 5A and C). In apoptosis process, mitochondrial outer membrane permeabilization is known as a 'point-of-no-return' and is closely regulated by Bcl-2 family proteins, especially Bax activation (25). Galangin apparently promoted Bax protein expression and pre-treatment with galangin significantly suppressed Bcl-2

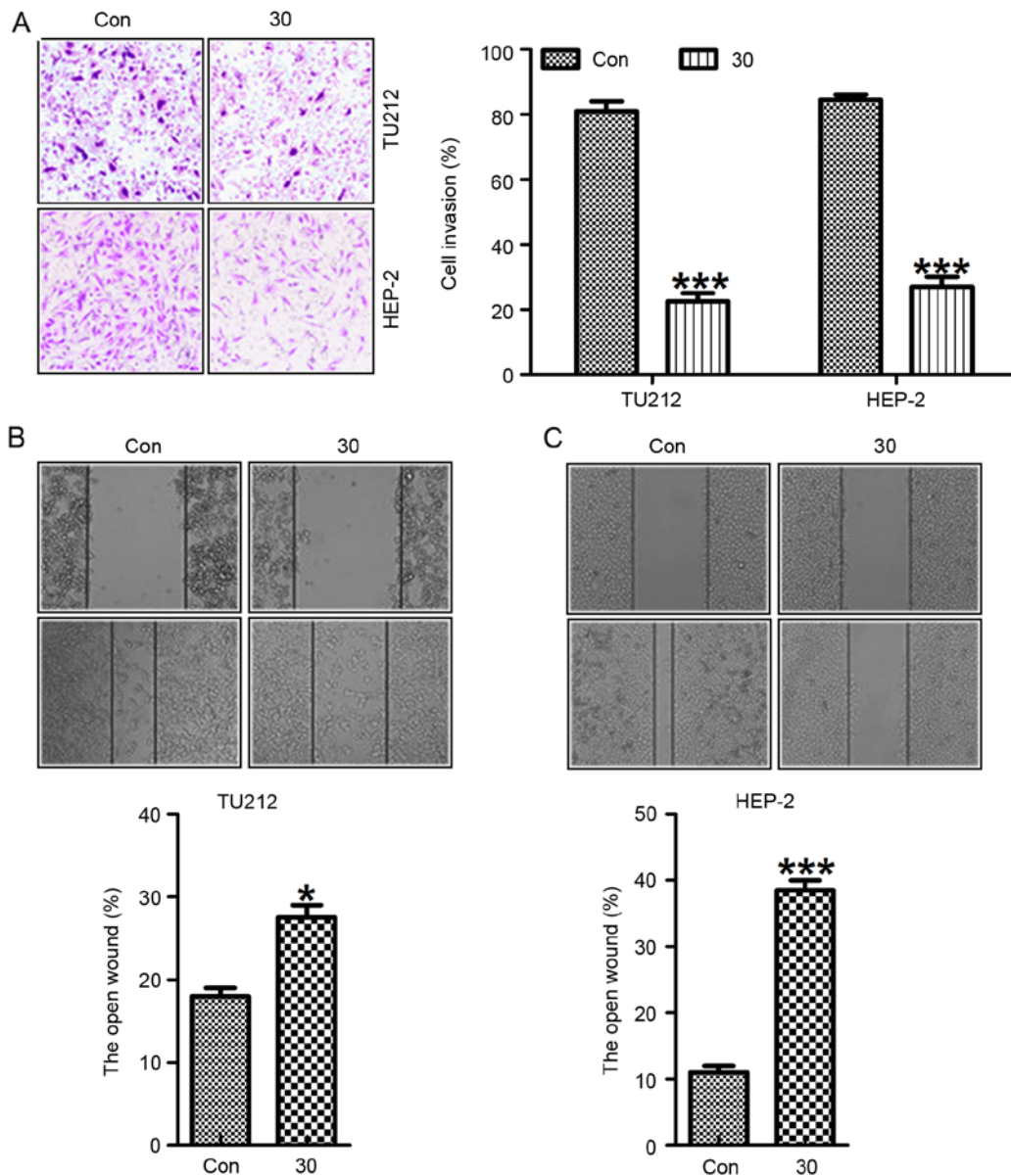


Figure 4. Galangin inhibits the laryngeal carcinoma cell invasion and proliferation. (A) TU212 and HEP-2 cells were treated with different concentrations of galangin, and then the invasive cells were calculated, which is shown in the left panel. After the scratching, (B) TU212 and (C) HEP-2 cells were treated with galangin at the indicated concentration for 24 h. Images were then acquired at 0 and 24 h under the microscope. The black lines represent the areas lacking the migratory cells. Representative pictures from three independent experiments are shown. All analysis was conducted in triplicate, and the results are mean  $\pm$  SEM of three independent experiments. \* $P < 0.05$ ; \*\*\* $P < 0.001$  (compared to the control/Con).

protein levels (Fig. 5D-F). The results from western blot analysis showed that galangin enhanced activation of cleaved caspase-3 and caspase-9, resulting in increased PARP cleavage in TU212 (Fig. 5D and E) and HEP-2 cells (Fig. 5D and F). Furthermore, Bax activation was confirmed to be increased for galangin treatment in TU212 (Fig. 5G and H) and HEP-2 cells (Fig. 5G and I). These results suggested that galangin treatment-triggered apoptosis was modulated through caspase activation.

*Galangin induces human laryngeal carcinoma cell death through autophagy regulation.* LCI, LC3II and Beclin 1 are required for the autophagy-mediated elimination of unfolded ubiquitinated long half-life proteins (26). As shown in Fig. 6A, western blot analysis was carried out to reveal a considerable

increase in LC3I, LC3II and Beclin 1 expression induced by galangin exposure compared to the control group without galangin treatment in TU212 (Fig. 6B) and HEP-2 (Fig. 6C) cells. The increased LC3I, LC3II and Beclin 1 further suggested that galangin stimulated autophagy and cell death in human laryngeal carcinoma (27). Similarly, in Fig. 6D-F, immunofluorescent assays further evidenced that LC3II was highly induced in galangin treatment. The data above indicate that galangin could result in human laryngeal carcinoma cell death, contributing to tumor suppression.

*Galangin inhibits human laryngeal carcinoma TU212 and HEP-2 cell proliferation via p38 and AKT/NF- $\kappa$ B suppression.* To further explore the potential molecular mechanism involved in galangin-induced TU212 and HEP-2 cell death,

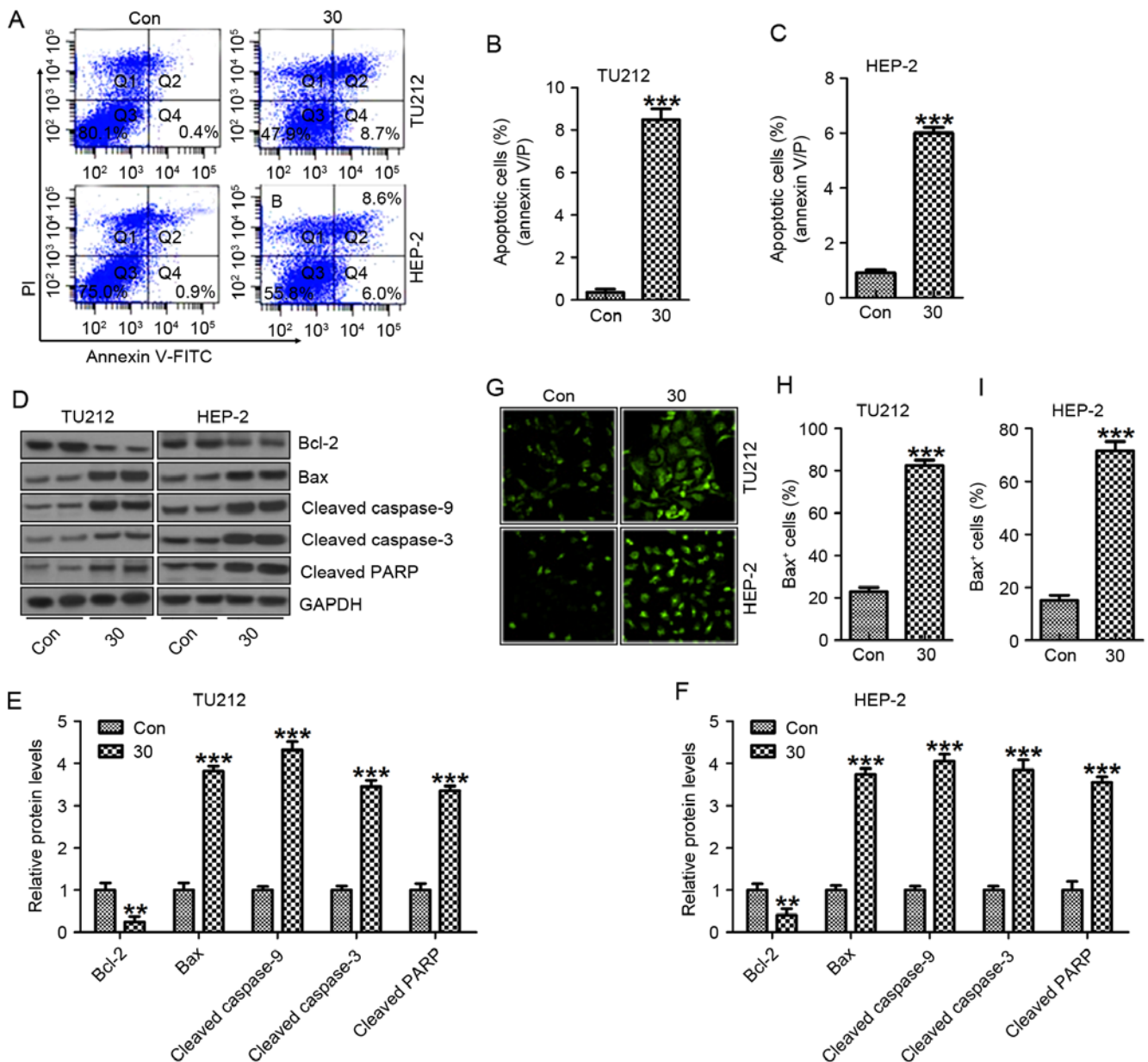


Figure 5. Galangin induces apoptosis in human laryngeal carcinoma TU212 and HEP-2 cells. (A) Flow cytometric assays were applied to determine the number of apoptotic TU212 and HEP-2 cells. (B) The quantification of TU212 apoptosis levels is shown. (C) The quantification of HEP-2 apoptosis levels was shown. (D) The signaling pathway, leading to apoptosis, was analyzed by western blot analysis in TU212 and HEP-2 cells. Protein levels of Bcl-2, Bax, cleaved caspase-9, cleaved caspase-3 and cleaved PARP were evaluated. (E) Protein levels of Bcl-2, Bax, cleaved caspase-9, cleaved caspase-3 and cleaved PARP in TU212 cells were quantified. (F) Protein levels of Bcl-2, Bax, cleaved caspase-9, cleaved caspase-3 and cleaved PARP in HEP-2 cells were quantified. (G) Immunofluorescence assays were carried out to determine Bax positive TU212 and HEP-2 cells. The quantification of Bax positive (H) TU212 and (I) HEP-2 cells are displayed. The analysis was conducted in triplicate, and the results exhibit the mean  $\pm$  SEM of three independent experiments. \*\* $P < 0.01$  and \*\*\* $P < 0.001$  (compared to the control/Con).

the protein expression of Raf, Ras and p38 were calculated. The results showed that Raf, Ras and p-p38 was highly expressed in the control group without galangin treatment in both TU212 (Fig. 7A and B) and HEP-2 cells (Fig. 7A and C). Importantly, galangin supplementation partially reversed the overexpression of Raf, Ras and p-p38. Enhancement of p38 activation plays an important role in modulating tumor migration, invasion as well as metastasis in some cancer cases (28). Altogether, galangin treatment-attenuated TU212 and HEP-2 cell progression partially relied on p38 signaling pathway.

NF- $\kappa$ B is a pleiotropic transcription factor, which is related to various biological processes, including inflammation, apoptosis as well as autophagy (29). NF- $\kappa$ B activation has been detected in >50% of tumors, regulated by PI3K/AKT signaling pathway (30). In this study, PI3K/AKT was expressed highly in the cancer cells without galangin treatment, which was downregulated after galangin administration in both TU212 (Fig. 7D and E) and HEP-2 (Fig. 7D and F) cells. These results indicated that galangin might play important roles in TU212 and HEP-2 cells as an antitumor agent inhibiting cancer cell proliferation by targeting p38 and NF- $\kappa$ B.

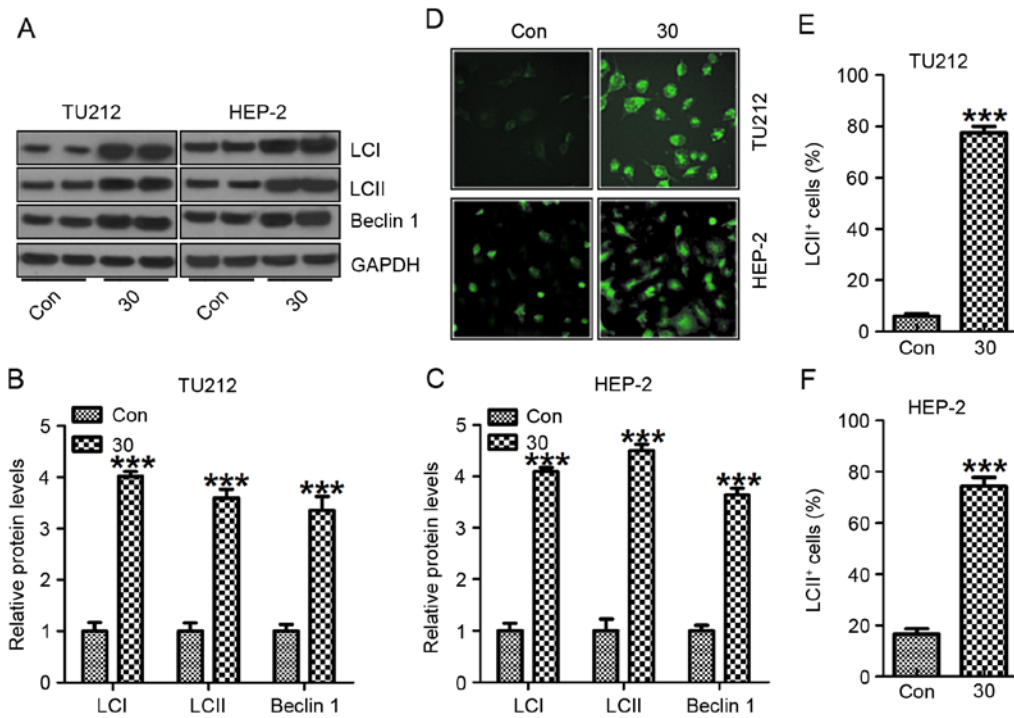


Figure 6. Galangin induces human laryngeal carcinoma cell death through autophagy regulation. (A) LCI, LCII and Beclin 1 protein levels were measured through western blot analysis. The quantification of LCI, LCII and Beclin 1 in (B) TU212 and (C) HEP-2 cells is shown. (D) Immunofluorescence assays were carried out to determine LCII positive TU212 and HEP-2 cells. The quantification of LCII positive cells (E) TU212 and (F) HEP-2 cells is shown. The analysis was conducted in triplicate, and the results are the mean  $\pm$  SEM of three independent experiments. \*\* $P$ <0.01 and \*\*\* $P$ <0.001 (compared to the control/Con).

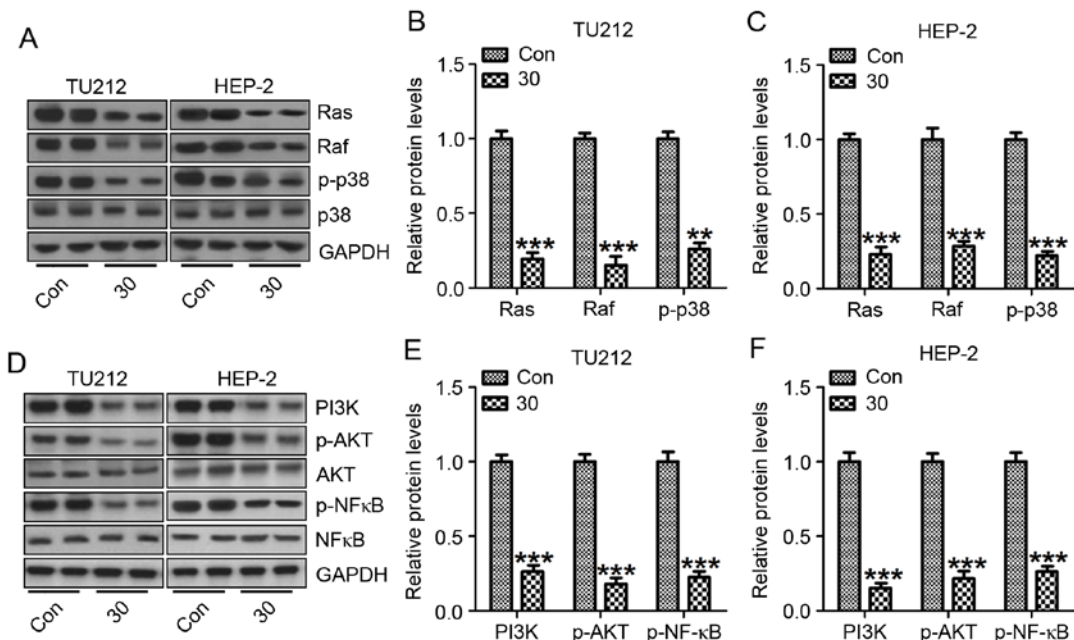


Figure 7. Galangin inhibits human laryngeal carcinoma TU212 and HEP-2 cell proliferation via p38 and AKT/NF- $\kappa$ B suppression. (A) Signals of Ras, Raf and p-p38 were determined by western blot analysis. (B) The quantification of Ras, Raf and p-p38 in TU212 cells. (C) The quantification of Ras, Raf and p-p38 in HEP-2 cells. (D) Signals of PI3K, p-AKT and NF- $\kappa$ B were measured by western blot assays. The quantification of PI3K, p-AKT and NF- $\kappa$ B in (E) TU212, and (F) HEP-2 cells following immunoblot analysis. The analysis was conducted in triplicate, and the results are the mean  $\pm$  SEM of three independent experiments. \*\* $P$ <0.01 and \*\*\* $P$ <0.001 (compared to the control/Con).

Galangin impedes human laryngeal carcinoma TU212 and HEP-2 cell proliferation via suppressing mTOR. Finally, suppressing the PI3K-Akt-mTOR signaling pathway has been considered as an essential molecular mechanism causing tumor

suppression (31). In this regard, we found that TSC1, an inhibitor of mTOR activation, was upregulated after the treatment with galangin. On the contrary, mTOR phosphorylated levels were reduced in TU212 and HEP-2 cells (Fig. 8A-C). Also,



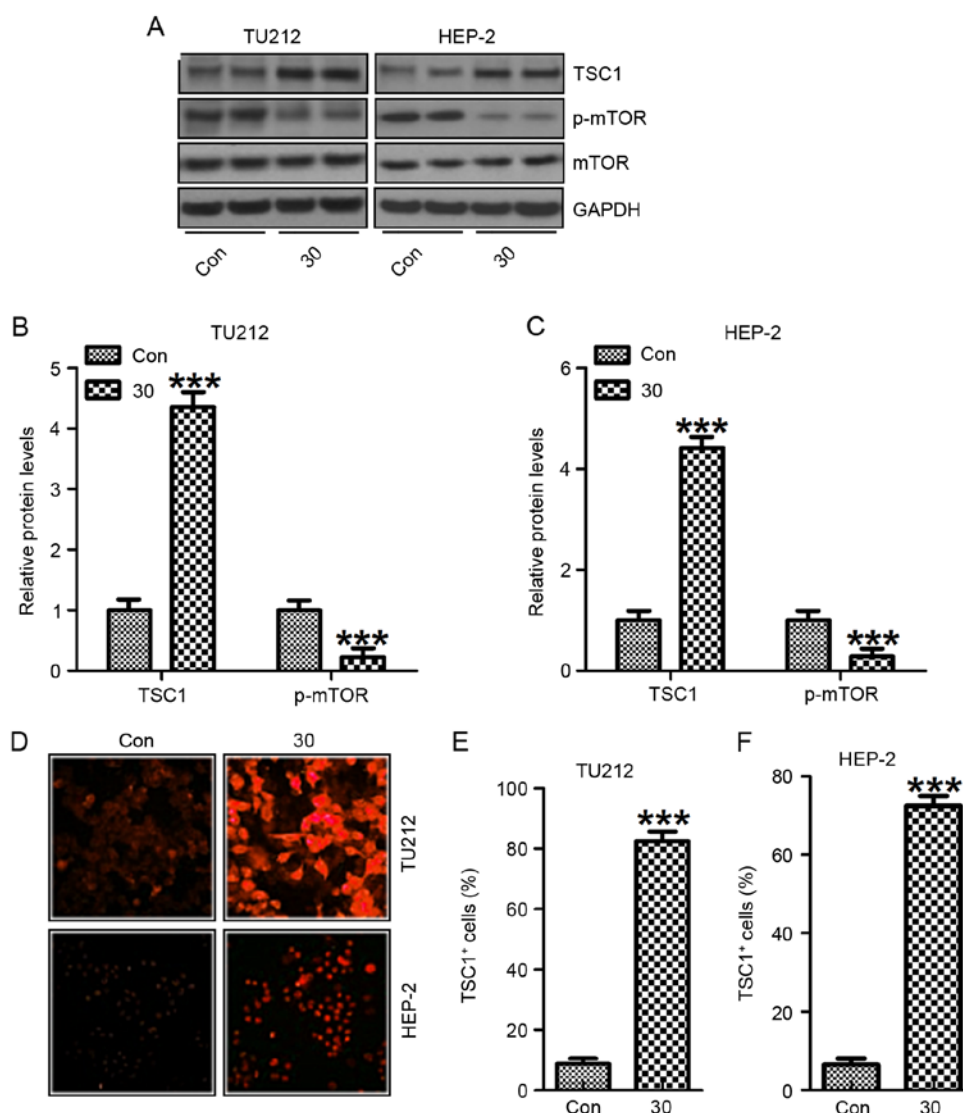


Figure 8. Galangin impedes human laryngeal carcinoma TU212 and HEP-2 cell proliferation via suppressing mTOR. (A) Western blot analysis was carried out to evaluate TSC1, p-mTOR and mTOR expression in TU212 and HEP-2 cells. TSC1 and p-mTOR protein levels in (B) TU212 and (C) HEP-2 cells were quantified after western blot assays. (D) Immunofluorescence assays were carried out to determine TSC1 positive cells in TU212 and HEP-2 cells. The quantification of TSC1 positive cells in (E) TU212 and (F) HEP-2 cells is shown. The analysis was conducted in triplicate, and the results are the mean  $\pm$  SEM of three independent experiments. \*\*\* $P < 0.001$  (compared to the control/Con).

immunofluorescent analysis was performed to explore TSC1 levels in both TU212 (Fig. 8D and E) and HEP-2 (Fig. 8D and F) cells. TSC1 were highly expressed for galangin treatment in comparison to the control ones. Collectively, these results indicated that galangin suppressed human laryngeal carcinoma cell growth via mTOR signaling pathway.

*Galangin promotes human laryngeal carcinoma growth inhibition in a xenograft tumor model in vivo.* To confirm the enhanced galangin-regulated inhibition of human laryngeal carcinoma growth, we analyzed the effects of galangin treatment on tumorigenicity *in vivo* using a TU212 xenograft mouse model. After administration with galangin for 42 days, both the tumor volume and weight were inhibited (Fig. 9A-C). Additionally, reduction of tumor Ki-67 (Fig. 9D and E) and upregulation of TUNEL (Fig. 9D and E) through IHC assays were also noted in the galangin-treated group in a dose-dependent manner. These results indicated that galangin

could promote suppression of xenografted human laryngeal carcinoma cell growth *in vivo*, which was in line with the data *in vitro*.

## Discussion

Human laryngeal cancer is known as one of the most common tumors of the head and neck region in the world (1,2,32). Patients suffering from laryngeal cancer show a poor survival rate with poor advance during the last decades (33). Thus, more effective treatments are needed to be explored for preventing laryngeal cancer. Natural compounds show an essential role in cancer and other disease prevention and treatment worldwide (4). Galangin is known as an anti-tumor agent, which is effective in preventing a broad range of tumors, such as liver, breast, as well as lung cancer (14-16,34). Galangin has been reported to perform its role in inhibiting tumor growth through apoptosis regulation (35). However, whether galangin

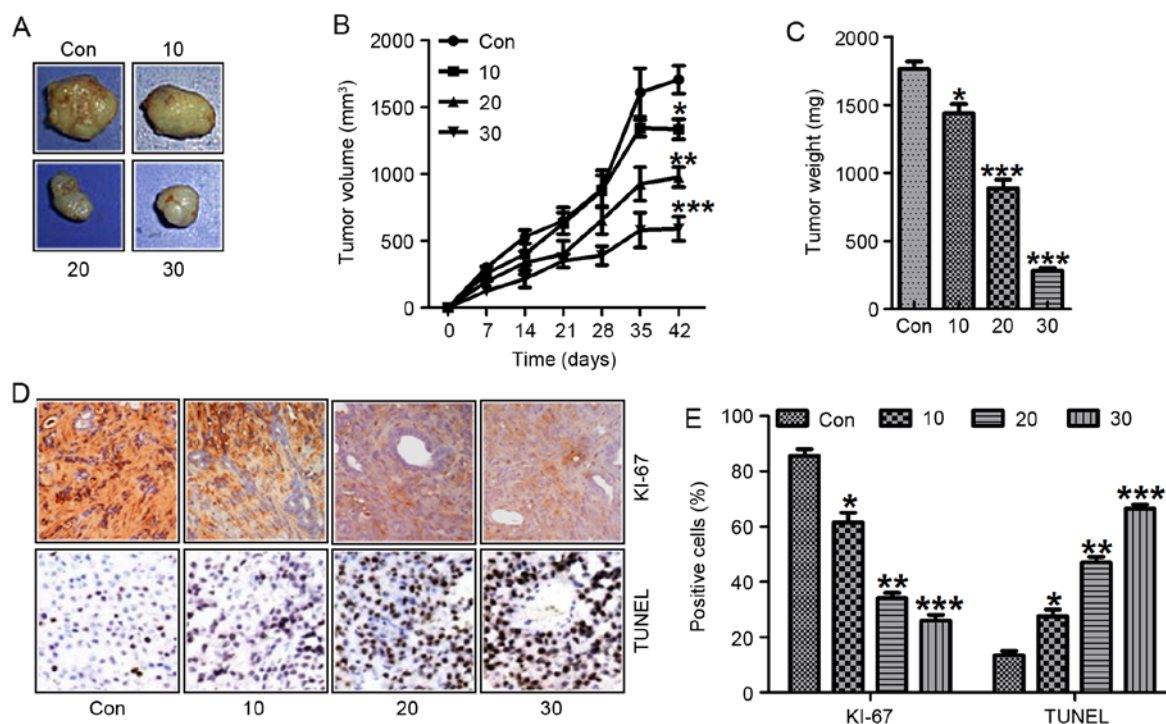


Figure 9. Galangin promotes human laryngeal carcinoma growth inhibition in a xenograft tumor model *in vivo*. (A) The tumor images are shown. (B) Tumor volumes were measured at different times. (C) Tumor weight was calculated at different treat-times. (D) Upper panel, the Ki-67 positive cells in a human laryngeal carcinoma sample were evaluated via immunohistochemical analysis. Lower panel, TUNEL positive cells in a tumor sample were assessed via immunohistochemical analysis. (E) The quantification of Ki-67 and TUNEL positive cells. The analysis was conducted in triplicate, and the results are the mean  $\pm$  SEM of three independent experiments. \* $P < 0.05$ , \*\* $P < 0.01$  and \*\*\* $P < 0.001$  (compared to the control/Con).

could be a potential compound for human laryngeal cancer prevention is not known. Thus, galangin was used here to provide possible therapeutic strategy for suppressing human laryngeal cancer. Notably, it is the first study to explore the molecular mechanisms of galangin-triggered human laryngeal cancer cell death. In our study, we found that exposure of galangin into TU212 and HEP-2 cells could partially suppress cancer cell proliferation, invasion and migration. Autophagy is a regulated process of degradation and recycling of cellular constituents; the process is important in organelle turnover and the bioenergetics management of starvation (36). During autophagy, the precursor form of LC3 is post-modified into LC3-I and LC3-II (37,38). LC3-I is localized in the cytosol, and LC3-II is a membrane-associated and a key hallmark for autophagosome formation (39). LC3-II can be used to estimate the abundance of autophagosomes before they are destroyed through fusion with lysosomes (40). Here, augmented LC3II was found after galangin treatment, indicating that the autophagy was induced. Obviously in *in vivo* study, the tumor size and weight was reduced for galangin administration. The results suggested that galangin has potential, to be developed as a therapeutic strategy for human laryngeal carcinoma.

PI3K/AKT signal pathway is crucial for regulating various cell activities, such as proliferation, cell growth, survival, chemotaxis, the inflammatory response and apoptosis (41). Enhanced activation of PI3K/AKT pathway is linked to the development and progression of many cancers, as well as resistance to chemotherapy (42). A previous study reported that inactivation of PI3K/AKT signaling pathway is involved in the success of chemotherapy-caused apoptosis in some cancer

cells (43,44). In the present study, PI3K was reduced in galangin treatment, subsequently downregulating the protein levels of phosphorylated AKT, being in line with previous results, indicating that galangin might perform its role in suppressing human laryngeal carcinoma through PI3K/AKT signaling pathway inactivation. Apoptosis constitutes a fundamental intrinsic mechanism of tumor suppression, as the resistance of apoptosis is a well-established hallmark of cancer (45,46). It is known that Bcl-2 protein is also a key regulator for apoptosis and its tumorigenic potential is supported by the finding of overexpression of Bcl-2 in various types of tumor, which is related to the activation of AKT (47). AKT is a key player in regulating cell signals that are important for cell death and survival. Activation of the AKT pathway promotes cell survival and is involved in the upregulation of Bcl-2 (48,49). In this study, we found that with the altered trend of AKT expressed levels, Bcl-2 was reduced in galangin treatment. The caspase-3 and PARP cleavage were highly improved, contributing to the death of TU212 and HEP-2 cells and leading to cell apoptosis and cell death due to galangin treatment. The results above suggested that galangin suppressed human laryngeal cancer development and progression through apoptosis induction.

NF- $\kappa$ B takes part in the information transfer process, including tissue damage, apoptosis, as well as stress, tumor suppression and cell differentiation, thus, it is an important nuclear transcription factor (50,51). NF- $\kappa$ B in different tumors or cancers is modulated by PI3K/AKT pathway (52). P-AKT activates I $\kappa$ B kinase (IKK $\alpha$ ), leading to inhibition of NF- $\kappa$ B degradation by I $\kappa$ B, allowing NF- $\kappa$ B to be transferred into the nucleus from the cytoplasm, where it activates its target

genes and promotes cell survival (53). In this study, NF- $\kappa$ B was markedly activated accompanied by AKT phosphorylation in the absence of galangin treatment. However, galangin administration downregulated NF- $\kappa$ B phosphorylated activity, leading to the upregulation of caspase-9, caspase-3 and PARP cleavage, indicating that galangin could inhibit human laryngeal cancer via AKT-mediated NF- $\kappa$ B signaling pathway. In addition, PI3K/AKT pathway is of great importance for the cell proliferation. One of the best-known downstream substrates of PI3K/AKT is the mammalian target of rapamycin (mTOR), inducing mammalian autophagy significantly, regulating protein translation (54). Previously it was indicated that the activation of PI3K-AKT-mTOR signal pathway may be involved in autophagy suppression (55). Impeding PI3K-AKT-mTOR signal pathway has been indicated to be effective for various cancers treatment (56,57). In our investigation, here the phosphorylated mTOR in high expression was suppressed in TU212 and HEP-2 cells for galangin administration, and in contrast, TSC1, an inhibitor of mTOR, was upregulated for galangin. The results illustrated that galangin prevented human laryngeal cancer through inhibiting laryngeal cancer cells proliferation and inducing autophagy via PI3K/AKT-regulated mTOR activity.

Collectively, our findings above demonstrated that galangin prevented human laryngeal cancer proliferation, invasion and migration by PI3K/AKT and p38 suppression, resulting in caspase activation, NF- $\kappa$ B dephosphorylation as well as mTOR inactivation with reduced Ki-67 expression and enhanced TUNEL levels. The present study indicated that the use of the dietary compound galangin might be a potential therapeutic strategy for human laryngeal carcinoma treatment.

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