



Cannabisin B induces autophagic cell death by inhibiting the AKT/mTOR pathway and S phase cell cycle arrest in HepG2 cells

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ABSTRACT

This study investigates the anticancer properties of cannabisin B, purified from hempseed hull, in HepG2 human hepatoblastoma cells. The results indicate that cannabisin B significantly inhibited cell proliferation by inducing autophagic cell death rather than typical apoptosis. Cell viability transiently increased upon the addition of a low concentration of cannabisin B but decreased upon the addition of high concentrations. Cannabisin B-induced changes in cell viability were completely inhibited by pre-treatment with 3-methyladenine (3-MA), indicating that the induction of autophagy by cannabisin B caused cell death. Additionally, cannabisin B induced S phase cell cycle arrest in a dose-dependent manner. Moreover, cannabisin B was found to inhibit survival signaling by blocking the activation of AKT and down-stream targets of the mammalian target of rapamycin (mTOR). These findings suggest that cannabisin B possesses considerable antiproliferative activity and that it may be utilised as a promising chemopreventive agent against hepatoblastoma disease.

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1. Introduction

Autophagy, which is described as the mass degradation of cellular components, is an essential and homeostatic process that enables cells to adapt to adverse situations, such as nutrient deprivation and invasion by microorganisms (Levine, Mizushima, & Virgin, 2011). Autophagic cell death (also known as type II programmed cell death) plays an important role in metabolism and stress responses as well as human diseases, including neurodegeneration, inflammation, muscular disorders, and resistance to pathogens (Mizushima & Levine, 2010). In eukaryotic cells, autophagy commonly occurs within a double-membraned vesicle, which is then hydrolysed by the lysosome. Although the molecular signaling pathways are poorly understood, several molecules have been discovered to regulate autophagy, including members of the BECN1, DAPK (death-associated protein kinase), DRP1 (death-associated related protein kinase 1), MAPK (mitogen-activated kinase), and PI3K-AKT-mTOR (phosphatidylinositol 3-kinase/AKT/mammalian target of rapamycin) pathways (Kelekar, 2008; Todde, Veenhuis, & van der Klei, 2009). Recent reports indicate that

autophagy mediates chemotherapy-induced cell death in cancer, functioning as a suppressor in early tumorigenesis but a stimulator in established tumors (Notte, Leclere, & Michiels, 2011).

Hepatoblastoma is a frequent liver tumor in Western countries often occurring in infants and children. The disease is most commonly diagnosed during a child's first three years and they are usually present with an abdominal mass (De Ioris et al., 2008). Hepatoblastomas originate from immature liver precursor cells. People with familial adenomatous polyposis (FAP), a syndrome of early-onset colonic polyps and adenocarcinoma, frequently develop hepatoblastomas (Hirschman, Pollock, & Tomlinson, 2005). Currently, there is no effective drug treatment for hepatoblastoma except for surgical resection. If metastases occur, the patient's condition is usually out of control. Therefore, the development of new agents, particularly phytochemicals from plants, is important for the treatment of this liver tumor.

Cannabisin B, a lignanamide from hempseed, was first identified in 1992 (Sakakibara, Ikeya, Hayashi, & Mitsuhashi, 1992). The bioactivity of this hempseed secondary metabolite remained unevaluated until recently, when we demonstrated that cannabisin B functions predominantly as an antioxidant in hempseed hull (Chen et al., 2012). Several reports have indicated that the antioxidant substances in plants exhibit antiproliferative activity in cancer cells (He & Liu, 2008; Prasad et al., 2010; Yang, Liu, & Halim, 2009). Thus, the objectives of the current study were (1) to

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examine the antiproliferative activity of cannabisin B and the crude extract of hempseed hull in HepG2 cells and (2) to investigate the autophagy-related molecules associated with the cell death signal transduction pathway.

2. Materials and methods

2.1. Materials

Dulbecco's modified Eagle's medium (DMEM) and fetal bovine serum (FBS) were purchased from GIBCO, NY, USA. The cell counting kit-8, cell cycle analysis kit, annexin V-FITC apoptosis detection kit, and DNA ladder extraction kit with spin columns were obtained from Beyotime Institute of Biotechnology, Jiangsu China. Monodansylcadaverine (MDC), 3-methyladenine (3-MA), and Hoechst 33258 were purchased from Sigma–Aldrich, USA. Acetonitrile (HPLC grade) was purchased from Fisher, USA.

2.2. Sample preparation

The hempseed hull was separated from the hull using a laboratory huller machine (model JLGJ4.5, Zhejiang Liangyi Co., China) and milled in a high speed grinder (FW135, Tianjin Taisite Instrument Co., China) then through 380 μm meshes. The hull powder (1.0 kg) was extracted using 60% EtOH water solution in an ultrasonic bath (KQ-100E, Jiangsu Kunshan Ultrasonic Instrument Co., China) at 40 kHz (20 L \times 30 min). The extract solution was then vacuum-evaporated to remove the EtOH, purified in a HPD-600 resin column, and eluted with water and 20% EtOH. Eluent of 40% EtOH was collected and lyophilized to produce the crude extract (1.0 g, yellow powder). The crude extract was then applied to a Sephadex LH-20 column and eluted using 20%, 40%, and 60% MeOH. The collected 60% MeOH eluent was finally purified by HPLC to obtain cannabisin B (70 mg).

2.3. HPLC measurement

Purification and quantification were conducted using a HPLC system equipped with a DAD (model LC-20A, Shimadzu, Japan). The sample was dissolved in 60% MeOH and filtered through a 0.45 μm filter. Column chromatography was performed on a C18 column (250 mm \times 4.6 mm i.d., 5 μm , model ST, Techmate Co., China) at 35 $^{\circ}\text{C}$ with a monitoring wavelength of 335 nm. The binary mobile phase consisted of (A) water–0.1% acetic acid and (B) acetonitrile–0.1% acetic acid. The gradient elution (at a flow rate of 1 ml/min) was as follows: 0–20 min, 10–30% B; 20–30 min, 30% B isocratic; 30–32 min, 10–30% B; 32–40 min, 10% B isocratic.

2.4. Cell culture and treatment

The HepG2 cell line was purchased from Shanghai Institute of Biochemistry and Cell Biology (SIBCB). Cells were grown in 90% DMEM supplemented with 10% fetal bovine serum (FBS), 100 units/ml penicillin, and 100 $\mu\text{g}/\text{ml}$ streptomycin. The cell cultures were maintained at 37 $^{\circ}\text{C}$ in a humidified atmosphere of 5% CO_2 . The medium was changed every other day, and the cells were subcultured at 1:2 every 3 days. The cells were then plated at an appropriate density according to each experimental scale.

2.5. Cell proliferation assay

WST-8 was employed to analyze cell proliferation because it is reduced the soluble formazan by mitochondrial dehydrogenase and exhibits little toxicity in cells. Cell viability was determined using the WST-8 assay kit (Beyotime Institute of Biotechnology,

China) according to the manufacturer's instructions. Briefly, 1×10^4 cells/well were seeded in a 96-well flat-bottomed plate, grown at 37 $^{\circ}\text{C}$ for 24 h, washed with PBS, and then treated with various concentrations (84, 167, 251, 335, or 503 μM) of cannabisin B extract dissolved in DMEM for 24, 48, or 72 h. Cells incubated in DMEM without cannabisin B extract served as the control. WST-8 dye was added to each well, and the cells were incubated at 37 $^{\circ}\text{C}$ for 2 h. The absorbance was finally determined at both 450 and 630 nm using a microplate reader (Bio-Rad 550, Japan). Values were expressed as the percentage of control cells.

2.6. Cell cycle analysis

The cells were seeded into a 6-well plate, incubated for 24 h, and then treated with concentrations of 167, 335, or 503 μM cannabisin B for 24 h. The cells were then fixed with 70% ice-cold methanol and transferred to the freezer until analysis. Upon analysis, the cells were washed with PBS and then stained with 50 $\mu\text{g}/\text{ml}$ propidium iodide in the presence of 25 $\mu\text{g}/\text{ml}$ RNase A at 37 $^{\circ}\text{C}$ for 30 min. The cell cycle distribution of 10,000 cells was recorded by a flow cytometer (BD FACS Calibur), and the percentage of cells in the G0/G1, S, and G2/M phases was analyzed with ModFit software.

2.7. Morphological detection of HepG2 cells

Cell morphology and nuclear morphology were both assessed with a fluorescent microscope (Olympus IX71, Japan). The cells were treated with 167, 335, or 503 μM for 24 h and remained untreated cells as controls. The cells were observed after being stained with 10 $\mu\text{g}/\text{ml}$ Hoechst 33258 for 15 min. The cells were also stained with MDC in the same manner as described above.

2.8. Analysis of apoptosis by Annexin V-FITC binding

The cells were seeded and treated in the same manner as described for cell cycle analysis. The cells were then collected, washed twice with PBS, and stained using the annexin V-FITC apoptosis detection kit (Beyotime Institute of Biotechnology, China) as described by the manufacturer's instruction (Jiang, Zu, Fu, Zhang, & Efferth, 2008). The fraction of the cell population that appeared in each quadrant was analyzed using quadrant statistics. The lower left quadrant contained intact cells, the lower right quadrant contained apoptotic cells, and the upper right quadrant contained necrotic or post-apoptotic cells. The results were expressed as the percentage of the total number of cells.

2.9. DNA fragmentation analysis

Fragmented DNA was isolated using the DNA extraction kit (Beyotime Institute of Biotechnology, China) according to the manufacturer's instructions. The samples were treated in the same manner as described for cell cycle analysis. The eluents containing DNA pellets were electrophoresed on a 1.0% agarose gel at 80 V for 1.5 h. The gel was examined and photographed using an ultraviolet gel imaging system.

2.10. Transmission Electron Microscopy (TEM)

The HepG2 cells were treated with cannabisin B for 24 h and then collected and centrifuged at 1000g for 5 min. The fixation procedures and the manufacturing of resin samples were manipulated according to a previous report (Lin et al., 2007). The thin slice was observed with a Hitachi 7650 electron microscope (Tokyo, Japan).

2.11. Western blot analysis

After treatment with cannabisin B, cells in 6 cm plate were carefully rinsed twice with PBS. After lysis in cold radioimmunoprecipitation assay buffer containing protease inhibitors, the cells were scraped and centrifuged at 13000g for 10 min at 4 °C. The liquid supernatants were then collected and mixed with electrophoresis loading buffer. The protein samples were normalized by loading equal amounts of total protein lysates. Protein samples were separated by SDS-PAGE, transferred onto a nitrocellulose membrane, and then probed with the primary antibody, which was followed by the addition of the secondary antibody conjugated with alkaline phosphatase. The bands were finally detected using the Alkaline Phosphatase Color Development Kit (Beyotime Institute of Biotechnology, China). Antibodies specific for β -actin, LC-3B, Akt, phospho-Akt (Ser473), mTOR, and phospho-mTOR were purchased from Beyotime Institute of Biotechnology (Jiangsu, China). The assay was repeated at least three times and the clearest one would be imaged for analysis.

2.12. Statistical analysis

Statistical analyses were performed using the SPSS 16.0 software package (SPSS Inc., Chicago, IL, USA). All experimental measurements were conducted in at least triplicate, and the data were expressed as the mean \pm standard deviation (SD). Differences between samples were evaluated by using an analysis of variance (ANOVA) and Duncan's multiple comparison method. Results were considered statistically significant if $p < 0.05$.

3. Results

3.1. Inhibition of Cannabisin B and crude extract on HepG2 cells

In our previous study, we confirmed that cannabisin B from hempseed hull extract possesses predominantly antioxidant activities (Chen et al., 2012). In this report, cannabisin B was separated and purified by solvent extraction, macroporous resin absorption,

LH-20 gel chromatography, and HPLC (Fig. 1) in order to be evaluated for antiproliferative activity in cancer cells.

The inhibitory effects of cannabisin B and crude extract from resin column purification on HepG2 cells are shown in Fig. 2A. Each sample significantly inhibited the growth of HepG2 cells in a concentration-dependent manner, with the exception being cannabisin B treatment for 24 h. Cannabisin B also exhibited a time-dependent antiproliferative activity when the concentration was greater than 251 μ M. Although the crude extract exhibited a stronger inhibition activity, the treated cells shape indicated it might be attributed to necrosis (data not showed).

Visible light pictures of the control cells and the cells treated with different doses of cannabisin B were taken to assess cell morphology (Fig. 2B). The cells shrunk and became round upon treatment with cannabisin B, and the samples that received 503 μ M cannabisin B exhibited obvious degradation compared with the control cells. These changes in cell morphology occurred in a dose-dependent manner. The results indicated that both cannabisin B and crude extract can inhibit the growth of HepG2 cells and that cannabisin B exerted its effects in a dose- and time-dependent manner.

3.2. Cannabisin B-induced S phase arrest in HepG2 cells

The effect of cannabisin B on cell cycle profile was analyzed by flow cytometry. The distribution of cells in different phases of the cell cycle is shown in Fig. 3. A significant and dose-dependent accumulation of cells was observed in the S phase, increasing from 21% in the control cells to 43.84% in the cells treated with 503 μ M cannabisin B. No obvious change was observed in the G2/M phase. The results indicated that cannabisin B could arrest HepG2 cells in S phase and subsequently block cell growth.

3.3. Cannabisin B can not induce apoptosis in HepG2 Cells

Apoptosis is a programmed death process that is defined by a number of morphological and biochemical changes, including cell shrinkage, chromatin condensation, DNA fragmentation, and the

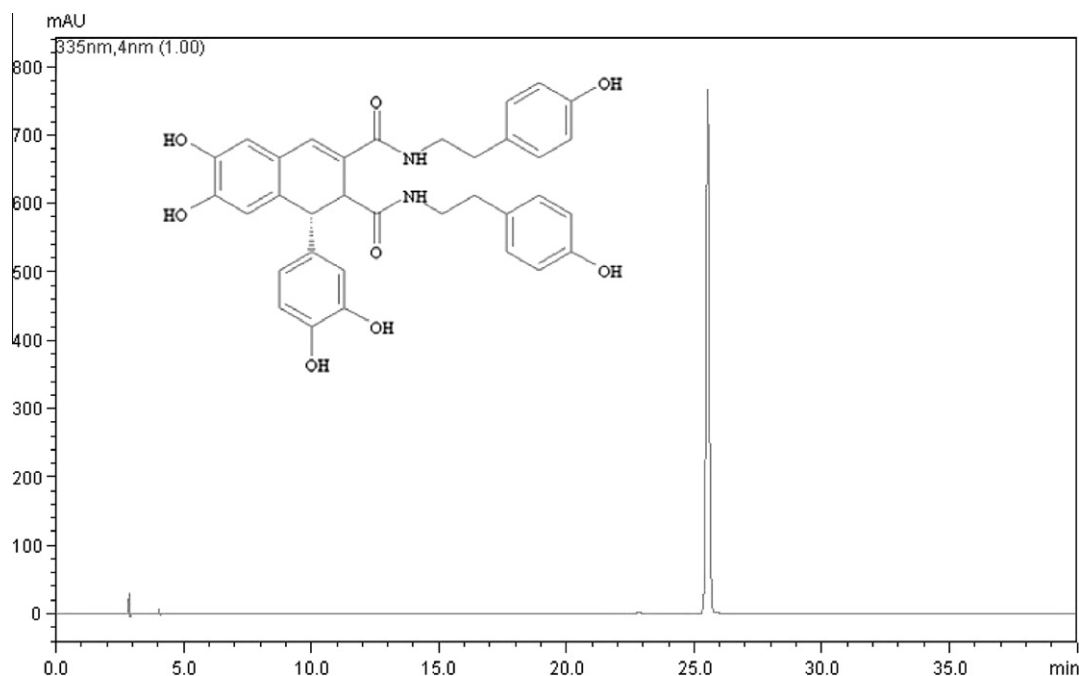


Fig. 1. The structure and HPLC chromatogram of cannabisin B.

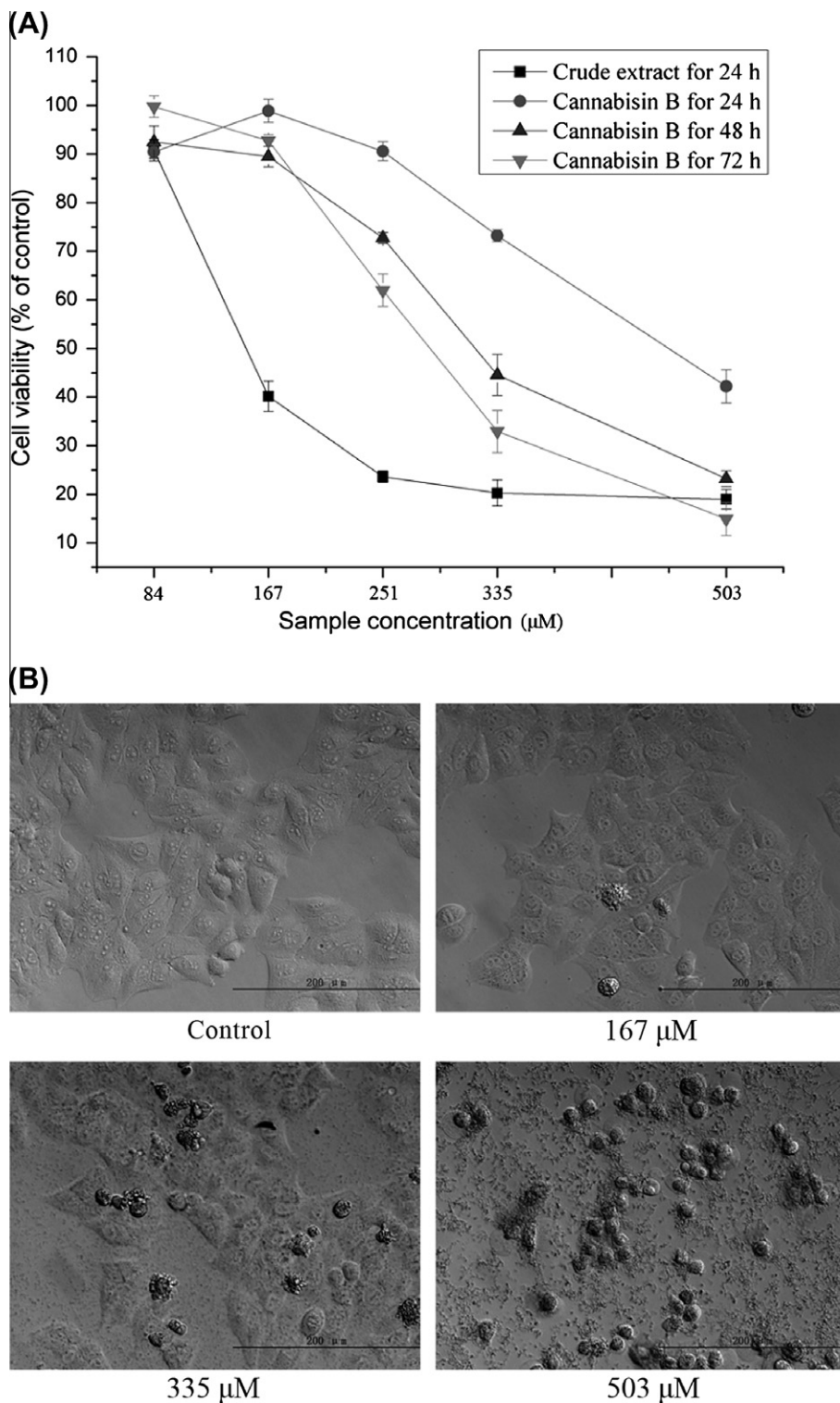


Fig. 2. The antiproliferative activity of cannabisisin B in HepG2 cells. (A) Viabilities of cells treated with various concentrations of crude extract or purified cannabisisin B for various lengths of time. (B) Images of cells treated with different concentrations of cannabisisin B for 24 h.

appearance of phosphatidylserine on the outer side of the plasma membrane (Wu et al., 2004). We previously observed rounding and shrinking in cells treated with cannabisisin B and consequently carried out a series of assays to detect apoptosis. Cells stained with Hoechst 33258 were photographed using a fluorescence microscope. As shown in Fig. 4A, the number of condensed nuclei in the cells treated with cannabisisin B was greater than that of the

control cells, and increased with higher doses of cannabisisin B. Although nuclear condensation is a phenomenon that is always associated with apoptotic cells, it is not a definitive indicator of apoptosis.

Flow cytometric analysis of cells stained with annexin V-FITC and propidium iodide (AP) can be used to determine the level of phosphatidylserine turnover from the inner to the outer lipid layer

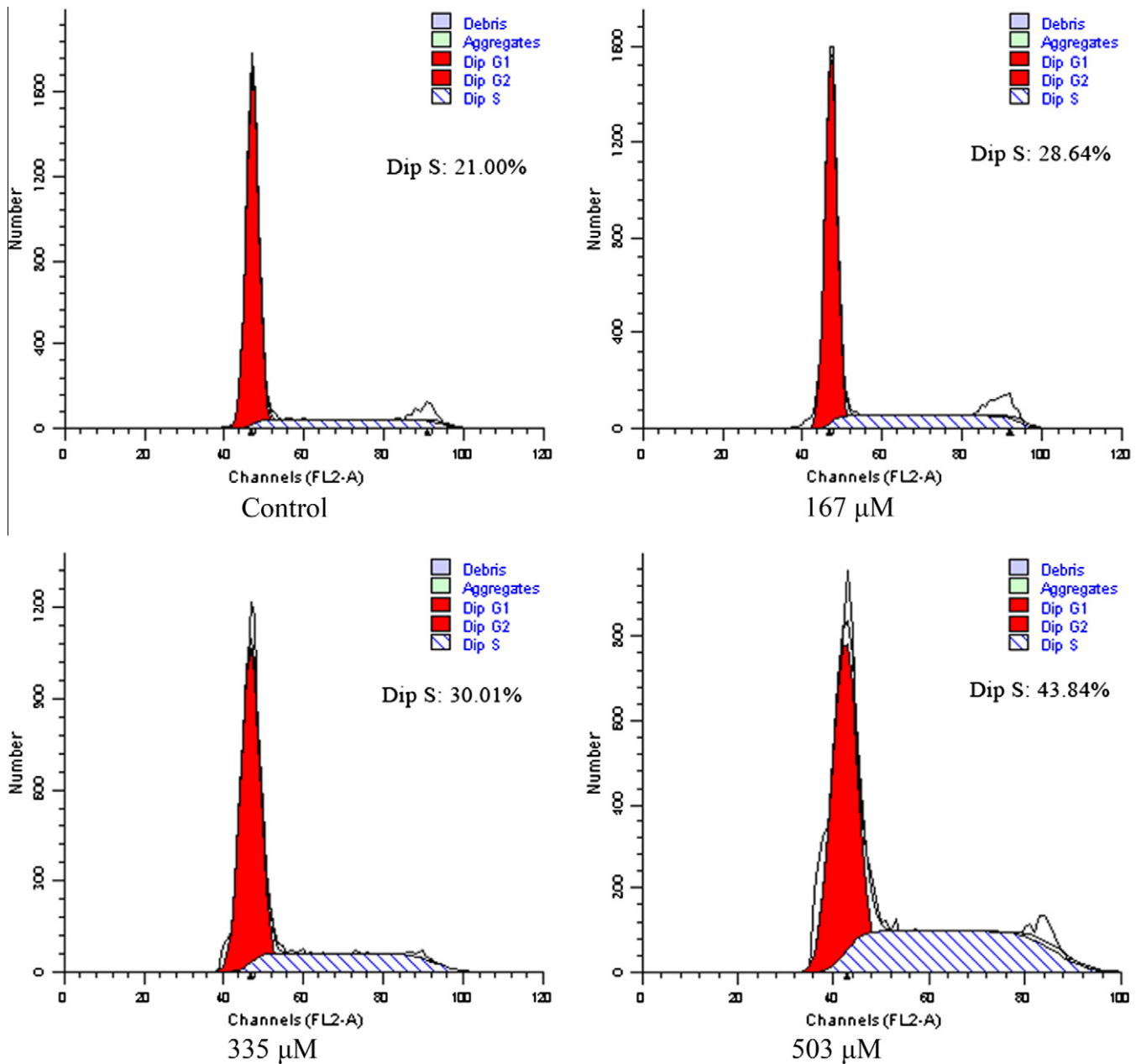


Fig. 3. The effect of cannabidiol on HepG2 cell cycle arrest, as determined by flow cytometry.

of the plasma membrane, which indicates an early stage of apoptosis. The cells that appear in the lower right quadrant are always regarded as apoptotic cells. Fig. 4B showed that there was only a small percentage (0.3–3.31%) of treated cells that were defined as apoptotic cells, and that the frequency of this population remained unchanged at higher doses of cannabidiol B. The results indicated that cannabidiol B (167–503 μM) could not induce typical apoptosis in HepG2 cells.

DNA fragmentation, a well-known marker of apoptosis, is indicated by the production of a DNA ladder pattern in an agarose gel following electrophoresis. As shown in Fig. 4C, we did not observe DNA fragmentation in the control samples or the samples treated with different doses of cannabidiol B. Though nuclear condensation was observed using the Hoechst 33258 staining assay, apoptosis could not be verified by flow cytometry or DNA ladder detection. The results suggested that the antiproliferative activity of cannabidiol B on HepG2 cells cannot be attributed to apoptosis.

3.4. A transient increase in Cannabidiol B-induced cell viability results from autophagy

We unexpectedly observed that the cells treated with low concentrations (84–167 μM) of cannabidiol B exhibited a transient increase in viability compared with the control cells (Fig. 2A), and this increase was not the result of necrosis. Non-apoptotic programmed cell death is predominantly attributed to autophagy (type II programmed cell death), which is a series of biochemical steps through which eukaryotic cells commit suicide by degrading their own cytoplasm and organelles (Reggiori & Klionsky, 2002). 3-MA, an inhibitor of autophagosomes, is commonly used to block autophagy. Fig. 5A showed that 3-MA could significantly inhibit the increase in viability induced by 167 μM cannabidiol B; moreover, 3-MA decreased the death rate of cells treated with 335 and 503 μM cannabidiol B. The cells treated with 3-MA are resistant to the antiproliferative activity of cannabidiol B.

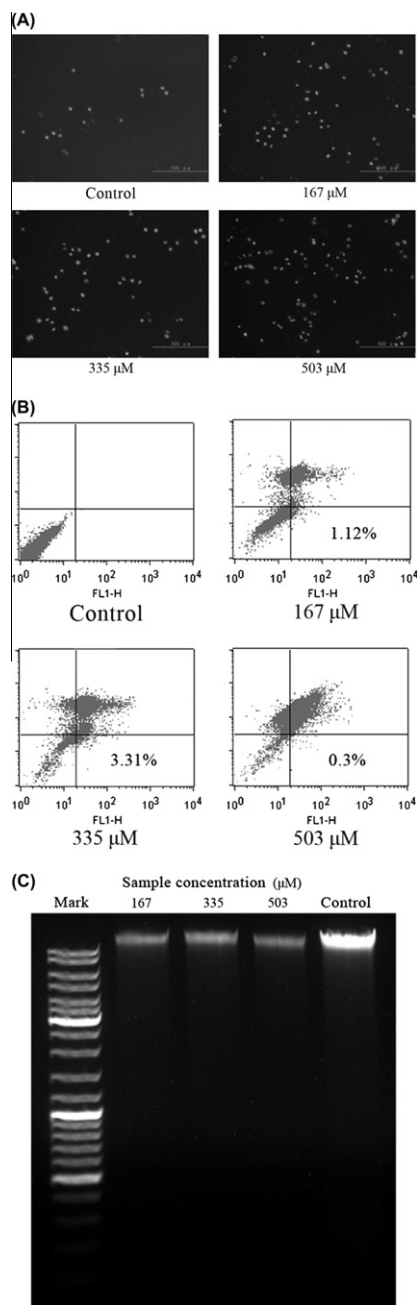


Fig. 4. The detection of apoptosis in HepG2 cells treated with cannabidiol. (A) Fluorescence microscopy was used to capture images of treated cells stained with Hoechst 33258. (B) The flow cytometry plots of cells stained with annexin V-FITC and propidium iodide. (C) DNA fragmentation was determined by agarose gel electrophoresis.

LC3, a group of proteins localized within autophagosome membranes, is regarded as a marker of autophagy (Mizushima et al., 2001). Two forms of cytoplasmic LC3 proteins exist, LC3-I and LC3-II. The amount of LC3-II correlates with the extent of autophagosome formation (Mizushima, Yoshimori, & Levine, 2010). Western blot analysis (Fig. 5B) indicated that LC3 was expressed in cells treated with cannabidiol alone as well as in cells treated with 3-MA and cannabidiol together. Increased LC3-II expression was only observed in cells treated with 167 μM cannabidiol, which also exhibited increased viability compared with the control cells (Fig. 5A).

To determine whether the increase in viability resulted from autophagy induced by cannabidiol, the treated cells were stained

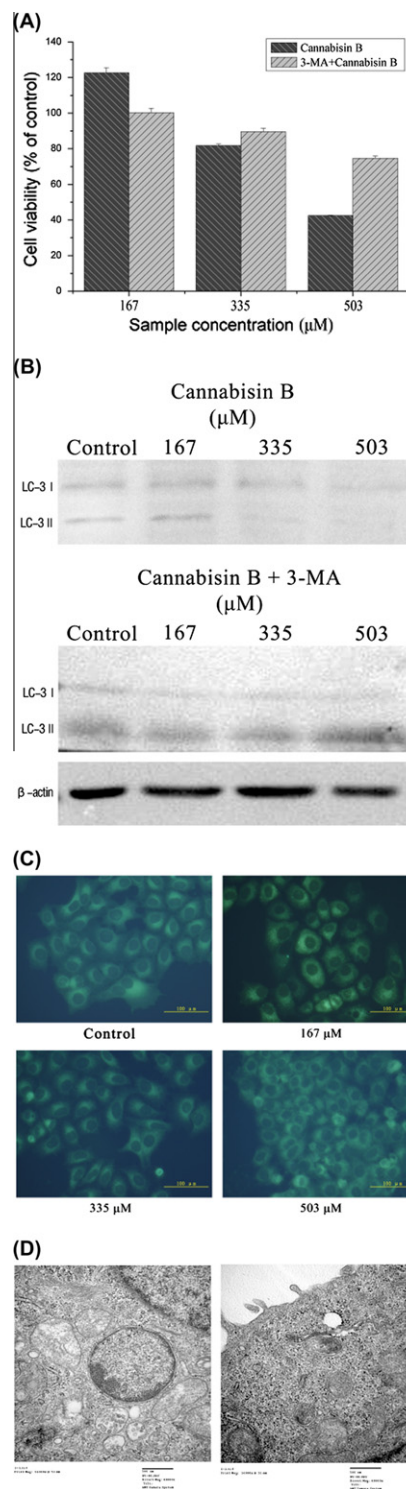


Fig. 5. Cannabidiol induces autophagy in HepG2 cells. Viability (A) and LC3 expression (B) of cells treated with cannabidiol alone or cells pre-treated with 3-MA and then treated with cannabidiol. (C) Fluorescence microscopy was used to capture images of treated cells stained with MDC. (D) The autophagocytic vacuoles examined by transmission electron microscopy.

with MDC and analyzed using a fluorescent microscope. As MDC accumulates in the mature autophagic vacuolar organelle (AVO), it produces a light spot in the cytoplasm that is indicative of autophagy. As shown in Fig. 5C, the cells treated with 167 μM cannabidiol exhibited interspersed staining of AVO, but no significant staining was observed in cells treated with 335 and 503 μM

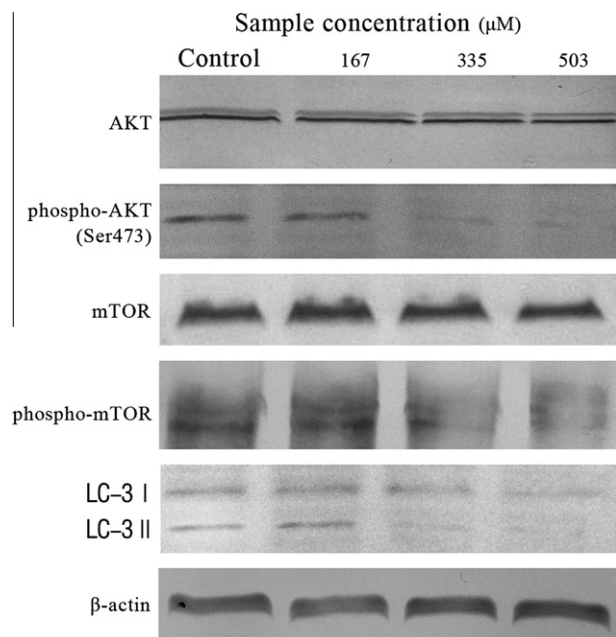


Fig. 6. The effect of cannabidiol on the AKT/mTOR pathway. Total AKT and mTOR, as well as the phosphorylated forms of these proteins, were examined by western blot analysis.

cannabidiol. The existence of an autophagosome was further confirmed by transmission electron microscope (TEM), which is regarded as a gold standard assay for identifying autophagy. In Fig. 5D, cells treated with 167 μM cannabidiol exhibit many membrane-bound vesicles that contain organelles and cellular fragments. Collectively, the LC3 expression levels, MDC staining, and TEM demonstrated that cannabidiol transiently promoted the proliferation of HepG2 cells at 167 μM and then elicited autophagic death in a dose-dependent manner.

3.5. Cannabidiol inhibited the AKT/mTOR pathway

The AKT/mTOR signaling pathway was investigated to determine whether it was involved in cannabidiol-mediated autophagic cell death. Fig. 6 shows the expression levels of AKT, mTOR, and their phosphorylated (activated) forms in cells. Cannabidiol had no impact on total AKT protein but significantly suppressed AKT phosphorylation (Ser 473) in a concentration-dependent manner. Additionally, exposure of HepG2 cells to cannabidiol reduced the levels of the phosphorylated (activated) form of mTOR, a downstream target of AKT, which may have inhibited cell growth and induced autophagy. No effect was observed on total mTOR expression. The results suggested that the autophagy induced by cannabidiol was regulated through the AKT/mTOR pathway.

4. Discussion

Cannabidiol is an unusual polyphenol that is found in hempseed, but it has not been identified in any other plants thus far. Cannabidiol possesses considerable antioxidant capacity because of its poly-phenolhydroxyl structure, which was discovered in our previous study (Chen et al., 2012). Numerous studies have demonstrated that polyphenols isolated from plants possess both antioxidant activities and anticancer activities (Alesiani et al., 2010; Carvalho et al., 2010; Kim et al., 2010). In this report, the anticancer properties of cannabidiol were confirmed for the first time, and the results indicate that cannabidiol and the crude extract of hempseed hull can both significantly inhibit the growth

of HepG2 cells in a dose-dependent manner. However, the crude extract presents a likely necrosis effect against tumor cells, probably because of other cytotoxic compounds in the extract. Moreover, under this experimental conditions, cannabidiol can arrest cells in the S phase of the cell cycle and induce autophagic cell death, but not typical apoptosis, in HepG2 cells.

Autophagy is often induced upon treatment with various chemotherapeutic agents and is considered an alternative way to kill tumor cells (Coward et al., 2009; Gao et al., 2008). Autophagy begins with the sequestration of cytosolic components, which are ultimately engulfed and digested within double-membraned vacuoles called autophagosomes (Todde et al., 2009). An interesting observation was made during this study: a low concentration of cannabidiol transiently increased cell viability whereas higher concentrations of cannabidiol induced cell death. This observation was ultimately attributed to autophagy after confirming AVO formation by MDC staining, detecting autophagosomes with TEM, and determining that the LC3 marker of autophagy was over-expressed in the cells. Furthermore, 3-MA, an autophagy inhibitor, is able to slow the decline in viability rather than increase or decrease viability sharply. In contrast, the common indicators of apoptosis, such as DNA fragmentation and flow cytometric detection of annexin V-FITC staining, are either nonsignificant or slightly significant and therefore do not support the occurrence of apoptosis in cannabidiol-treated HepG2 cells.

Cell cycle arrest is often observed during programmed cell death. Many reports show that apoptosis always occurs simultaneously with cell cycle arrest in the G2/M phase via interfering with the particular cyclin involved (Lee et al., 2004; Zhao et al., 2011). However, the results of this study indicate that cannabidiol can arrest cells in the S phase of the cell cycle in a dose-dependent manner while simultaneously blocking cell growth. The probable explanation may be that cannabidiol impacts DNA synthesis and/or a cyclin that regulates S phase. Although a positive relationship between S phase arrest and the level of autophagy was identified in this experiment, the mechanism behind this relationship has not been determined and should be investigated in future studies.

Recent studies have confirmed that the inhibition of the AKT/mTOR pathway is correlated with triggering autophagy in cancer cells (Kim et al., 2006). Activated AKT, which provides key signals to downstream molecules such as mTOR, can inhibit autophagy and promote cell survival (Yap et al., 2008). Our results show that cannabidiol can suppress the phosphorylation of AKT at Ser 473 (the activated form) and consequently inactivate phosphorylated mTOR, though it has no effect on total AKT or mTOR. These findings indicate that cannabidiol induces autophagic cell death by inhibiting the AKT/mTOR pathway.

In summary, the present study demonstrates (a) that cannabidiol, a bioactive compound from hempseed hull, possesses antiproliferative activity in human hepatocarcinoma HepG2 cells; (b) that cannabidiol arrests cells in the S phase in a dose-dependent manner; and finally, (c) that cannabidiol induces autophagic cell death through the regulation of the AKT/mTOR pathway. The above findings *in vitro* provide inspiring information of cannabidiol to be utilized against liver tumor. In order to clarify the bioactivity, future research will need to focus on investigating the antiproliferative effects of cannabidiol on other cancer cells as well as the activities *in vivo*.

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