Cnidium officinale Makino extract induces apoptosis through activation of caspase-3 and p53 in human liver cancer HepG2 cells

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Received August 17, 2016; Accepted July 21, 2017

DOI: 10.3892/etm.2017.4916

Abstract. A number of diverse studies have reported the anticancer properties of Cnidium officinale Makino (CO). However, the apoptotic effect of this traditional medicinal herb in human hepatocellular carcinoma cells (HepG2) remains to be elucidated. Therefore, the present study investigated the ability of CO to reduce cell viability through apoptotic pathways. Cell viability was determined using the 2,3-bis[2-methoxy-4-nitro-5-sulfophenyl]-2H-tetrazolium-5-carboxanilide assay. CO extract-induced apoptosis in HepG2 cells was assessed by Hoechst 33258 staining. The cell cycle was monitored using fluorescence-activated cell sorting analysis with propidium iodide staining. Furthermore, the present study explored whether various signaling molecules associated with HepG2 cell death were affected by CO treatment, including caspase-3, B-cell lymphoma 2 (Bcl-2), tumor protein p53 (p53), cyclin-dependent kinase 4 (CDK4) and cyclin D. The expression levels of these genes were examined by reverse-transcription polymerase chain reaction and western blotting. The expression levels of caspase-3 and p53 were upregulated with CO extract treatment, whereas those of Bcl-2, CDK4 and cyclin D were significantly downregulated. Cleaved caspase-3 expression was upregulated following treatment with CO extract in a dose-dependent manner. Collectively, the data suggest that CO extract has the potential to induce apoptosis of HepG2 cells and may act by suppressing the cell cycle, which leads to caspase-3 cleavage and p53 signaling.

Introduction

Hepatocellular carcinoma is a well-known malignant tumor whose incidence and mortality rate are increasing worldwide (1). Multiple genetic and epigenetic modifications play an important role in this cancer (2), and some viral infections have been reported to promote liver cancer progression by disrupting the normal cellular mechanisms of apoptosis (3).

Apoptosis is programmed cell death, and it has been shown to play an important role in preventing cancer by repairing damaged DNA in response to mutation (4,5). Induction of apoptosis is a chemopreventive process that involves activation of the p53-dependent apoptotic cell death pathway (6-8). The tumor suppressor gene p53 is also a transcription factor that regulates cell cycle and DNA damage-induced cell mutation through the activation of caspase, a cysteine protease. Activation of anti-apoptotic proteins, such as B-cell lymphoma (Bcl-2), conversely, can delimit apoptosis reduction and deactivation of pro-apoptotic factors such as Bcl-2-associated X protein (Bax) (9).

Natural compounds have an abundance of polyphenols and can be used as chemoprevention agents against malignant cancers in humans (10). Cnidium officinale Makino (CO) has been used historically in traditional Eastern medicine to enhance stamina and manage pain. Owing to the pharmacological properties associated with the abundant polyphenol composition of CO, it has been suggested as a potential herbal treatment for use in metabolic diseases and various forms of cancer (11-14).

However, the efficacy of CO in the treatment or prevention of cancer has not been fully established. In this study, we investigated the methanol extract of CO as a new alternative treatment against liver cancer. Our study demonstrates the potent inhibitory effects of the extract against proliferative liver cancer via downregulation of the cell cycle and induction of apoptosis. Our results suggest that CO may have therapeutic applications in diverse forms of liver diseases.

Materials and methods

Materials and reagents. The plant extraction was performed as previously described by Jung et al (15) with a few modifications. CO was obtained from the Dongguk University Oriental
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Hospital (Korea). The roots of CO (50 g) were blended, and the crude powder was precipitated with 3,000 ml of methanol (80%) at 37°C for 3 days. The methanol extracts were concentrated using a rotary evaporator at 60°C under vacuum conditions. The extract was dissolved in 50 ml of sterile deionized water. The aqueous solution was then lyophilized by freeze-drying at -60°C. The cell culture materials were purchased from Thermo Fisher Scientific (Boston, MA, USA). Antibodies and other laboratory reagents were purchased from Cell Signaling Technology Inc. (Beverly, MA, USA) and Sigma-Aldrich (St. Louis, MO, USA), respectively.

Cell culture. Human hepatoma cells (HepG2) were purchased at the Korean Cell Line Bank (Seoul, Korea). The Chang liver cells were obtained from Seoul National University (Seoul, Korea). HepG2 cells and the Chang liver cells were cultured in Dulbecco's modified Eagle's medium (DMEM) containing 10% fetal bovine serum (FBS) and 1% penicillin-streptomycin. The cells were maintained under an atmosphere of 95% air/5% CO₂ at 37°C. These cells were trypsinized and maintained every 1-2 days.

Determination of cell viability. The effects of different concentrations of CO (100-1,000 µg/ml) on viability of the Chang liver cells and HepG2 cells were investigated by the 2,3-bis [2-methyloxy-4-nitro-5-sulphophenyl]-2H-tetrazolium-5-carboxanilide (XTT) assay (EZ-Cytox cell viability assay kit; Daeil Lab Service, Seoul, Korea), respectively. Briefly, 5x10⁴ cells were seeded in a 96-well microplate and incubated in DMEM under 5% CO₂ at 37°C for 24 h. Then the cells were treated with different concentrations of CO (dissolved in DMEM) for 24 h. To measure cell viability, 10 µL EZ-Cytox was added to each well and the absorbance was measured at 460 nm using a microplate spectrophotometer (Molecular Devices, Sunnyvale, CA, USA). The viability of CO-treated cells was expressed as a percentage of the control cell viability.

Apoptotic analysis with Hoechst 33258 staining. CO extract-induced apoptosis in HepG2 cells was assessed by Hoechst 33258 staining. The cells were seeded at a density of 5x10⁴ cells/well in a 96-well culture plate and were incubated in DMEM at 37°C in 5% CO₂ for 12 h. Then the cells were pretreated with CO extract (0, 100, 250, 500, 750 and 1,000 µg/ml) and incubated again for 24 h. Following treatment, the media were removed and stained with 1 µM Hoechst 33258 staining at room temperature for 60 min. After which, all the solutions were removed and washed twice with 100 µl PBS, and observed by the Cytation 3 Cell Imaging Multi-Mode reader (BioTek Instruments, Inc., Winooski, VT, USA).

Cell cycle analysis. Flow cytometry analysis was used to measure the proportion of HepG2 cells in the different stages of the cell cycle. HepG2 cells were seeded in 6-well plates at 1x10⁶ cells/ml and incubated for 24 h. The cells were incubated with CO extract (0-1,000 µg/ml) for 24 h, and were then harvested and washed twice with phosphate-buffered saline (PBS). Each sample was fixed in 1 ml of 70% ethanol for 2 h at -20°C and the samples were centrifuged. The collected cells were resuspended in PBS containing 50 µg/ml propidium iodide (PI) and 100 µg/ml RNase A and incubated in the dark for 30 min at room temperature. Cell cycle was analyzed using the BD FACSCalibur flow cytometer (BD Biosciences, Franklin Lakes, NJ, USA). The data were analyzed using the BD CellQuest Pro software.

Reverse transcription-polymerase chain reaction (RT-PCR). Total RNA was isolated from HepG2 cells pre-treated with varying concentrations of the extract using TRI-reagent (MRC, Cincinnati, OH, USA) according to the manufacturer's instructions. Briefly, cDNA was synthesized using 0.5 µg of total RNA and the Improm II reverse transcription system with oligo-deoxymyridine (oligo-dT) primers (Promega, Madison, WI, USA). The following semi-quantitative reverse transcription-polymerase chain reaction (RT-PCR) conditions were used for amplification; initial denaturation at 95°C for 10 min and 32 cycles of denaturation at 94°C for 1 min, annealing at 60°C for 1 min and extension at 72°C for 1 min. The following primers were used for RT-PCR; caspase-3, sense primer, 5'-TTT TTC AGA GGG GAT CGT TG-3'; and antisense primer, 5'-CGG TTA ACC CCG GTA AGA AT-3'; Bcl-2, sense primer, 5'-CTG CGA AGA ACC TTG TGT GA-3'; and antisense primer, 5'-TGG TCC TAC CAA CCA GAA GG-3'; p53, sense primer, 5'-GCT CTG ACT GTA CCA CCA TCC-3'; and antisense primer, 5'-CTC TCG GAA CAT CTC GAA GGG-3'; CDK4, sense primer, 5'-ATG GCTGCCACTCGATATGACCC-3'; and antisense primer, 5'-GTACCAGACGCTTAACCACACAGG-3'; cyclin D, sense primer, 5'-AGA CCT GCC CGC CCT CGG TG-3'; and antisenprimer, 5'-GTA GTA GGA CAG GAA GTT GTT C-3'; and GAPDH, sense primer, 5'-CCA GAT CCC TTC AAA ATC AA-3'; and antisense primer, 5'-AGG TCC ACC ACT GAC ACG TT-3'. Transcripts were quantified using electrophoresis on 2% agarose gels. All assays were performed in triplicate.

Western blotting. The lysates from HepG2 cells were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) according to previously described protocols (16). Supernatant protein concentrations were determined using the Bio-Rad DC protein assay reagents (Bio-Rad, Hercules, CA, USA). The proteins were transferred to polyvinylidene difluoride (PVDF) membranes (Amersham Pharmacia Biotech, Buckinghamshire, UK), which were blocked overnight at 4°C in Tris-buffered saline containing 0.1% Tween-20 and 5% bovine serum albumin (BSA). The membranes were then incubated overnight at 4°C with specific antibodies diluted at 1:1,000. Immune complexes were further incubated with their respective peroxidase-conjugated secondary antibodies diluted at 1:1,000 for 1 h. The membranes were analyzed using chemiluminescent reactions (ECL Plus kit; Amersham Pharmacia Biotech), after which protein expressions were visualized and analyzed using ImageJ software (National Institutes of Health, Bethesda, MD, USA).

Determination of standard components by high-performance liquid chromatography (HPLC). The methanol extract of CO was analyzed using an HPLC system (Agilent 1100 series;
Agilent Technologies, Santa Clara, CA, USA) equipped with a Phenomenex column (C18, 4.06_250 mm). An isocratic mixture of CH$_3$CN:MeOH:H$_2$O (60:30:10) was used as the mobile phase at a flow rate of 1.0 ml/min. Absorbance was measured at 210 nm. Standard stock solutions of cnidilide, ligustilide, neocnidilide, butylphalide, senkyunolide, tetramethylpyrazine, caffeic acid, ferulic acid, and perlolyrine were prepared by dissolving 1 mg/ml of analytical standard in methanol and stored at 4˚C prior to use.

Statistical analysis. The results were expressed as mean ± standard deviation (SD) of three independent experiments. Statistical differences among means were calculated by using ANOVA (one-way) followed by Duncan’s multiple range test. The results of cell cycle analysis were tested using Pearson's Chi-square test. Differences with P<0.05 were considered significant. The statistical software package SPSS 10.0 (SPSS Institute, Chicago, IL, USA) was used for these analyses.

Results

Determination of active components in CO. The composition of CO extract was analyzed by determining the presence of nine standard compounds, including cnidilide, ligustilide, neocnidilide, butylphalide, senkyunolide, tetramethylpyrazine, caffeic acid, ferulic acid, and perlolyrine, by HPLC. Ferulic acid (1), cnidilide (2), senkyunolide (3), and ligustilide (4) were identified from CO extract.

Cell viability. The effect of CO on the Chang liver cell viability was determined by XTT assay. The results indicated that CO was not cytotoxic at concentrations between 100-1,000 µg/ml (Fig. 2A). The viability of HepG2 cells treated for 24 h with increasing concentrations of CO extract was assessed. As shown in Fig. 2B, HepG2 cell viability significantly decreased after treatment with CO extract at 100, 250, 500, 750, and 1,000 µg/ml.

Effects of CO on apoptotic cell morphology. In order to investigate whether CO has an apoptotic effects, HepG2 cells treated with CO extract were stained by Hoechst 33258 staining and identified cells undergoing apoptosis by cell imaging reader. As shown in Fig. 3, the marked cells (yellow color) indicate the apoptotic cells. Apoptotic cells in the group treated with CO extract were increased compared with the untreated group in a dose-dependent manner.

Inhibitory effect of CO on cell cycle. To further examine the apoptotic characteristics in HepG2 cells treated with CO extract, the cell cycle phases were analyzed. As shown in Fig. 4, the percentage of S phase cells reduced to 3.5, 4.4, 5.9, 7.3, and 9.9% at 24 h following treatment with 100, 250, 500, 750, and 1,000 µg/ml CO extract, respectively but did not show any significant difference.

Effect of CO on the expression of cell cycle- and apoptosis-related genes in HepG2 cells. As shown in Fig. 5,
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Figure 2. Effects of varying concentrations of Cnidium officinale Makino (CO) extract on the viability of the Chang liver cells and human hepatocellular carcinoma (HepG2) cells. (A) Chang liver and (B) HepG2 cells were treated with 100-1,000 µg/ml CO extract for 24 h and cell viability was assessed using the EZ-Cytox cell viability assay kit. Data are expressed as means ± standard deviation (SD) of three independent experiments. Untreated Chang liver cells and HepG2 cells were considered 100% viable. Values with different superscripted letters are significantly different at P<0.05 by Duncan's multiple range test.

Figure 3. Effects of Cnidium officinale Makino (CO) extract on apoptotic morphology in HepG2 cells. HepG2 cells were seeded and incubated for 12 h. Then, cells were treated with different concentrations of CO (0-1,000 µg/ml) for 24 h and stained with Hoechst 33258 staining. (A) The yellow cells mean apoptotic cells. Apoptotic cells automatically showed yellow and were counted by the machine program. (B) Data are mean ± SD (n=3), values with different superscripted letters are significantly different at P<0.05 by Duncan's multiple range test.

Figure 4. Effects of Cnidium officinale Makino (CO) extract on HepG2 cell cycle progression. The cells were incubated with CO extract (0-1,000 µg/ml) for 24 h. Cell cycles were determined by fluorescence-activated cell sorting (FACS) analysis. Each item was derived from representative experiments, where data was obtained from at least 10,000 events. (A) The cell populations in G0/G1, S, and G2/M phases were determined using the BD CellQuest Pro software. (B) The graph indicates the fluorescence intensity of incorporated propidium iodide (PI, %) in each phase of cell cycle.
mRNA levels of caspase-3 increased to 11.4, 34.5, 49.4, 80.0, and 84.7% and those of p53 to 17.1, 39.4, 51.0, 63.5, and 70.7%, respectively, after treatment with 100, 250, 500, 750, and 1,000 µg/ml CO. Conversely, the same concentrations of CO decreased the levels of Bcl-2 by 9.2, 20.0, 47.6, 62.0, and 64.7%, respectively. In addition, CDK4 expression decreased after treatment with the same concentrations of CO to 14.1, 23.9, 66.0, 87.1, and 86.4%, respectively. In addition, CDK4 expression decreased after treatment with the same concentrations of the CO extract to 14.1, 23.9, 66.0, 87.1, and 86.4%, respectively, and those of cyclin D decreased by 13.6, 24.2, 64.2, 92.8, and 92.0%, respectively.

**Effect of CO on the expression of cell cycle- and apoptosis-related proteins in HepG2 cells.** We analyzed the expression of cell cycle- and apoptosis-related proteins to investigate the molecular mechanism by which CO extract triggered cell cycle regulation and apoptosis in HepG2 cells. The protein levels of caspase-3 and p53 increased after CO treatment in a dose-dependent manner. In contrast, those of Bcl-2 and cell cycle enzymes decreased in a dose-dependent manner after treatment with the same concentrations of CO extract (Fig. 6).

**Effect of CO on cleaved caspase-3 in HepG2 cells.** The expression levels of caspase-3 in HepG2 cells were determined following treatment with CO extract using western blot analysis. As shown in Fig. 7, the percentage of cleaved caspase-3 increased to 32.4, 64.3, 93.5, 127.0, and 133.5% following a 24-h treatment with CO extract at concentrations of 100, 250, 500, 750, and 1,000 µg/ml, respectively.

**Discussion**

Apoptosis is regarded as an important cellular mechanism that occurs in response to abnormalities and cell damage (6,9). Anomalous apoptotic responses are prominent in the development and progression of many types of human cancers (17,18). One of the chemopreventive strategies in the treatment of cancer is the induction of apoptotic cell death (7). Natural compounds are rich sources of anticancer agents that can induce apoptosis and DNA repair in response to mutations (10). Previous studies have shown that essential oils and compounds from medicinal herbs can be used to treat human diseases owing to their potent biological activities against abnormal physiological reactions in the body (19-22). Our research confirmed that the major components of CO, including ferulic acid, cnidilide, senkeyunolide, and ligustilide, induced apoptotic cell death in HepG2 cells. It has been established that rhizomes of CO contain 1-2% of essential oils and active compounds, such as cnidilide, ligustilide, neocnidilide, butylphalide, senkeyunolide, tetramethylpyrazine, caffeic acid, ferulic acid, and perlollyrine (23). In addition, several recent reports have highlighted...
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Figure 6. Effects of Cnidium officinale Makino (CO) extract on caspase-3, Bcl-2, p53, CDK4, and cyclin D protein expressions in HepG2 cells. HepG2 cells were seeded in 6-well plates and incubated in the presence or absence of CO extract (100-1,000 µg/ml) for 24 h. (A) The expression levels of caspase-3, Bcl-2, p53, CDK4, and cyclin D were determined by western blot analysis. (B-F) The graph shows the densitometry scanning analysis of the ratios of caspase-3, Bcl-2, p53, CDK4, and cyclin D. Untreated HepG2 cells were considered to have 100% viability. Data are mean ± SD (n=3), values with different superscripted letters are significantly different at P<0.05 by Duncan's multiple range test.

Figure 7. Effects of Cnidium officinale Makino (CO) extract on caspase-3 in HepG2 cells. HepG2 cells were seeded and incubated in the presence or absence of CO (100-1,000 µg/ml) for 24 h. (A) Apoptosis regulatory proteins, such as caspase-3 and cleaved caspase-3, were assayed by western blot analysis. (B) Densitometric quantification of data, which are expressed as means ± SD of three independent experiments. The viability of untreated HepG2 cell was considered 100%. Values with different superscripted letters are significantly different at P<0.05 by Duncan's multiple range test.

the anti-inflammatory and anticancer effects of CO (14,24). Lee demonstrated that ferulic acid, a phenolic compound, induced apoptosis in HepG2 cells through the generation of reactive oxygen species (ROS) (25). Kan et al also confirmed the anticancer effects of senkyunolide A and Z-ligustilide in colon cancer (26).

Cell growth is regulated by CDK and its binding proteins, such as the cyclins. In G1 phase, CDK4 and cyclin D enzymes are known to initiate cell division (27). Interestingly, Musgrove et al claimed that cyclin D is one of the therapeutic target molecules in cancer therapy (28). Cyclin D 1 plays an important role in the progression of G1 to S phase of the cell cycle in normal cells through the activation of CDK4 (14). In this study, treatment with the methanol extract of CO significantly reduced HepG2 cell viability. In addition, the Hoechst 33258 staining demonstrated that apoptosis in HepG2 cell were increased with the increasing CO extract concentrations. We confirmed G0/G1 arrest and reduction of cell population in S phase via the suppression of CDK4 and cyclin D expression in the CO-treated HepG2 cells.

The p53 tumor suppressor gene is a pivotal molecule mediating cell cycle arrest and apoptosis (29). Furthermore, this regulation subsequently inhibits the activity of cell cycle-regulating enzymes and the expression of the anti-apoptotic protein Bcl-2 (8). Reduction in the expression of Bcl-2 signaling molecules has been reported as anticancer mechanisms in diverse types of cancers (4). RT-PCR and western blot data showed that CO extract induced the activation of the tumor
Makino extract mediated through apoptosis


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vation of p53 and reduction of Bcl-2. Therefore, CO is effectively induce apoptosis of HepG2 cells through acti

p53-dependent signaling pathways that regulate caspase-3 expression (31). Our data also revealed that CO extract activated the p53 gene, causing a cascade of events leading to the activation of caspase-3 and finally cell death.

Taken together, we suggest that CO extract may effectively induce apoptosis of HepG2 cells through activation of p53 and reduction of Bcl-2. Therefore, CO is a promising and potential source of polyphenol compounds that may prove effective as chemoprevention agents for hepatocellular carcinoma.

References


