RESEARCH ARTICLE

Dentatin from *Clausena excavata* Induces Apoptosis in HepG2 Cells via Mitochondrial Mediated Signaling

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Abstract

Hepatocellular carcinoma (HCC) is a primary liver cancer with high global incidence and mortality rates. Current candidate drugs to treat HCC remain lacking and those in use possess undesirable side effects. In this investigation, the antiproliferative effects of dentatin (DTN), a natural coumarin, were evaluated on HepG2 cells and DTN's probable preliminary molecular mechanisms in apoptosis induction were further investigated. DTN significantly (p<0.05) suppressed proliferation of HepG2 cells with an IC₅₀ value of 12.0 μ g/mL, without affecting human normal liver cells, WRL-68 (IC₅₀> 50 μ g/mL) causing G₀/G₁ cell cycle arrest via apoptosis induction. Caspase colorimetric assays showed markedly increased levels of caspase-3 and caspase-9 activities throughout the treatment period. Western blotting of treated HepG2 cells revealed inhibition of NF- \varkappa B that triggers the mitochondrial-mediated apoptotic signaling pathway by up-regulating cytoplasmic cytochrome c and Bax, and down-regulating Bcl-2 and Bcl-xL. The current findings suggest DTN has the potential to be developed further as an anticancer compound targeting human HCC.

Keywords: Dentatin - HepG2 - apoptosis - mitochondrial-mediated signaling pathway

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Introduction

Hepatocellular carcinoma (HCC), known as malignant hepatoma is a primary cancer of the liver. HCC is the third most common cause of cancer-related deaths globally after lung and stomach cancer. As per GLOBOCAN report, in 2008, 749,000 recent cases of HCC were identified globally and its related deaths were recognized to be 695,000 accounting for the high incidence to mortality ratio (Ferlay et al., 2013). Initially, the focus of HCC treatment was on systemic, nonspecific, high-dose chemotherapy, but currently there is a need to invent a drug that balances minimal adverse effects with maximal antitumor activity (Abou-Jawde et al., 2003).

Breakthroughs in plant-derived anticancer agents had begun to show significant growth in drug discovery (Sultana et al., 2014). Analyses that include cytotoxicity testing provide crucial means of ranking compounds for consideration in anticancer drug discovery whereas, multiparametric *in vitro* analysis, which evaluates apoptosis induction is able to provide important data regarding undesirable side-effects and further to conclude that the compound under investigation is tumor specific, non-redundant, and able to hinder tumor progression (Niles et al., 2008; Zhang et al., 2014). *Clausena excavata* Burm.f., a wild shrub, is extensively distributed throughout Southeast Asia belonging to the family of Rutaceae and has been used as folk medicine for the treatment of several disorders and diseases (Mohan, 2012). Their therapeutic properties have been mainly attributed to the presence of carbazole alkaloids and coumarins (Wu and Furukawa, 1982; Wu et al., 1996; Sunthitikawinsakul et al., 2003; Taufiq-Yap et al., 2007; Kumar et al., 2012).

DTN is known for its cytotoxicity against various cancer cells as previously reported, which included human prostate cancer cells (LNCaP and PC-3) (Arbab et al., 2012) and human breast cancer cells (MCF-7) (Arbab et al., 2013). In this current investigation, the antiproliferative effects of DTN were evaluated on HCC cells, HepG2 and the probable molecular mechanism of DTN that causes cell death in HepG2 cells via apoptosis induction was further investigated.

Materials and Methods

Compound isolation and purification

C. excavata was sourced from Pendang, Kedah in December 2010 and identified at the Herbarium Unit, Institute of Bioscience, University Putra Malaysia (UPM).

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The roots (1.5 kg) of *C. excavata* were air-dried and ground before extraction. DTN was isolated and purified by one of the co-author. The compound was sent for infrared (IR) and nuclear magnetic resonance (NMR) analyses at the Laboratory of Spectroscopic Analysis, Faculty of Science, UPM. The purity (>96.42%) was determined by means of HPLC and LC-MS. Recently, DTN has been isolated and characterized using IR and NMR (Arbab et al., 2012; Sharif et al., 2013). The spectral data were found to be similar as previously reported by Xin and the team (Xin et al., 2008).

Reagents

DTN was dissolved in DMSO and added to the cell culture with a final DMSO concentration of 0.1% w/v. Only analytical grade reagents were used.

Cell culture and maintenance

HepG2 cells and WRL-68 cells were sourced from American Type Culture Collection (ATCC). Culturing of both cells in RPMI 1640 (Nacalai Tesque, Japan) was done, supplemented with 10% Fetal Bovine Serum (Life Technologies, USA) and 1% Antibiotic-Antimycotic (Nacalai Tesque, Japan), maintained at 37°C/5% CO₂.

Cell proliferative assay

Cell proliferation assay was quantified by the ability of living cells to reduce the yellow MTT dye by mitochondrial succinate dehydrogenase, resulting in the production of formazan purple crystals as described by Mosmann, 1983. Briefly, cell suspension of HepG2 (100 μ L/well) was seeded at densities of 2 x 10⁵ cells/mL into 96 well microtiter plates and treated with DTN of different concentrations (100 µg/mL, 50 µg/mL, 25 µg/mL, 12.5 μ g/mL, 6.25 μ g/mL and 3.125 μ g/mL) and followed by incubation in atmosphere of 5% CO₂ at 37°C for 68 h. MTT stock solution (5 mg/mL) at a volume of 10% $(20 \ \mu L)$ of the medium volume was added per well and further incubated at 37°C for 4h, away from light. This was followed by the removal of media and the formazan crystals that resulted were solubilized by adding 100 µL of DMSO with gentle swirling. The absorbance was measured at 595 nm with a microliter plate reader (Tecan Sunrise basic, Groedig, Austria). The viability of the cell was expressed as a percentage of absorbance relative to vehicle-treated cell control. DTN was also tested for its cytotoxicity on WRL-68 cells and paclitaxel was used as positive control.

Morphological assessment via phase contrast inverted microscope

HepG2 cells at 2×10^5 cells/mL density then underwent treatment with DTN (IC₅₀) for 24 h, 48h and 72h. The changes in the morphology of treated cells were compared to untreated control cells observed under Phase Contrast Inverted Microscope equipped with NIS-Elements D software (Nikon).

Confocal microscopic Assessment using AO/PI Double Staining

Briefly, HepG2 cells were seeded at a density of 2×10^5

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cells/mL and treated with DTN (IC₅₀) for 24h, 48h and 72 h. Cells not treated served as negative control. This was followed by the addition of 10 μ L (10 μ L) of fluorescent dyes, (AO/PI) with acridine orange (AO), (10 μ g/mL) and propidium iodide (PI), (10 μ g/mL) into the harvested cell pellet in equal volumes of each. Appropriate volume of freshly stained cell suspension was dropped onto a glass slide and covered by coverslip. The coverslip was then sealed with nail polish. Concurrently, the coverslip was observed under the Confocal Laser Scanning Microscope equipped with Qfluoro software (Zeiss, Germany) within 30 minutes before the fluorescent color started to fade.

Annexin V-FITC/PI assay

Loss of membrane asymmetry was detected using Annexin V-FITC Kit according to manufacturer's protocol (MACS Miltenyi Biotech, USA). HepG2 cells at a density of 1 x 10⁶ cells/mL were treated with DTN (IC₅₀) in time-dependent manner (6h, 12h, 24h, 48h, and 72h). Untreated cells served as the negative control. Harvested cells were washed, resuspended in 1X binding buffer and incubated with Annexin-V FITC and PI prior to the analysis on Fluorescence-activated cell sorting (FACS) by flow cytometer (BD Bioscience, California) equipped with CellQuest Pro software.

Analysis of cellular DNA content

Briefly, HepG2 cells of 2 x 10⁶ cells/mL density were treated with DTN (IC₅₀) for 24h, 48h and 72h with negative control served by untreated cells. Harvested cells were fixed with 70% cold ethanol and stored frozen (-20°C) up to 1 week. Cells were then incubated in staining solution (0.2mg/ml RNAse A and 50 µg/mL PI). Samples were kept in the dark at room temperature for 30 minutes, prior to the analysis on FACS by flow cytometer, equipped with CellQuest Pro software.

Caspase-3/8/9 activity

Caspase activity was determined using Caspase-3/8/9 Colorimetric Assay Kit as per manufacturer's protocol (GenScript, USA). Seeding of HepG2 cells was done at a density of 4 x 10⁶ cells/mL and followed by treatment with DTN (IC₅₀ c for 12h, 24h, 48h and 72h. At the same time, negative control was set. Resuspension of the cells was done in cold lysis buffer followed by incubation on ice for 1 h. After centrifugation, 10 μ L of supernatant was used to assay the protein concentration by Bradford method. Equal amounts of protein were mixed with 2X reaction buffer and 200 μ M DEVD-pNA substrate (caspase-3)/ IETD-pNA substrate (caspase-8)/ LEHD-pNA substrate (caspase-9) and further incubated at 37°C for 4h, away from light. The plates were read at 405 nm.

Western blotting

HepG2 cells underwent treatment with DTN (IC₅₀) for 12h, 24h and 48h. Untreated cells served as the negative control. Protein extraction was done using CytobusterTM Protein Extraction Reagent following manufacturer's protocol (Novagen, USA) and the protein concentrations were quantified by Bradford method. Protein samples (20

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Table 1. Flow Cytometric Analysis of Annexin V-FITC/PI of HepG2 Cells Treated with DTN (IC₅₀) in a Time Dependent Manner

	Control	6 H	12 H	24 H	48 H	72 H
VIABLE	97.19±0.65	95.42±0.81*	93.80±0.5*	84.35±0.15*	70.70±0.20*	42.01±0.21*
EARLY APOPTOSIS	2.27±0.11	3.39±0.26*	3.87±0.17*	10.74±0.14*	16.82±0.44*	20.62±0.13*
LATE APOPTOSIS	0.42±0.11	0.76±0.17	1.80±0.19*	3.42±0.30*	9.42±0.10*	24.52±0.44*
SECONDARYNECROSIS	0.12±0.07	0.43±0.12	0.52±0.20	1.49±0.20*	3.06±0.19*	12.85±0.41*

Table shows percentage of cell count ± SD of three independent experiments, each in triplicate. '' Indicates a significant difference (p<0.05)

 μ g) were separated by SDS-PAGE and transferred to a PVDF membrane using a wet transfer system (Bio-Rad, USA). After blocking with 5% non-fat dry milk and 0.05% Tween-20 in Tris-buffered saline (TBS), the membranes were incubated overnight with the primary antibody [Betaactin (1:1000), Bcl-2 (1:2000), Bax (1:2000), Bcl-xL (1:1000), Cytochrome c (1:1000) and NF-xB (1:1000)] followed by incubation with Goat Anti-Mouse IgG (H+L) Alkaline Phosphatase (AP) Conjugate (Bio-Rad, USA). AP Conjugate Substrate Kit (Bio-Rad, USA) was used for the color development and photographs were taken while the membrane was wet to enhance the purple color. *ImageJ* software was then used to quantify the intensity of the .bands

Statistical analysis

Each experiment was performed in triplicate. Data were presented as mean±SD and analyzed by One Way ANOVA and Tukey's Test using IBM SPSS Statistics 20. Differences with p<0.05 were considered statistically significant.

Results and Discussion

DTN exhibits potent antiproliferative effect on HepG2 and not on WRL-68

DTN was screened for cytotoxicity on several cancer cells, namely HepG2, CEMss, HeLa and HT29, using the MTT colorimetric assay. Results of the assay (data not shown) showed that, DTN exerted highest cytotoxicity effect on HepG2 cells with IC₅₀ value of 12.03 μ g/mL. Treatment of normal human liver cells, WRL-68 cells with DTN showed no evidence of significant inhibition up to 50 μ g/mL after 72 h of incubation. Paclitaxel, used as positive control, exhibited an inhibitory effect on HepG2 cells at IC₅₀ value of 2.32±0.05 μ g/mL after 72 h treatment.

DTN shows distinct cell death related morphological features

Microscopy assessment of cellular morphological changes attributed to DTN treatment in HepG2 cells was made. In respect to this, morphological changes in response to DTN's (IC_{50}) treatment in HepG2 cells at exposure time of 24h, 48h and 72h were compared to untreated HepG2 cells using phase contrast inverted microscope. Figure 1 shows features of cell shrinkage and membrane blebbing following DTN's treatment in a time-dependent manner. In addition to using this evaluation, other microscopic assessment using dyes to detect cellular morphological changes possibly relating to apoptosis induction were later used to provide evidence of



Figure 1. Phase-Contrast Inverted Micrographs of HepG2 Cells Treated with DTN (IC_{50}). a) Control, show an epithelial-like morphology, presented normal, healthy form demonstrated by clear skeletons b) 24h, cells lose contact with adjacent cells, as rapid cell shrinkage occurs concurrently to nuclear changes, forming a blebbing surface on the cell c) 48h, rapid growth inhibition leading to detachment of the cell and membrane blebbing and d) 72h, detachment of most of the cells with apparent cell shrinkage. VI: Viable Cell, MB: Membrane Blebbing, CS: Cell Shrinkage



Figure 2. AO/PI Stained Confocal Micrographs of DTN (IC₅₀) **Treated HepG2 Cells. a)** Control **b**) 24 h, c) 48 h and **d**) 72 h. VI: Viable Cell, MB: Membrane Blebbing, CC: Chromatin Condensation, LA: Late Apoptosis, SN: Secondary Necrosis

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Figure 3. Scatter Profile of Flow Cytometric Analysis of Annexin V-FITC/PI of DTN (IC₅₀) Treated HepG2 Cells in Time Dependent Manner. a) Control, b) 6h, c) 12h, d) 24h, e) 48h and f) 72h. QI: viable cells, QII: early apoptotic cells, QIII: late apoptotic cells, QIV: secondary necrotic cells



Figure 4. Flow Cytometric Analysis of Cellular DNA Content Frequency Histograms of HepG2 Cells Treated with DTN (IC_{50}) in Time Dependent Manner. a) Control, b) 24h, c) 48h and d) 72h. Sub G_0/G_1 denotes apoptotic DNA with hypodiploid DNA content

this phenomenon. Viable, early apoptosis, late apoptosis and secondary necrosis states of cells were determined using acridine orange (AO) and propidium iodide (PI) fluorescent dyes (Foglieni et al., 2001). Following this, confocal microscopic examination of DTN (IC_{50}) treated HepG2 cells in a time-dependent manner after AO/PI staining revealed typical cellular morphological changes of HepG2 cells associated with cell death via probable apoptosis induction (Figure 2). Untreated control showed healthy HepG2 cells possessing weak green nuclei of proliferated uncontrolled growth (Figure 2-a). After 24



Figure 5. Western Blotting Analysis Showing Regulation of Bax, Bcl-2, Bcl-xL, Cytochrome c, NF- \times B and β -actin (Loading Control) protein expression in DTN (IC₅₀) Treated HepG2 Cells after 12h, 24h and 48h with Untreated Control

h treatment, HepG2 cells were bright green with yellow nuclei, denoting evidence of condensed chromatin (Figure 2-b). Early apoptosis and late apoptosis features were observed after respective 48h (Figure 2-c) and 72h treatments (Figure 2-d), which include bright green cells of yellow nucleic, membrane blebbing and orange intact cells. Apoptotic cells going through secondary necrosis were found following 72h treatment where cells displayed reddish orange intact nucleus (Figure 2-d).

DTN induces loss of membrane asymmetry

The key features of loss of membrane asymmetry are the exposure of phosphotidylserine (PS) from the inner face of plasma membrane to the surface of the cell (Van Engeland et al., 1996; Van Engeland et al., 1998). Thus, in order to discriminate the integrity of cell membrane, FITC labelled Annexin V together with a dye exclusion of PI were performed to target and detect early apoptotic cells before a cell loses its membrane integrity (Vermes et al., 1995; Van Engeland et al., 1996). Apoptotic cells were counted as early or late apoptotic cells, which are shown repeatedly in the lower right (Annexin V/FITC+PI-) and upper right (Annexin V/FITC+PI+) quadrants of the scatter profile of DTN (IC50) treated HepG2 cells in increasing treatment time (Figure 3). There was a significant increase (p<0.05) of the apoptotic cells from an untreated control (2.69%), such increase occurring in a time-dependent manner at 4.15%, 5.67%, 14.16%, 26.24% and 45.14% after 6h, 12h, 24h, 48h and 72h treatments respectively.

DTN induces G_d/G_1 cell cycle arrest

DTN induced a concomitant accumulation of cells in the G_0/G_1 phase, with an apparent increase (p<0.05) from untreated HepG2 cells (50.98±0.33%) to treated cells; 53.84±0.32%, 58.17±0.20% and 62.04±0.56% after 24h, 48h and 72h of treatment respectively, with subsequent decreases in S phase (Figure 4). These accumulations could be due to G1 checkpoint that made it impossible for the division of the DTN treated HepG2 cells, which then entered the G0 resting phase, thus signaling the G_0/G_1 phase arrest (Pucci et al., 2000). These cells were possibly eliminated through apoptosis induction (Pucci et al., 2000) as it correlated with a significant (p<0.05) increase of apoptotic cells (sub G0/G1), as treatment time with DTN increased in HepG2 cells.

DTN induces apoptosis via mitochondrial mediated signaling pathway

The Bcl-2 family proteins are implicated in the regulation of the mitochondrial-mediated apoptosis pathway (Shamas-Din et al., 2013; Czabotar et al., 2014; Volkmann et al., 2014). Thus, fluctuations in the pro-apoptotic and anti-apoptotic Bcl-2 proteins levels were investigated in DTN (IC_{50}) treated HepG2 cells. The Western Blotting analysis demonstrated increased pro-apoptotic Bax protein in a time-dependent manner (Figure 5). In contrast, anti-apoptotic proteins of Bcl-2 and Bcl-xL decreased upon DTN treatment (Figure 5).

An important stage in the mitochondrial pathway is the subsequent release of cytochrome c from the intermembrane space (Estaquier et al., 2012; White, 2013). Upon releasing cytochrome c into the cytosol, the permeability of mitochondria is controlled by Bcl-2 proteins by either promoting or inhibiting apoptosis directly on mitochondrial permeability transition pores (Narita et al., 1998). Pro-apoptotic protein, Bax forms pores, while anti-apoptotic protein Bcl-2 and Bcl-xL hinder their formation (Crompton, 1999). Figure 5 shows significant (p<0.05) time-dependent increase of cytochrome c accumulation in the cytosol upon DTN treatment in HepG2 cells.

Transduction of apoptotic signals needs to be activated by a cascade involving caspases. Caspase-3, particularly, has a key role in executing apoptosis as it catalyzes specific cleavage of several important cellular proteins (Brentnall et al., 2013). Increase in caspase-3 activity is crucial for drug-induced apoptosis as the activation of caspase-3 is a hallmark of apoptosis and can be used in quantifying the caspase cascade (Logue and Martin, 2008). Caspase-3 is triggered by either caspase-8 or caspase-9, which acts as a convergence point for different apoptosis signaling pathways. In this current investigation, the activity of caspase-8 remained at basal level while the activity of caspase-9 significantly (p<0.05) increased in a time dependent manner. Similarly, increase in caspase-3 activity was observed in DTN-treated HepG2 cells in a time-dependent manner and this evidently proved the use of the active form of caspase-3 in detection events leading toward apoptotic induction. The current results correlate well with the mitochondrial signaling pathway, and as the cytochrome c is released, it binds to Apaf-1 protein and ATP, which in turn binds to pro-caspase-9, forming an apoptosome (Wurstle et al., 2012). The apoptosome then triggers the pro-caspase-9 to induce the formation of active caspase-9, which simultaneously activates the caspase-3 (Logue and Martin, 2008; White, 2013).

NF- α B is now becoming an interesting therapeutic target for the treatment of cancer. NF- α B, an apoptosis

suppressing protein, stimulates the expression of the IAPs and other proteins from the anti-apoptotic Bcl-2 family (De Almagro and Vucic, 2012; Ouyang et al., 2012). The IAPs constantly suppress apoptosis by direct inhibition of caspases-3, while anti-apoptotic members of the Bcl-2 family counteract the role of the pro-apoptotic members (De Almagro and Vucic, 2012). In this current study, the down-regulated NF-xB (Figure 5) suppresses the expression of IAPs, consequently decreases cell survival and hence, triggers apoptosis. Inhibition of NF-xB will increase the susceptibility to apoptosis, lead to higher cell mortality, in which NF-xB regulates anti-apoptotic genes, particularly the TRAF1 and TRAF2 and, thus, activate the activities of the caspases, which are the central to most of the apoptotic processes (Dolcet et al., 2005; Kim et al., 2006).

In summary, the present study demonstrates the ability of DTN to induce apoptosis against HepG2 cells mediated by mitochondrial signaling pathway and this effect is associated with the inhibition of NF-xB. Further studies will focus on addressing its effects in animal models, which warrants DTN as a probable chemotherapeutic agent to treat human hepatocellular carcinoma.

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