

Cytisine induces endoplasmic reticulum stress caused by calcium overload in HepG2 cells

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Abstract. Cytisine, a quinolizidine alkaloid, is one of the major bioactive components found in the small tree *Sophora Alopecuraides* L., and is a traditional Chinese medicine that is used for treating hepatitis and liver cancer. In the 1960s, quinolizidine alkaloids were reported to exhibit inhibitory effects on tumour cell proliferation in several types of cancer cells. However, few studies have investigated the effect of cytosine on liver cancer. Our team confirmed that cytosine induced apoptosis in HepG2 cells via a mitochondrial pathway. The primary aim of the present study was to evaluate the endoplasmic reticulum (ER) stress caused by calcium overload in cytosine-induced apoptosis in HepG2 cells and the molecular mechanisms of this phenomenon. In addition, the present study was undertaken to evaluate the expression of $\alpha 7$ -nAChR when apoptosis was induced by cytosine in HepG2 cells. In the present study, transmission electron microscopy was used to observe the morphological appearance of HepG2 cells. The apoptosis of the cells with cytoplasmic vacuolization was significant under electron microscopy. Apoptotic bodies, the expansion of the ER,

and swelling of mitochondria were observed in the HepG2 cells after cytosine treatment. Flow cytometric analysis demonstrated that the apoptosis rate of HepG2 cells was upregulated. In addition, the intracellular calcium concentration was detected by laser confocal fluorescence microscopy. The laser confocal fluorescence microscopy showed that the calcium concentration was increased in a dose-dependent manner. The activity of caspase-4 was evaluated by an enzyme-linked analyser, and the expression levels of CHOP, JNK, p-JNK and $\alpha 7$ -nAChR were assessed via western blot analysis. In the present study, we observed that cytosine induced ER stress-inducing factors and CHOP and p-JNK1/2 protein expression, and it increased the JNK protein expression in the HepG2 cells. Furthermore, $\alpha 7$ -nAChR protein expression was promoted in a dose-dependent manner after cytosine treatment. These findings suggest that cytosine induced the ER stress-mediated apoptotic pathway via activation of CHOP, JNK and caspase-4 in HepG2 cells, and cytosine is a potential new target compound for nAChRs (nicotinic acetylcholine receptors) to treat liver cancer.

Introduction

Quinolizidine alkaloids have previously been demonstrated to exhibit antitumour activities *in vitro* and *in vivo* (1,2). Cytisine, a quinolizidine alkaloid from *Sophora Alopecuraides* L., has been widely used for the treatment of central nervous system diseases. It has previously been demonstrated that cytosine may significantly inhibit neuronal apoptosis induced by NMDA exposure by reversing intracellular Ca^{2+} overload and balancing the expression levels of Bcl-2 and Bax (3). Furthermore, several *in vitro* studies have also confirmed the antitumour activities of cytosine. Cytisine was found to inhibit human cancer cell lines, including HepG2, A549, K562, Ec109, HL-60 and U937 cells (4). Our team confirmed the molecular mechanisms underlying cytosine-induced apoptosis of HepG2 via a mitochondrial pathway; however, the specific mechanism underlying endoplasmic reticulum (ER) stress remains to be elucidated.

Apoptosis is a crucial mechanism for cell clearance. It is now accepted that cell apoptosis has mainly three apoptotic

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Abbreviations: PI, propidium iodide; ER, endoplasmic reticulum; nAChRs, nicotinic acetylcholine receptors; UPR, unfolded protein response; EOR, endoplasmic reticulum overload response; HCPT, 10-hydroxycamptothecin; SDS-PAGE, sodium dodecyl sulphate-polyacrylamide gel electrophoresis

Key words: cytosine, HepG2 cells, apoptosis, endoplasmic reticulum stress, calcium overload, CHOP pathway, JNK pathway

signalling pathways, including the mitochondrial, the death receptor- and the ER-associated pathway (5-7). In recent years, researchers have discovered that ER stress-mediated apoptosis is a new signalling pathway for apoptosis (8). The endoplasmic reticulum (ER), a membrane-bound organelle, plays an essential role in eukaryotic cell processes, including synthesizing, sorting, assembling, modifying and trafficking proteins, and maintaining intracellular Ca^{2+} homeostasis (9). The normal functioning of the ER signalling cascades requires high concentrations of free calcium ions within the ER lumen. Impairment of Ca^{2+} homeostasis or other noxious stimuli may trigger an ER stress response that is manifested by either an unfolded protein response (UPR) or an ER overload response (EOR) within the ER lumen (10-13). However, if the stress persists on the ER for a longer period of time, the ER stress triggers cell death (typically apoptosis) mainly through three pathways, including CHOP (14), JNK (15) and caspase-12 (16-18). Although caspase-12 has been implicated in ER stress-induced apoptosis in rodents, it is controversial whether similar mechanisms operate in humans. Recent research has indicated that caspase-4, which is one of the closest paralogs of rodent caspase-12, plays an important role in cell death (19). Thus, the present study investigated the ER stress effects of cytosine on HepG2 cells in relation to CHOP, JNK and caspase-4.

Additional studies have indicated that cytosine can be used as an nAChR partial agonist in clinical use as a smoking cessation aid. Therefore, cytosine is considered to be superior to nicotine-replacement therapy for smoking cessation (20-22). Nicotinic acetylcholine receptors (nAChRs) consist of various types of subunits, such as $\alpha 2$ - $\alpha 10$, $\beta 2$ - $\beta 4$, which are a complex of five subunits forming hetero- or homo-pentamers that form a central ion channel. nAChRs were originally identified to be only expressed in the nervous system and at neuromuscular junctions (muscle type nAChRs). However, the discovery of widespread expression of nAChRs in the cell membrane of all mammalian cells, including cancer cells, suggested its direct role in the development and progression of cancers (23). Therefore, nAChRs are viewed as a novel drug target for the prevention and treatment of various forms of cancers. Recently, increasing evidence has shown that $\alpha 7$ nicotinic receptors exhibit the highest permeability for Ca^{2+} , promoting cell death via a release of Ca^{2+} from intracellular stores and intracellular Ca^{2+} overloading (24,25).

The major aim of the present study was to elucidate the effects of cytosine on ER stress in HepG2 cells in terms of the intracellular calcium concentration and the three pathways involved in ER stress, including CHOP, JNK and caspase-4. Additionally, the present study investigated the effect of cytosine on the expression of nAChR.

Materials and methods

Instruments and reagents. Cytosine (no. 20111101C; Shanxi River Pharmaceutical Co., Ltd., Xi'an, China); human hepatocellular carcinoma cell line, HepG2 (Life Sciences and Environmental Sciences Development Centre of Harbin University of Commerce, Harbin, China); RPMI-1640 medium (no. 20130507; Gibco Co., Carlsbad, CA, USA); fetal calf serum (FCS) (no. 20140206; Zhejiang Tianhang Biotechnology

Co., Ltd., Zhejiang, China); propidium iodide (PI) (no. P4170; Sigma Co., St. Louis, MO, USA); fluorescence microscope (Olympus Corporation, Tokyo, Japan); HCPT (no. H109197; Aladdin Reagent Co. Ltd., Shanghai, China); monoclonal antibody of β -actin (no. 141125), p-JNK (no. A1514) (both from ZSGB-BIO Co., Ltd., Beijing, China); monoclonal antibody of GADD153 (no. 980787W; Beijing Biosynthesis Biotechnology Co., Ltd., Beijing, China), JNK (no. 150116; Wanleibo Co., Ltd., Shanghai, China); caspase-4 test kit (no. 20140526; Beyotime Institute of Biotechnology, Haimen, China); JEM-1220 transmission electron microscope (JEOL Ltd.); EPICS XL-MCL flow cytometry (Beckman Coulter, Inc., Brea, CA, USA); 680 enzyme-linked analyser (Bio-Rad Laboratories, Inc., Hercules, CA, USA); peroxidase-conjugated AffiniPure goat anti-rabbit IgG (H+L; cat. no. ZB-2301) and peroxidase-conjugated AffiniPure goat anti-mouse IgG (H+L; cat. no. ZB-2305) (both from ZSGB-BIO Co., Ltd.).

Effect of cytosine on the morphological appearance of HepG2 cells. Transmission electron microscopy was used to observe the morphological appearance of the HepG2 cells after drug treatment (26). Cells were gently harvested after trypsin digestion and washed twice with PBS. Approximately 3×10^5 cells were treated in single wells of 6-well plates. The cultured HepG2 cells were exposed to cytosine at concentrations of 2.5, 5 and 10 mM, or they were exposed to HCPT at concentrations of 60 μM for 24 h at 37°C with 5% CO_2 . Since cytosine and HCPT both belong to the same family of alkaloids, HCPT was used as the positive drug in the present study to verify that the experimental method was correct. A blank control group was added with the same volume of RPMI-1640 culture medium, and the cells were collected after 24 h. Next, 2% glutaraldehyde was used to fix the cells for >2 h, and they were immobilized with osmium acid. After alcohol gradient dehydration, all groups were treated with epoxy resin embedding, ultrathin sectioning, uranyl acetate and lead citric acid double staining and then imaged under the transmission electron microscope.

Apoptosis assays. Quantitative analysis of the apoptosis was evaluated via flow cytometry with PI staining according to prior guidelines (27). Cells were gently harvested after trypsin digestion and washed twice with PBS. The cells were then centrifuged at 1,000 rpm for 10 min. Cells (1×10^6) were treated approximately into a single cell suspension with PBS solution and fixed with 70% ice-cold ethanol at 4°C overnight. Afterwards, the cells were washed twice with PBS and stained with a solution containing 800 μl of PI (PI staining contains sodium citrate 33.4 mg, PI 5 mg, RNase A 1 mg, joined Triton X-100 0.5 ml) for 30 min in the dark at room temperature. A minimum of 10,000 cells were maintained for all of the samples. The cells were later analysed for their DNA content using flow cytometry. By analysing the DNA histograms, the percentage of the cells in different cell cycle phases was evaluated using MlytiCycle for Windows 32-bit software (Beckman Coulter, Inc.). Finally, the cells with sub- G_0/G_1 DNA (sub- G_1) were calculated as apoptotic cells. Apoptosis was recorded using the laser line of 488/630 nm.

Confocal laser scanning microscope (CLSM) analysis. The accumulation of Ca^{2+} ions in the cytoplasm was studied using

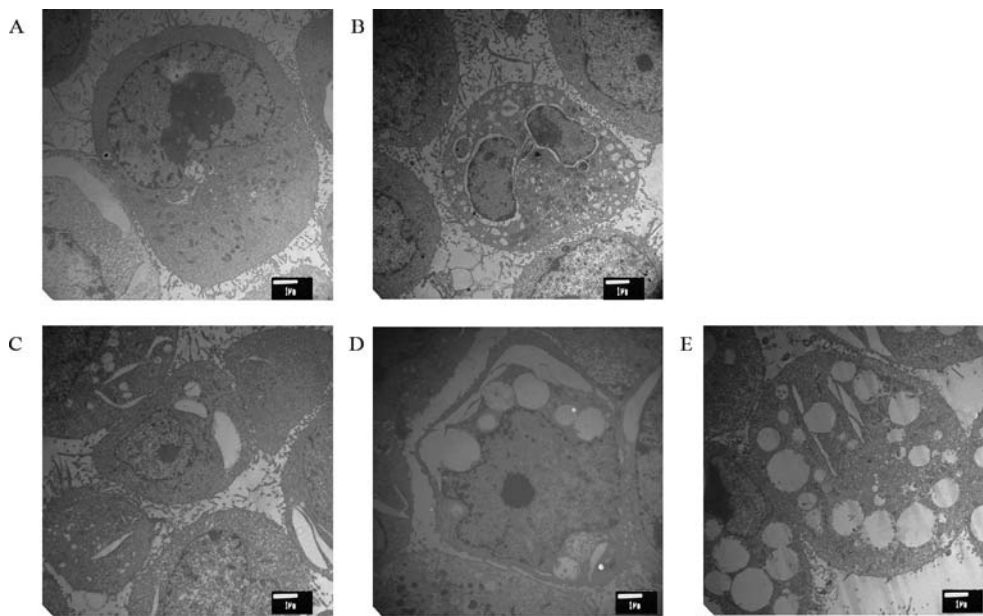


Figure 1. Effect of cytosine on the morphological appearance of HepG2 cells. In all groups, cells were observed by a transmission electron microscope. (A) HepG2 cells treated with RPMI-1640 medium. (B) HepG2 cells treated with 60 μ M HCPT. (C) HepG 2 cells treated with 2.5 mM cytosine. (D) HepG2 cells treated with 5 mM cytosine. (E) HepG2 cells treated with 10 mM cytosine.

the fluorescent dye, Fluo3 AM (28). Approximately, 3×10^5 cells were seeded per well in 6-well plates and incubated for 24 h before treatment with various concentrations of cytosine and a further 24 h of incubation at 37°C with 5% CO₂. Cytosine at concentrations of 2.5, 5 and 10 mM or HCPT at a concentration of 60 μ M was added, and the plate was incubated at 37°C in 5% CO₂. The HepG2 cells were resuspended with PBS at 1,500 rpm for 10 min after trypsin digestion. After fixation, the cells were incubated with 200 μ l of Fluo3 AM (4 μ g/ml) to detect the intracellular Ca²⁺ at 37°C at ~30 min. Next, the cells were washed twice with PBS. The fluorescence emitted by the Fluo3 AM when bound to Ca²⁺ ions was recorded using the laser line of 488/570 nm.

Western blot analysis. Cells were harvested after the treatment with test drugs for 24 h. After the indicated treatments, the HepG2 cells were harvested and lysed with a cell lysis solution at 4°C for 30 min. Whole cellular proteins were extracted and cytosolic fractions were prepared following the procedure described by the manufacturer. Total proteins were quantitatively assayed using the BCA method. A loading buffer was added to the cytosolic extracts, and it was boiled for 10 min. Afterwards, 50 μ g of protein was loaded onto a 15% sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) gel and run at 80 V for 30 min and 120 V for 1 h. Proteins were transferred onto nitrocellulose membranes. Afterwards, incubation was performed for 1 h in a blocking solution (5% non-fat dry milk in 20 mM of TBS with 0.1% Tween) at room temperature. Next, the membranes were gently washed with diluted TBS-T and subsequently incubated for 24 h with the primary antibody of GADD153 (1:500), JNK (1:500), p-JNK (1:500) and α 7-nAChR (1:500) at 4°C, respectively. The secondary antibody was added at 1:2,000 dilutions and incubated at room temperature for 2 h. Cell membranes were removed and coloured with DAB agent (29,30). Membranes

were examined via chemiluminescence detection using a photographic film. The image was captured using a Tannon gel imaging system, and the hybrid band was quantitatively analysed with Gel-Pro Analyzer 3.1 density analysis software.

Caspase activity assay. The enzymatic activity of caspase-4 was determined using a caspase colourimetric assay kit according to the manufacturer's protocol. Briefly, the cells were lysed in a lysis buffer for 15 min in an ice bath. The lysed cells were centrifuged at 20,000 \times g for 10 min, and 50 μ l of the protein was incubated with 40 μ l of a reaction buffer and 10 μ l of the colourimetric tetrapeptides, Ac-DEVD-pNA for caspase-4 at 37°C for 1 h. The optical density of the reaction mixture was quantified spectrophotometrically at a wavelength of 405 nm. The caspase enzymatic activity in the cell lysate is directly proportional to the colour reaction.

Statistical analysis. Differences in proliferation between different cell lines were analysed using one-way ANOVA. Statistical analysis was performed using SPSS 19.0 software. (SPSS, Inc., Chicago, IL, USA). $P < 0.01$ was considered to be a statistically significant difference.

Results

Effect of cytosine on the morphological appearance of HepG2 cells. The effect of cytosine on the ultra-structure of the HepG2 cells is shown in Fig. 1. Cells from the control group had the following characteristics: a clear structure, complete cell structure, a small number of microvilli on the cell surface, large nuclei, abundant euchromatin and obvious nucleolus. In addition, we also observed many free ribosomes, and the rough ER and mitochondria were scattered in the cytoplasm. In the positive control group, the mitochondria were reduced, and the cell processes were obviously decreased. The ER swelled

Table I. Apoptosis rate of HepG2 cells following treatment with cytosine as detected by flow cytometry (mean \pm SD, n=3).

Groups	Concentration	Number of cells	Apoptosis rate (%)
Control	-	1x10 ⁶	0.97 \pm 0.29
HCPT	60 μ mol/l	1x10 ⁶	37.79 \pm 1.55 ^a
Cytosine	2.5 mmol/l	1x10 ⁶	10.83 \pm 0.45 ^a
	5 mmol/l	1x10 ⁶	16.98 \pm 1.42 ^a
	10 mmol/l	1x10 ⁶	34.28 \pm 1.11 ^a

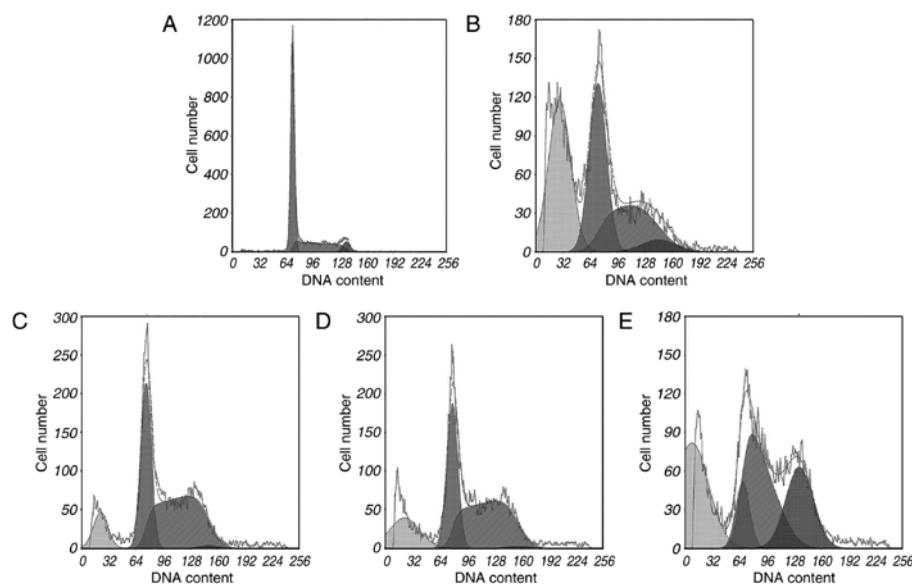
^aCompared with control P<0.01.

Figure 2. Apoptosis rate of HepG2 cells by flow cytometry (FCM). Cells were treated with cytosine at concentrations of 2.5, 5 and 10 mM or HCPT at a concentration of 60 μ M for 24 h. The cells were then stained with propidium iodide and analysed using an EPICS XL-MCL flow cytometer (Beckman Coulter). (A) HepG2 cells treated with RPMI-1640. (B) HepG2 cells treated with 60 μ M HCPT. (C) HepG2 cells treated with 2.5 mM cytosine. (D) HepG2 cells treated with 5 mM cytosine. (E) HepG2 cells treated with 10 mM cytosine.

and expanded, and typical apoptotic bodies appeared. The cell surface microvilli decreased or disappeared after different concentrations of cytosine. Under electron microscopy, the apoptosis of the cells with cytoplasmic vacuolization was found to be significant. Retraction of the cytoplasm, chromatin condensation, expansion of the perinuclear space, rupture of the membrane, expansion of the ER, and swelling of the mitochondria were observed in HepG2 cells after drug treatment.

Apoptosis analysis. During apoptosis, activation of certain nucleases results in DNA degradation. The sub-G₁ method relies on the fact that after DNA fragmentation, there are small fragments of DNA capable of being eluted following washing in either PBS or a specific phosphate-citrate buffer. This finding means that after staining with a quantitative DNA-binding dye, cells that have lost DNA may take up less stain and may appear to the left of the G₁ peak. Rates of apoptosis were analysed using PI staining and flow cytometric analysis. The FCM results indicated that cytosine was capable of inducing apoptosis at concentrations of 2.5, 5 and 10 mM (Table I). In particular, HCPT induced a higher percentage (37.79 \pm 1.55%) of apoptosis at 60 mM compared to the control group (P<0.01).

Table II. Effects of cytosine on the concentration of [Ca²⁺]_i in the HepG2 cells (mean \pm SD, n=3).

Groups	Concentration	Variation in [Ca ²⁺] _i (fluorescent intensity)
Control	-	7.39 \pm 0.17
HCPT	60 μ mol/l	27.17 \pm 0.76 ^a
Cytosine	2.5 mmol/l	17.98 \pm 0.56 ^a
	5 mmol/l	21.54 \pm 0.26 ^a
	10 mmol/l	30.69 \pm 0.80 ^a

^aCompared with control, P<0.01.

As shown in Fig. 2, treatment with cytosine increased the percentage of cells in the sub-G₁ phase (P<0.01). The preincubation of HepG2 cells with cytosine (2.5, 5 and 10 mM) significantly increased the sub-G₁ cell population (P<0.01). The data demonstrated (Table I and Fig. 2) that cytosine increased HepG2 cell apoptosis rates in a dose-dependent manner.

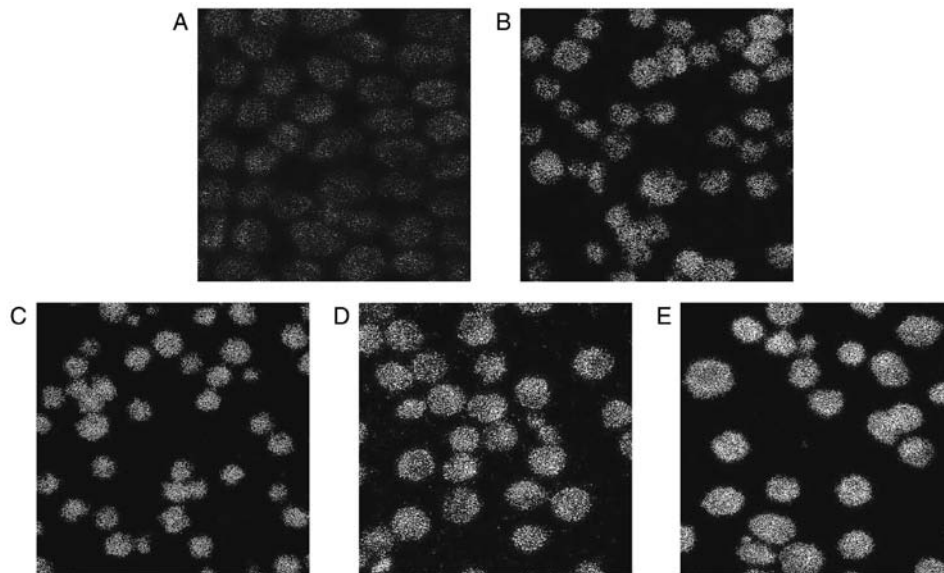


Figure 3. Concentration of $[Ca^{2+}]_i$ in HepG2 cells by laser confocal fluorescence microscopy. In all groups, cells were stained with Fluo3 AM and observed by laser confocal fluorescence microscopy. (A) HepG2 cells treated with RPMI-1640. (B) HepG2 cells treated with 2.5 mM cytosine. (C) HepG2 cells treated with 5 mM cytosine. (D) HepG2 cells treated with 10 mM cytosine. (E) HepG2 cells treated with 60 μ M HCPT.

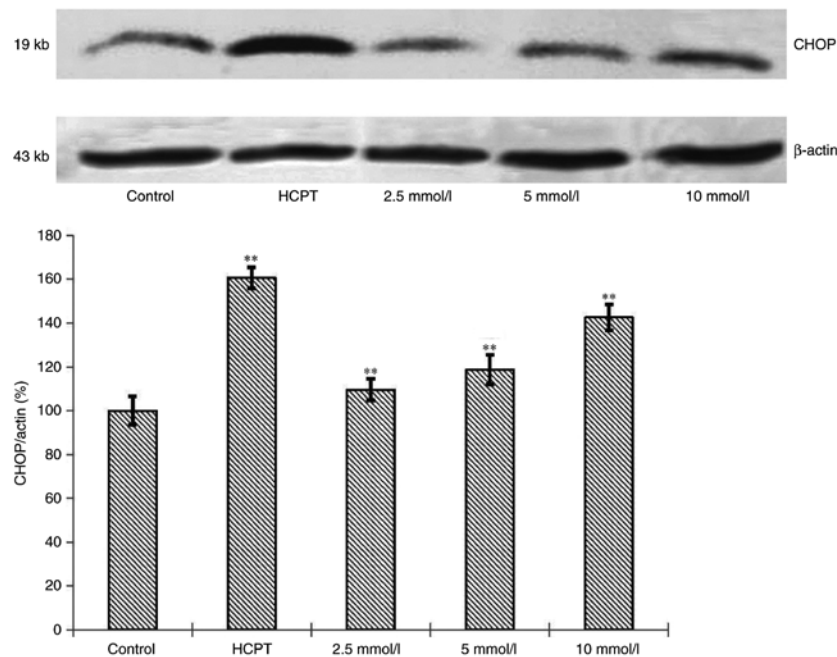


Figure 4. Effect of cytosine on the expression level of CHOP protein in HepG2 cells. The CHOP level in HepG2 cells was determined after 24 h of incubation with or without cytosine treatment and HCPT by western blotting. Each blot is a representative of three similar experiments. **Compared with control, $P < 0.01$.

Laser confocal fluorescence microscopy analysis. To determine the contribution of Ca^{2+} released from intracellular stores to the cytosine-increased $[Ca^{2+}]_i$ content, the HepG2 cells were treated with cytosine for 24 h. Our results clearly indicated that cytosine caused an increase in the cytoplasmic Ca^{2+} levels in a dose-dependent manner (Table II). Cytosine (10 mM) significantly increased the cytosine-increased $[Ca^{2+}]_i$ content by 30.69 ± 0.80 , and cytosine (2.5 mM) slightly increased the cytosine-increased $[Ca^{2+}]_i$ content by 17.98 ± 0.56 ($P < 0.01$). The green fluorescence intensity increased with the increase in drug concentration, as shown in Fig. 3.

Effects of cytosine on the expression of CHOP protein. To confirm our observation that cytosine induced ER stress, we performed immunoblotting analyses on ER-regulated protein, CHOP. CHOP level in the HepG2 cells in the cytosine group was increased compared with the normal group. Significant differences were noted ($P < 0.01$, Fig. 4).

Effects of cytosine on the expression of JNK protein. Western blot analysis revealed that exposure of the HepG2 cells to cytosine for 24 h induced a slight upregulation of expression of the p-JNK1 and p-JNK2 proteins ($P < 0.01$, Fig. 5). In contrast,

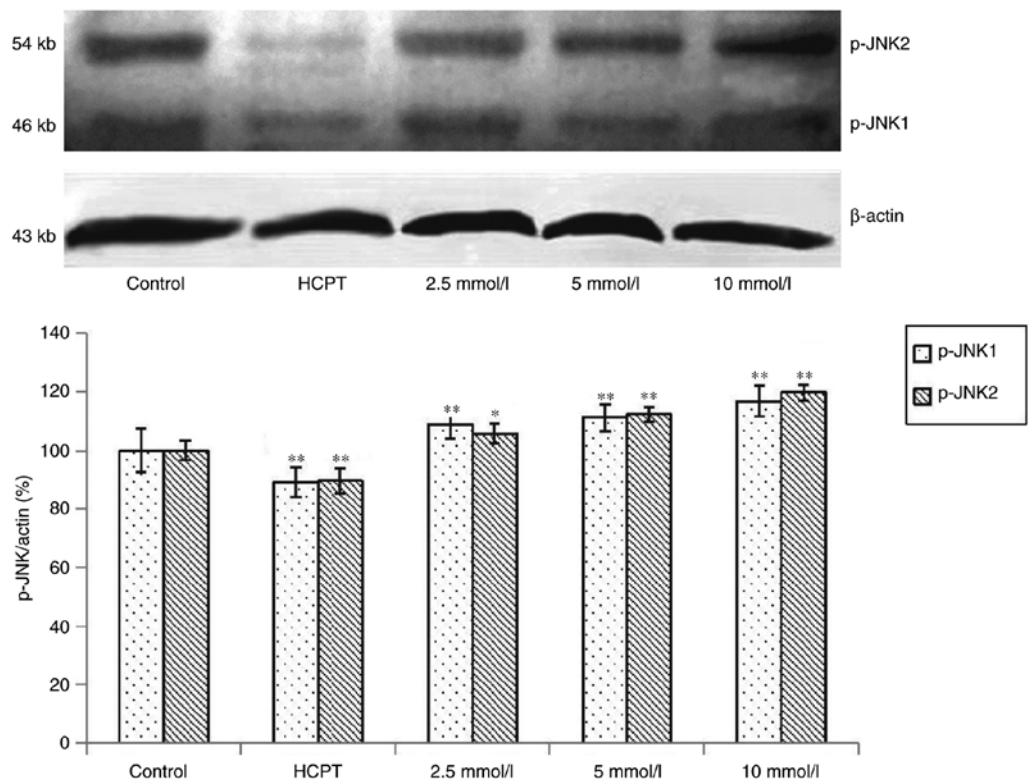


Figure 5. Effect of cytosine on the expression level of p-JNK protein in HepG2 cells. Expression levels of p-JNK1 and p-JNK2 proteins. After treatment, the fractions were resolved by SDS-PAGE, transferred onto nitrocellulose membranes, and probed with primary antibody for p-JNK and secondary antibody for IgG (H+L). The amount of β -actin was measured as an internal control. *Compared with control, P<0.05; **compared with control, P<0.01.

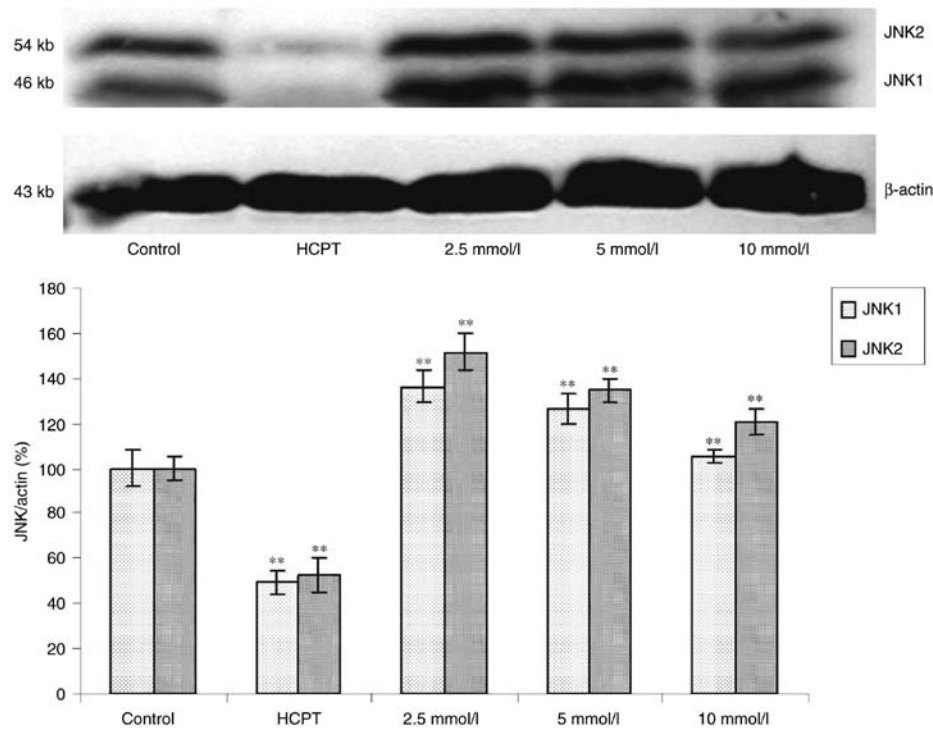


Figure 6. Effect of cytosine on the expression level of JNK protein in HepG2 cells. Expression levels of JNK1/2 proteins. After treatment, the fractions were resolved by SDS-PAGE, transferred onto nitrocellulose membranes, and probed with the primary antibody for JNK and secondary antibody for IgG (H+L). The amount of β -actin was measured as an internal control. **Compared with control, P<0.01.

the expression levels of JNK1 and JNK2 were significantly increased in the HepG2 cells in a dose-dependent manner (P<0.01, Fig. 6).

Effect of cytosine on the activation of caspase-4. As shown in Table III, after treatment of the HepG2 cells with cytosine for 24 h, the activities of caspase-4 were significantly increased

Table III. Effects of cytosine on caspase-4 activity in the HepG2 cells (mean \pm SD, n=3).

Groups	Concentration	OD value ($\bar{x} \pm s$)	pNA (μ M)	Caspase-4 activity (%)
Control	-	0.1093 \pm 0.006	29.25	-
HCPT	60 μ mol/l	0.3177 \pm 0.008 ^a	81.35	178.12
Cytosine	2.5 mmol/l	0.1500 \pm 0.325 ^a	39.43	34.79
	5 mmol/l	0.2337 \pm 0.007 ^a	60.35	106.32
	10 mmol/l	0.3247 \pm 0.007 ^a	83.10	184.10

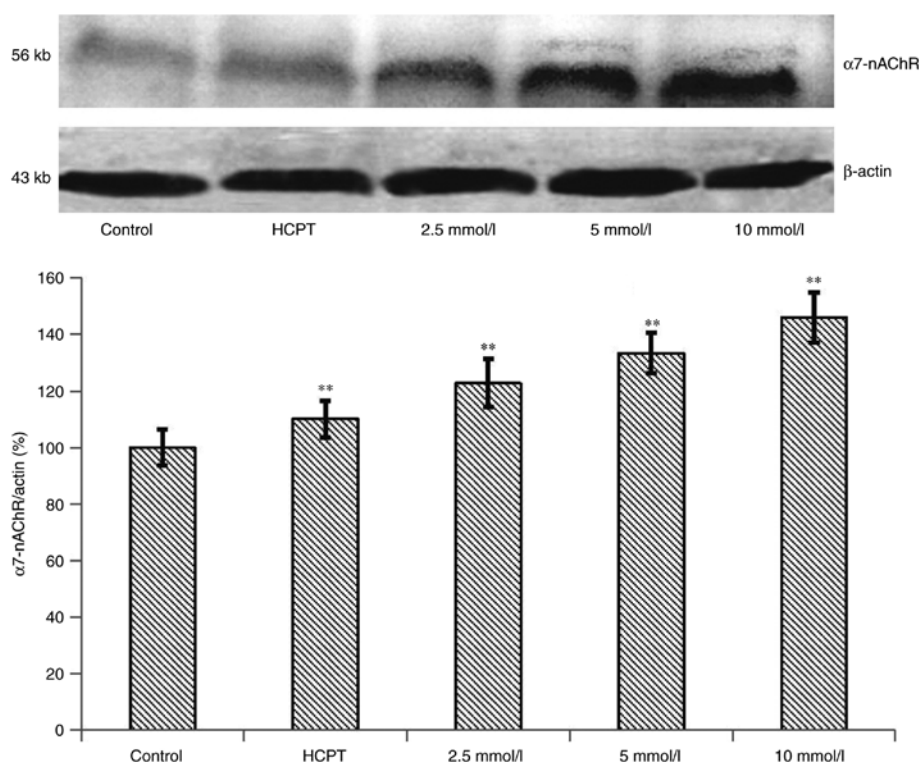
^aCompared with control P<0.01.

Figure 7. Effect of cytosine on the expression level of $\alpha 7$ -nAChR protein in HepG2 cells. Expression levels of $\alpha 7$ -nAChR were determined by western blotting. After treatment, the fractions were resolved by SDS-PAGE, transferred onto nitrocellulose membranes, and probed with primary antibody for $\alpha 7$ -nAChR and secondary antibody of IgG (H+L). The amount of β -actin was measured as an internal control. **Compared with control, P<0.01.

compared with the control group (P<0.01). Cytosine led to a significant increase in the level of caspase-4 and significant upregulation of caspase-4 expression.

Effects of the expression of $\alpha 7$ -nAChR protein in HepG2 cells.

To examine the $\alpha 7$ -nAChR expression induced by the overloading of Ca^{2+} , western blot experiments were conducted. As shown in Fig. 7, the $\alpha 7$ -nAChR expression level was significantly increased by treatment with 10 mM of cytosine for 24 h compared with the control group (P<0.01). Western blot analysis revealed that the exposure of the HepG2 cells to HCPT for 24 h induced a slight upregulation of the expression of the $\alpha 7$ -nAChR protein compared with the control group. The expression levels of $\alpha 7$ -nAChR were significantly upregulated in the HepG2 cells in a dose-dependent manner (P<0.01).

Discussion

Apoptosis induction is one of the mechanisms proposed for the anticancer therapeutic effects of cytosine. Our team previously confirmed that cytosine could induce apoptosis in HepG2 cells via a mitochondrial pathway (31). However, the molecular mechanisms of the effects of ER stress due to cytosine have not been thoroughly elucidated to date. ER stress, a newly defined signalling pathway, initiates apoptosis (32-34). In the present study, we demonstrated that Ca^{2+} overload induced by cytosine promoted endoplasmic reticulum (ER) stress, consequently leading to HepG2 cell apoptosis. In a previous study, we showed that cytosine, which is a naturally occurring quinolizidine alkaloid, had an anticancer effect. Our prior MTT assay investigation proved that cytosine could inhibit the proliferation of human HepG2 cells in a dose-dependent manner (31).

In the present study, morphological changes of apoptotic cells were observed after treatment with different concentrations of cytosine. Cell surface microvilli decreased and disappeared with an increase in drug concentration. Different amounts of spherical protrusions, apoptotic bodies, cell membrane rupture and cytoplasm overflow were observed in the apoptotic cells with transmission electron microscopy. ER dilatation and mitochondrial swelling were observed in the HepG2 cells after a high dose of cytosine. In the positive control group, the mitochondria were reduced, and the cell processes were obviously decreased. Typical apoptotic cells and apoptotic bodies were observed in the HepG2 cells. These results suggested that cytosine induces the apoptosis of the HepG2 cells (Fig. 1).

The ER is associated with protein synthesis, post-translational modification, and absorption and release of calcium ions. However, when the quality control system is overloaded under various stresses, including disruption of Ca^{2+} homeostasis, ER functions are impaired, and unfolded proteins are accumulated in the ER lumen. This situation is called ER stress. The ER is the primary site of intracellular Ca^{2+} storage (35-37). Calcium homeostasis is central to all cellular functions and has been studied for decades (38-40). Ca^{2+} is a critical second messenger that mediates many physiological cellular signalling pathways (41). The ER controls the calcium ion through the ryanodine receptor, inositol triphosphate receptor and sarco/endoplasmic reticulum Ca^{2+} -ATPase (41). Moreover, research has clearly confirmed that $[\text{Ca}^{2+}]_i$ accumulation contributes to cell death (42). Studies have shown that cytosine can give rise to Ca^{2+} overload (43). Whether Ca^{2+} overload induced by cytosine activates apoptosis in HepG2 cells remained undetermined. We hypothesized that cytosine stimulation would cause ER stress due to Ca^{2+} overload, leading to apoptosis. To determine whether the effects of cytosine on ER resulted in apoptosis, HepG2 cells were treated for 24 h with various concentrations of cytosine. Flow cytometry with PI staining revealed that the drug treatment increased the proportion of apoptotic cells (Table I and Fig. 2), confirming the cytosine-induced apoptosis in HepG2 cells. Next, laser confocal fluorescence microscopy was used to examine the intracellular Ca^{2+} concentration. Our data showed that cytosine increased the intracellular Ca^{2+} concentration in a cytosine-dependent manner (Table II and Fig. 3). These results suggested that cytosine increased the intracellular Ca^{2+} concentration by releasing Ca^{2+} from the ER, thereby causing ER stress. These findings revealed that calcium as a starting factor could induce ER stress, resulting in HepG2 cell apoptosis. It was concluded that HepG2 cellular exposure to cytosine resulted in a direct and immediate consequence of Ca^{2+} overload that led to disturbance in Ca^{2+} homeostasis in the ER, which caused ER stress leading to apoptosis.

When ER stress is prolonged, pro-apoptotic signaling pathways are activated, such as the CCAAT/enhancer-binding protein (C/EBP) and homologous protein (CHOP/GADD153), caspase-12 and JNK-dependent pathways (44,45). The caspase family plays an important role in apoptosis. Previous research has shown that caspase-12 is involved in ER-mediated apoptosis (46). Overexpression of caspase-12 only occurs during ER stress. It has been reported that caspase-12 triggers cellular apoptosis through the caspase-9/caspase-3 pathway. Cleaved

caspase-12 reportedly activates caspase-9, followed by activation of caspase-3 (47,48). In a previous study, we found that human caspase-4, one of the closest paralogs of rodent caspase-12, plays an important role in cell death via ER stress. However, the relevance of caspase-12 to ER-induced apoptosis has been questioned due to the absence of caspase-12 in most humans (49). Upregulation of caspase-4 in HepG2 cells had been demonstrated in the present study. When cells were incubated with cytosine, the expression of caspase-4 was significantly higher than that noted in the control (Table III). Thus, we conclude that caspase-4 plays an important role in cytosine-induced apoptosis that is associated with ER stress. It is suggested that Ca^{2+} overload, as an apoptosis-inducing factor, may cause ER stress to increase caspase-4 expression, which ultimately activates caspase-3 (this result has been confirmed by our team) (31), leading to apoptosis.

Another mediator of programmed cell death in ER-stressed cells is the transcription factor, CHOP. In line with previous reports, we found that significant overexpression of CHOP/GADD153 promotes cell death or DNA damage (50). Elevated levels of CHOP can promote the transcription of Bim, BAX and DR5 while suppressing the induction of Bcl-2 in ER stress (51). In addition, upregulation of the expression of CHOP is involved in caspase activation and mitochondrial events (50). In the present study, the expression of CHOP (Fig. 4) was continuously increased after cytosine treatment in the HepG2 cells. In the present study, calcium overload was hypothesized to result in a high expression of the CHOP gene, which promoted the accumulation of Bim, BAX and DR5 while suppressing the induction of Bcl-2. Indeed, upregulation of the expression of CHOP is involved in ER stress-induced apoptosis, and although it is normally undetectable in proliferating cells, it becomes highly synthesized in cells exposed to cytosine that perturb the homeostasis of the ER, which is linked to the development of apoptosis.

JNK, a member of the Ser/Thr protein kinase family, is induced by stress conditions, such as ER stress (52). Additionally, JNK participates in cell proliferation, survival, and apoptosis (53). It has been confirmed by previous research that inhibition of JNK phosphorylation results in decreased levels of ER stress-induced caspase-3/-7 (54). JNK, which has three isoforms, i.e., JNK1, JNK2 and JNK3, is also one of the signaling pathways of ER. JNK1 and JNK2 are associated with cell death and play an important role in apoptosis (55). To investigate the role of JNK in ER stress that is induced by cytosine, western blotting was used to examine expression levels of JNK1/2 and p-JNK1/2. We found that the expression levels of p-JNK1 and p-JNK2 were significantly enhanced in HepG2 cells in a dose-dependent manner (Fig. 5). JNK1/2 was increased by cytosine treatment (Fig. 6). These results suggested that c-Jun phosphorylation by Ca^{2+} overload in the HepG2 cells was mainly associated with JNK1/2. Our data suggested that the ER stress-JNK pathway is involved in Ca^{2+} overload and that cytosine induced ER stress via Ca^{2+} overload, thereby promoting cell death due to subsequent JNK activation.

nAChRs play an important role in synaptic transmission; thus, the primary focus of nicotine and its receptors has been its physiological effects within the nervous system. However, there is growing evidence suggesting that nAChRs are not

only expressed in the nervous system but also in nonneuronal cells, including cancer cells (23,56). In support of our findings, gastric, bladder, colon and non-small lung cancer cells have been shown to express nAChR subunits, and nicotine has been shown to be mitogenic for vascular smooth muscle cells (57). nAChRs are known to play several important roles in cancer cells, such as proliferation, inflammation, angiogenesis and invasion (58,59). Thus, the expression of nAChRs in non-neuronal cells indicates that they possess functions independent of neurotransmission. Therefore, nAChRs are viewed as a novel drug target for the treatment of cancer. Current research indicates that the α -7 nicotinic receptor, which is one of the various types of subunits, is related to intracellular Ca^{2+} overload (60). However, the actual role of the α -7 nicotinic receptor in HepG2 cells is unknown. Determining the relationship between the calcium ion concentration and α -7-nAChR in HepG2 cells is crucial. Our data indicated that α -7-nAChR was expressed at a high level under cytosine conditions (Fig. 7). After the cells were treated with cytosine, an increase in intracellular calcium concentration in the HepG2 cells was noted suggesting that intracellular Ca^{2+} overload was related to the upregulation of the expression of α -7-nAChR. As a result, cytosine, which is a partial agonist to nicotinic acetylcholine, has a high affinity to α -7-nAChR. Cytosine activates nAChRs, and the confocal laser scanning results indicated that cytosine can induce HepG2 cell calcium overload. The results of the present study showed that cytosine activates nicotinic acetylcholine leading to calcium overload. However, the specific relevance between nicotinic acetylcholine activation and Ca^{2+} overload remains to be studied.

Our team previously confirmed that cytosine induced apoptosis of HepG2 cells via the mitochondrial pathway. Following treatment with cytosine, mitochondrial permeability may increase, which may subsequently lead to mitochondrial matrix expansion, outer membrane rupture and the release of cytochrome *c*. In addition, a previous study confirmed that increased cytochrome *c* release into the cytoplasm caused activation of caspase-3 (31). The present study investigated whether cytosine induced apoptosis via the ER pathway.

In summary, the results of the present study indicate that cytosine induces the apoptosis of tumour cells via an ER pathway. Our data clearly suggest that calcium overload promotes ER stress-induced apoptosis in cytosine-induced HepG2 cells, which occurs through modulating the CHOP/GADD153, JNK and caspase-4 pathways. Finally, the caspase cascade is activated to induce apoptosis of HepG2 cells. Cytosine has affinity to nAChRs, and it can activate α -7-nAChR expression. In conclusion, cytosine induced ER stress-mediated apoptotic pathway by activating CHOP, JNK and caspase-4 in HepG2 cells.

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