

Apoptosis induced by ardipusilloside III through BAD dephosphorylation and cleavage in human glioblastoma U251MG cells

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Abstract Ardipusilloside III is a saponin newly isolated from *Ardisia pusilla* A.DC. Since saponins have exhibited broad anti-cancer and pro-apoptotic activity, we investigated the ability of ardipusilloside III to induce apoptosis in human glioblastoma U251MG cells, as well as the involvement of apoptotic signaling pathways. Ardipusilloside III markedly suppressed proliferation of U251MG cells in a time- and dose-dependent manner ($P < 0.05$, $IC_{50} = 8.2 \mu\text{g/ml}$), but did not affect the growth of primary cultures of human astrocytes. Ardipusilloside III-treated U251MG cells underwent typical apoptotic changes. Exposure to a low dose of ardipusilloside III provoked G₂/M-phase cell cycle arrest, which preceded apoptosis characterized by the appearance of cells with sub-G₁ DNA content. However, a higher dose of ardipusilloside III induced apoptosis without first causing cell cycle arrest. In addition, ardipusilloside III exposure resulted in time-dependent BAD dephosphorylation and cleavage as well as activation of caspase-8 and caspase-3. Therefore, both the intrinsic pathway of apoptosis, mediated by BAD dephosphorylation and cleavage, and the extrinsic pathway of apoptosis, mediated by caspase-8 and caspase-3 activation,

were involved in ardipusilloside III-induced apoptosis. These data suggest that ardipusilloside III is a reliable candidate for chemotherapeutic treatment of human glioblastomas, and should be investigated further.

Keywords Ardipusilloside III · Glioblastoma · Apoptosis · BAD · Caspase

Abbreviations

IC ₅₀	50% inhibitory concentration
BAD	Bcl-xL/Bcl-2-associated death promoter homolog
DMEM	Dulbecco's modified Eagle's medium
MTT	3-(4, 5-dimethylthiazol-2-yl)-2, 5-diphenyltetrazolium bromide

Introduction

Glioblastoma multiforme (WHO Grade IV), composed of poorly differentiated neoplastic astrocytes, is the most common and lethal primary brain malignancy [1]. Despite the combination of treatments offered to patients, including selective surgery, radiotherapy, and chemotherapy, residual cancer cells inevitably invade the surrounding normal brain tissue, leading to cancer recurrence and a poor prognosis. Since glioblastomas are relatively resistant to radiation and chemotherapy due to genetic and epigenetic alterations, development of effective drugs to reverse its drug resistance and induce apoptosis is critical [2, 3].

The discovery of novel anti-cancer agents has progressed considerably in recent years. Compelling data from laboratory studies, epidemiological investigations, and human clinical trials all have demonstrated that saponins have significant potential in cancer chemoprevention and chemotherapy [4–6]. Saponins also have a neuroprotective

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effect [7] as well as an immunological adjuvant function [8], which could also provide relief from the development of serious lesions in patients with glioblastoma. However, the mechanisms by which saponins induce apoptosis are not clear.

Tumor growth is the result of either improper cell proliferation or cell death. Necrosis and apoptosis are the two principal mechanisms of cell death. Apoptosis, in contrast to necrosis, does not lead to cell lysis and inflammatory responses in cancer chemotherapy. Avoiding and controlling intense inflammatory responses could prevent fatal encephaledema. Therefore, increasing apoptosis without increasing necrosis during chemotherapeutic treatment of glioblastoma is ideal. The intracellular pathways leading to apoptotic cell death are complicated, but most involve activation of specific proteases known as caspases. There are two primary mechanisms by which caspases are activated: the extrinsic, or death receptor-mediated, pathway and the intrinsic, or mitochondrial-mediated, pathway [9]. The majority of known apoptotic signals are part of the mitochondrial pathway, including both the pro-apoptotic and the anti-apoptotic Bcl-2 family members, loss of mitochondrial transmembrane potential, and release of caspase activators. For death receptor-induced apoptosis, the crucial events are the formation of a Death Inducing Signaling Complex (DISC) and caspase-8 activation [10].

BAD, a member of the Bcl-2 homology 3 (BH3)-only subgroup of the Bcl-2 family as well as a ligand of the prosurvival protein 14-3-3, is a switch protein for the mitochondrial-induced apoptotic pathway [11]. Lacking a C-terminal hydrophobic domain essential for targeting the mitochondrial outer membrane, BAD distributes exclusively in cytoplasm, heterodimerized with 14-3-3. Both BAD dephosphorylation and cytochrome c release are essential to induce apoptosis [12]. In response to cell death stimuli, dephosphorylated BAD translocates to the outer mitochondrial membrane, where it displaces Bax from the Bcl-xL:Bax complex. The newly released Bax then forms a transmembrane pore across the outer mitochondrial membrane, leading to loss of membrane potential and efflux of cytochrome c and the apoptosis-inducing factor (AIF) [9, 13]. BAD also can promote the formation of the mitochondrial permeability transition pore directly [14] and mediate cell cycle progression by regulating the expression of cyclins [15]. During apoptotic cell death, BAD is cleaved by a caspase(s) at its N terminus to a truncated form. The 15-kDa truncated BAD termed by tBADS is a more potent inducer of apoptosis than uncleaved BAD. Previous studies have shown that many anti-cancer agents such as doxorubicin and genistein induce apoptosis through the BAD apoptotic signaling pathway, as well as endogenous signals such as transforming growth factor (TGF)- β 1

[16–18]. However, the role of these pathways in saponin-induced apoptosis has not been verified.

Ardisia pusilla A. DC (of the family Myrsinaceae), a Chinese medicinal herb also known as Jiu Jie Long, is the source of the triterpenoid saponins called ardisiposiliosides. Ardisiposilioside I and ardisiposilioside II have known anti-cancer activity in both Lewis pulmonary carcinoma and hepatocarcinoma [19, 20]. Recently, the structure of ardisiposilioside III, a newly discovered saponin, has been determined by extensive use of nuclear magnetic resonance (NMR) techniques as well as chemical evidence, although this work has not been published yet. Ardisiposilioside III is distinguished from ardisiposilioside I by the extra 28 α -hydroxy group in its aglycone, and different sugar moieties. Since the effect of this new agent on cancer cells has not been tested previously, we investigate herein the ability of ardisiposilioside III to induce apoptosis in human glioblastoma U251MG cells. In addition, we seek evidence of the molecular mechanism by which this potential chemotherapeutic agent kills cancer cells.

Materials and methods

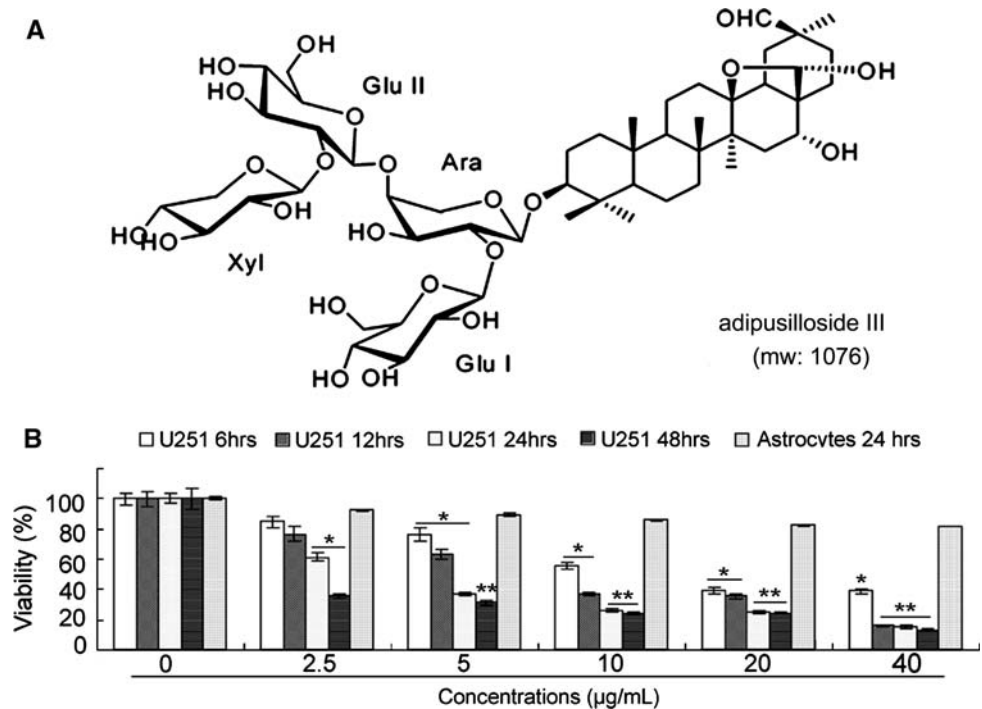
Sample preparation

Purified ardisiposilioside III (Purity > 98%) was supplied by the Department of Pharmacy of Xijing Hospital, the Fourth Military Medical University, Xi'an, P.R.China. Ardisiposilioside III was isolated from *A. pusilla* and established as 3-O- $\{\beta$ -D-xylopyranosyl-(1 \rightarrow 2)- β -D-glucopyranosyl-(1 \rightarrow 4)- $[\beta$ -D-glucopyranosyl-(1 \rightarrow 2)]- α -L-arabinopyranosyl]-3 β , 16 α , 28 α -trihydroxy-13 β , 28-epoxy-oleanan-30-al with the molecular formula C₅₂H₈₄O₂₃ (molecular weight: 1,076). The structure of this compound is shown in Fig. 1a. Stock solutions were prepared by dissolving the dry material in dimethylsulfoxide (DMSO) (Gibco/Invitrogen, NY, USA) and storing aliquots at -20°C . These were later diluted to the final concentration in fresh medium before each experiment. In all experiments, the final DMSO concentration did not exceed 1% (v/v), so as not to affect cell growth. Cells incubated with ardisiposilioside III-free medium were used as controls.

Cell lines and cell culture

U251MG human glioblastoma cells (maintained in our laboratory, originally obtained from Uppsala, Sweden) were cultured in DMEM medium supplemented with 10% newborn calf serum (both from Gibco/Invitrogen, NY, USA) in a 37 $^{\circ}\text{C}$ incubator with a humidified atmosphere of 5% CO₂–95% O₂.

Fig. 1 In vitro treatment of human glioblastoma cells with adipusilloside III inhibits proliferation in a dose- and time-dependent manner: (a) The chemical structure of adipusilloside III; (b) Dose- and time-dependent effects of adipusilloside III on the viability of human glioblastoma U251MG cells and primary cultured human astrocytes. Columns, mean percent of viable cells from three experiments; Bars, \pm SE; * $P < 0.05$; ** $P < 0.01$, versus control group (adipusilloside III -free)



Cultured primary astrocytes were obtained from a slightly impaired brain tissue fragment of a volunteer with cerebral trauma who consented to the procedure. Acquisition of the tissue was done following approval by the local medical research ethics committee at Xijing Hospital, the Fourth Military Medical University, Xi'an, P.R. China. The fragment was dissected for its grey matter by a surgeon when it was removed, washed in phosphate buffered sodium (PBS) and dispersed repeatedly. The resulting cell suspension was filtered and cultured in DMEM with 10% newborn calf serum. After 2 weeks in culture, the remaining cells were mostly astrocytes [21].

Twenty-four hours before the experiments, the cells were transferred to serum free medium. Adipusilloside III of different concentrations was added to the culture medium for different incubation periods as indicated in the Results section.

MTT assay for cytotoxicity

The number of viable cells was determined by the 3-(4, 5-dimethylthiazol-2-yl)-2, 5-diphenyltetrazolium bromide (MTT) assay as described previously [22]. Briefly, the cells were cultured in 96-well plates (Greiner Bio-One GmbH, Frickenhausen, Germany) at a density of 4×10^3 cells/well in the presence of adipusilloside III at the indicated concentrations. After incubation for 6, 24, 48, or 72 h, the MTT (Sigma, St. Louis, MO, USA) dissolved in PBS was added to each well at a final concentration of 5 mg/ml and

then incubated at 37°C for 4 h. The water-insoluble dark blue formazan crystals that formed during MTT cleavage in actively metabolizing cells were dissolved in DMSO. The optical density was measured at a wavelength of 490 nm with a Bio-Rad 680 microplate reader (Bio-Rad, California, USA). The reduction in viability of the adipusilloside III-treated U251MG cells and astrocytes was expressed as the percentage compared with adipusilloside III-free control cells. All experiments were done in triplicate.

Cell cycle analysis

Cell cycle distribution was analyzed by flow cytometric analysis as described previously [23]. Briefly, following incubating for 12, 24, 48, or 72 h, U251MG cells treated with or without adipusilloside III were trypsinized with 0.25% trypsin (Sigma, St. Louis, MO, USA), counted, centrifuged at 300g for 5 min, and fixed in ethanol at 4°C overnight. Then the cells were washed and centrifuged. The resulting cell pellets were resuspended in an RNase solution (0.02 mg/ml; Sigma) containing propidium iodide (0.02 mg/ml; Sigma), and incubated at 4°C for 30 min. The DNA content of approximately $1-2 \times 10^5$ stained cells were analyzed using a FACScan flow cytometer equipped with the FACStation data management system running Cell Quest software (Becton Dickinson, San Jose, CA, USA). The results are expressed as a plot of fluorescence intensity versus cell number.

Electron microscopy

U251MG cells were cultured in T-150 flasks (Greiner Bio-One GmbH, Frickenhausen, Germany) (3×10^6 cells/cm²) and treated with 8.2 µg/ml ardisiposilioside III for 24 h, and then trypsinized with 0.25% trypsin and centrifuged at 1,400g for 15 min. The pellets were fixed in 0.1 M PBS (pH 7.4) with 2.5% glutaraldehyde, then postfixed in 2% buffered osmium tetroxide for 2 h and finally dehydrated through a series of graded ethyl alcohols from 70% to 100%. The schedule is as follows: 70% for 10 min., 95% for 10 min. and three changes of 100% for 5 min each. Specimens used for transmission electron microscopy were embedded in epon resin. Thin sections were cut on an ultramicrotome and double stained with uranyl acetate and lead citrate. Electron micrographs were taken on an electron microscope (JEM-2000EX, JEOL Ltd., Tokyo, Japan) operating at 80 kV.

Annexin V/propidium iodide

To determine a number of apoptotic cells, Annexin V assays were performed using an apoptosis detection kit (Annexin V-FITC/PI Staining Kit; Immunotech Co., Marseille, France). Briefly, 1.5×10^5 cells were plated in 24-well plates, and treated with 3.7 and 8.2 µg/ml ardisiposilioside III for 6 and 12 h, respectively. Cells were harvested by treatment with 0.02% EDTA in PBS or by scraping, washed in cold PBS (0.01 mol/l, pH = 7.4), incubated for 15 min with fluorescein-conjugated annexin V and propidium iodide, and analyzed using the same flow cytometer and software used for the cell cycle analysis. PI-negative but annexin V-positive cells were considered to be early apoptotic, while cells that were both PI and annexin V negative were considered normal.

Analysis of DNA fragmentation

Cellular DNA was extracted from the cells according to Gong's modified method [24]. Briefly, cells treated with or without 8.2 µg/ml ardisiposilioside III treatment for 24 h were trypsinized with 0.25% trypsin and collected by centrifugation (200g, 10 min), washed twice in cold PBS (0.01 mol/l, pH = 7.4) and resuspended at a density of 4×10^6 cells/400 µl in hypotonic lysing buffer (5 mM Tris, 20 mM EDTA, pH 7.4) containing 0.5% Triton X-100 for 30 min at 4°C. The lysates were centrifuged at 13,000g for 15 min at 4°C. Fragmented DNA was extracted from the supernatant with phenol-chloroform-isoamylalcohol, precipitated by addition of 2 vol. of absolute ethanol and 0.1 vol. of 3 mM sodium acetate, and treated with RNase

A (500 U/ml) at 37°C for 3 h. The pattern of DNA fragmentation was visualized by electrophoresis in 1.5% agarose gel containing ethidium bromide in 40 mM Tris-acetate buffer (pH 7.5) at 50 V for 4 h and photographed under UV light.

Cellular and nuclear morphology

Apoptotic U251MG cells were identified based on alterations in their nuclear morphology detected by staining with Hoechst 33342 as described previously [25]. Briefly, cells were grown in 24-well plates (1×10^4 cells/well) in the presence of 3.7 and 8.2 µg/ml ardisiposilioside III for 24 h. Cells were observed with an inverted microscope (Leica Microsystems, Wetzlar, Germany), then were washed in PBS and fixed in 70% ethanol for 2 h at 4°C. Cell nuclei were stained with 5 µg/ml Hoechst 33342 (Sigma, St. Louis, MO, USA). After final washing in PBS, the changes in nuclear morphology were visualized by fluorescence microscopy (Leica Microsystems, Wetzlar, Germany) using excitation at wavelength of 330–380 nm.

Western blot analysis

Protein extracts of U251MG cells (ardisiposilioside III-free control and cells treated with 8.2 µg/ml ardisiposilioside III for 6, 12, or 24 h) were prepared by lysing the cells in RIPA buffer (150 mM NaCl, 1% Nonidet P-40, 0.5% sodium deoxycholate, 0.1% SDS, 50 mM Tris-HCl, pH 8) containing 10 mM EDTA, 1 mM sodium orthovanadate and 1 mM phenylmethylsulfonylfluoride (PMSF; all from Sigma) for 30 min at 4°C. Samples were then centrifuged at 14,000g for 25 min at 4°C. The protein concentration in the supernatant was determined by BCA Protein Assay Kit (HyClone-Pierce, Utah, USA). Equivalent amounts (60 µg protein/lane) of protein lysates were separated by 12% SDS-polyacrylamide gel electrophoresis and transferred onto a nitrocellulose membrane (0.22 µm, Invitrogen, NY, USA) in a transfer tank (Bio-Rad, California, USA) using the submerged method. The membrane was blocked for 2 h at room temperature in PBS containing 0.1% Tween-20 (Sigma) and 5% non-fat dried milk (Carnation, city, state, USA), and then incubated with the primary antibody in dilution buffer ($1 \times$ TBS, 0.1% Tween-20 with 5% BSA) with gentle agitation overnight at 4°C. Primary antibodies were: anti-BAD (diluted 1:400, rabbit polyclonal), anti-phospho-BAD (Ser136) (diluted 1:200, rabbit polyclonal), anti-Caspase-8 (1C12) (diluted 1:600, mouse monoclonal), anti-caspase-3 (diluted 1:600, rabbit polyclonal, all from Cell Signaling Technology, Beverly, MA, USA), and anti-β-actin (diluted 1:400, mouse monoclonal C-2, Santa Cruz,

CA, USA). Then, the membranes were washed in dilution buffer ($1 \times$ TBS, 0.1% Tween-20 with 5% BSA) and incubated for 1 hour with horseradish-peroxidase (HRP)-conjugated secondary antibodies, and finally developed by an ECL system (Cell Signaling Technology, Beverly, MA, USA). Secondary antibodies were: HRP-conjugated anti-rabbit IgG (diluted 1:2000) and HRP-conjugated anti-mouse IgG (diluted 1:2000, both from Cell Signaling Technology). The Western blotting was performed following Laemmli's method [26] and the grayscale values of each band on the blots were measured using BandScan4.3.

Statistical analysis

Data are expressed as the mean \pm the standard error of mean (SEM) of separate experiments ($n \geq 3$, where n represents the number of independent experiments). All data were tested for significance by one-way analysis of variance (ANOVA) followed by Fisher's post hoc test using SPSS13.0 software. Results with a P value less than or equal to 0.05 were considered statistically significant.

Results

Cytotoxicity of ardisipilloside III in both U251MG cells and primary astrocytes

The effects of ardisipilloside III on the viability of U251MG cells and primary cultured human astrocytes were assessed by the MTT assay (Fig. 1b). Ardisipilloside III inhibited the growth of U251MG cells in a dose- and time-dependent manner. Viability of U251MG cells treated with 5 $\mu\text{g/ml}$ ardisipilloside III for 6 h decreased to 71.3% ($n = 10$, $P < 0.05$). After U251MG cells were treated with 40 $\mu\text{g/ml}$ ardisipilloside III for 72 h, the percent of viable cells significantly decreased to 12.6% ($n = 10$, $P < 0.05$). The concentrations at which ardisipilloside III inhibited cell growth by 25% (IC_{25}) and 50% (IC_{50}) at 24 h were 3.7 and 8.2 $\mu\text{g/ml}$, respectively. However, exposure of the astrocytes to different concentrations of ardisipilloside III (5–40 $\mu\text{g/ml}$) for 24 h did not result in any statistically significant change in cell viability, with the viability ranging from 81.7% to 92.3% ($n = 10$, $P > 0.05$). Therefore, ardisipilloside III reduced the viability of U251MG cells, but was only slightly toxic to the primary cultured human astrocytes.

Cell cycle change in U251MG cells treated with ardisipilloside III

The cellular DNA content was analyzed by flow cytometry to detect changes in the cell cycle distribution. DNA

histogram analysis showed that, following incubation with ardisipilloside III at a low concentration (3.7 $\mu\text{g/ml}$), the percentage of S phase cells reduced from 44.2% to 14.6%; while that of cells in the G_2/M phase increased from 8.9% to 26.6% within 48 h, suggesting G_2/M phase arrest. At 72 h, the presence of a significant sub- G_1 phase (1.4%) fraction suggests induction of apoptosis in the U251MG cells. Ardisipilloside III arrested the cell cycle at the G_2/M phase transition prior to the induction of apoptosis in the glioblastoma cells. However, at a higher concentration (8.2 $\mu\text{g/ml}$), ardisipilloside III quickly induced apoptosis, characterized by the appearance of cells with sub- G_1 DNA content, without arresting the cell cycle (Fig. 2a, Table 1).

Morphological alteration of U251MG cells treated with ardisipilloside III

Hoechst 33342 nuclear staining show that U251MG cells underwent marked morphological changes in a time- and dose-dependent manner after incubation with ardisipilloside III. The U251MG cells shrank, aggregated, and detached from the surface of the culture flask. Apoptotic U251MG cells showed characteristic morphological and biochemical features such as chromatin aggregation, nuclear and cytoplasmic condensation, and the partition of cytoplasm and nucleus into membrane bound-vesicles known as apoptotic bodies (Fig. 3). Electron microscopy revealed the apoptotic changes of intracellular structures in most ardisipilloside III—treated cells. In comparison with the normal intracellular morphology of the control cells, apoptotic U251MG cells lost their microvilli, had increased numbers of lysosomes, and had condensed, fractured, and marginalized chromatin. Also, numerous membrane-enclosed vacuoles containing cytoplasm and parts of the fractured nucleus were found in the treated cells, which would most likely become apoptotic bodies (Fig. 2b).

Ardisipilloside III increased phosphatidylserine exposure and induced DNA cleavage in U251MG cells

To quantify early apoptotic effects, the cells were stained with annexin V and propidium iodide (PI) following ardisipilloside III exposure. The right lower quadrant of the density dot plots from the flow cytometric analysis represented early apoptotic cells. Phosphatidylserine externalization occurred after ardisipilloside III treatment in a time- and dose-dependent manner (Table 2). The number of apoptotic cells increased at both concentrations (3.7 and 8.2 $\mu\text{g/ml}$) with time (6 and 12 h). Despite the fact that the number of early apoptotic cells increased after treatment with the higher concentration, the late

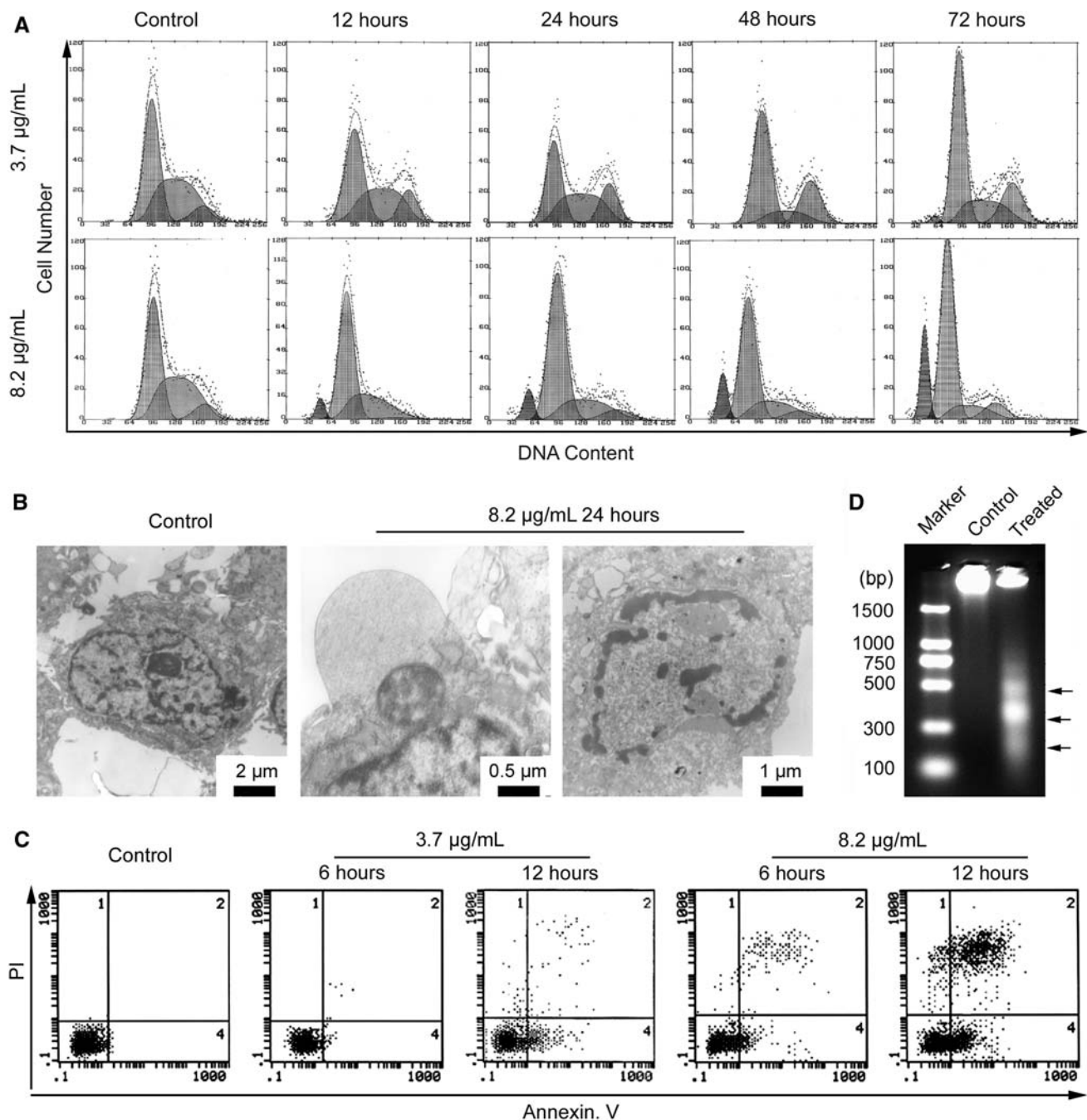


Fig. 2 Effect of ardisiposilide III on cell cycle progression, intracellular structures, phosphatidylserine externalization and DNA fragmentation of human glioblastoma U251MG cells: (a) Flow cytometric analysis of cell cycle phase distribution of U251MG cells after treatment with different concentrations of ardisiposilide III for the indicated time periods; (b) Electron microscopy showed the organization of subcellular organelles of U251MG cells, including the presence of vacuoles containing cytoplasm and nucleolus fragments budding off the plasma membrane (center, magnification $\times 15,000$)

apoptotic cells and necrotic cells also increased to some extent, which suggested that treatments should not exceed the IC_{50} , in order to avoid necrosis *in vitro* (Fig. 2c).

and marginal chromatin condensation (right, magnification $\times 7,500$); (c) Representative dot plots showing flow cytometric analysis of glioblastoma U251MG cells treated with different concentrations of ardisiposilide III for the indicated time periods then stained with FITC-conjugated Annexin V and propidium iodide; (d) The characteristic DNA ladder suggesting apoptotic DNA damage in the ardisiposilide III-treated U251MG cells is shown in Lane 3. Cells were treated with 8.2 µg/ml ardisiposilide III for 24 h

Electrophoresis of cellular DNA revealed a distinctive ladder pattern of DNA cleavage in the apoptotic U251MG cells, which represents the discrete fragments of 180–200

Table 1 Cell cycle distribution of ardisiposilioside III-treated U251MG cells (%)

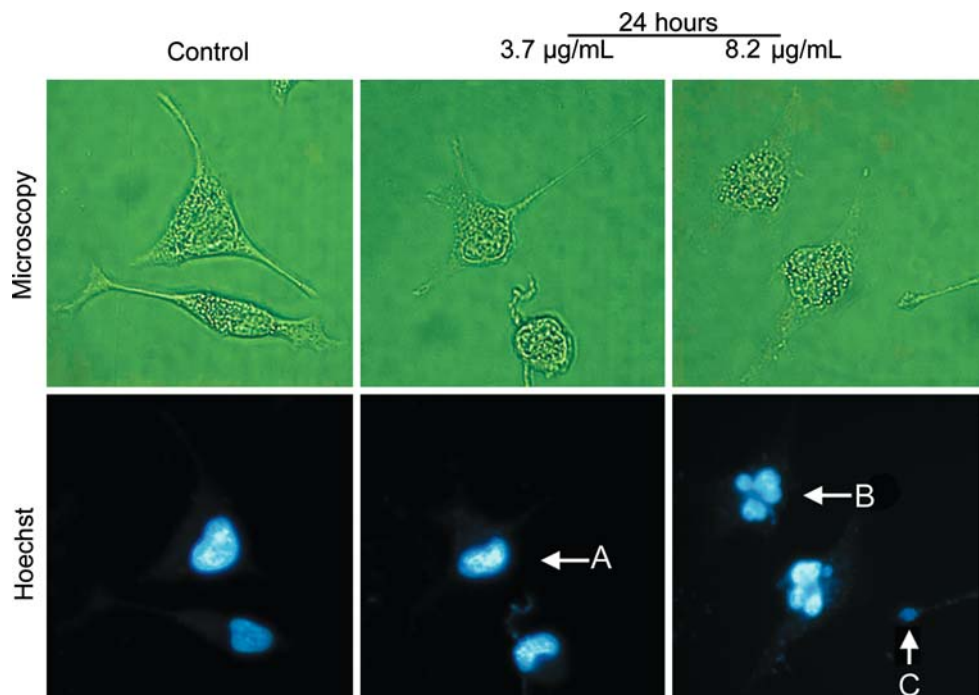
Cell cycle status	Control	3.7 µg/ml				8.2 µg/ml			
		12 h	24 h	48 h	72 h	12 h	24 h	48 h	72 h
Sub-G ₁	0	0	0	0	1.4	5.7	7.3	12.3	18.0
G ₀ /G ₁	46.9	46.0	39.2	58.8	54.3	64.5	67.1	67.2	72.8
S	44.2	38.8	39.3	14.6	24.1	33.5	25.3	25.1	16.0
G ₂ /M	8.9	15.2	21.5	26.6	21.5	1.9	7.6	7.7	11.2

base pairs or multiples thereof that result from non-random cleavage between the nucleosomes. The ladder pattern did not appear in the DNA isolated from control cells (Fig. 2d). Electrophoresis of cellular DNA revealed that ardisiposilioside III induced apoptosis specific DNA cleavage in U251MG cells.

Dephosphorylation and cleavage of BAD in ardisiposilioside III-induced apoptosis

To determine the involvement of BAD in ardisiposilioside III-induced apoptosis, the steady-state levels of phospho-BAD (serine 136) were measured by Western blot analysis (Fig. 4a). The expression of endogenous phospho-BAD was decreased slightly ($p\text{-BAD}/\text{actin} = 89.9\% \pm 0.4\%$, $n = 3$, $P < 0.01$) after ardisiposilioside III treatment for 6 h and decreased even further ($78.4\% \pm 0.1\%$, $n = 3$, $P < 0.01$) after 24 h of treatment, while the ardisiposilioside III-free control group showed no difference.

Fig. 3 Morphological alterations of U251MG cells treated with different concentrations of ardisiposilioside III for 24 h. Treated cells shrank, aggregated, and detached from the bottom of culture flask as shown in the upper panels (magnification $\times 600$). Nuclear morphology was detected using Hoechst 33342, and visualized by fluorescence microscopy as shown in the lower panels (magnification $\times 600$): (a) Cell condensation and fracturing of marginal chromatin; (b) The lobulated gemmules of the cell nucleus; (c) Apoptotic body

**Table 2** Apoptosis of ardisiposilioside III-treated U251MG cells (%)

Apoptosis status	Control	3.7 µg/ml		8.2 µg/ml	
		6 h	12 h	6 h	12 h
Normal	93.5	78.1	74.8	64.6	30.3
Early apoptosis	4.1	17.0	17.3	17.3	39.0
late Apoptosis	1.9	3.2	5.4	14.4	21.8
Necrosis	0.5	1.7	2.6	3.8	8.9

Treatment of U251MG cells with 8.2 µg/ml ardisiposilioside III also resulted in Bad cleavage. As early as 6 h after treatment, the tBADs/actin ratio increased significantly ($78.8\% \pm 0.4\%$, $n = 3$, $P < 0.01$) and continued to increase up to 24 h ($87.8\% \pm 0.2\%$, $n = 3$, $P < 0.01$). We detected no cleaved BAD in cells cultured without ardisiposilioside III (Fig. 4a). In addition, total BAD expression began to decrease slightly after 12 h of treatment ($94.6\% \pm 0.9\%$, $n = 3$, $P < 0.01$), continuing to decrease up to 24 h ($93.6\% \pm 0.9\%$, $n = 3$, $P < 0.01$).

Western blot analysis well described the dephosphorylation and cleavage of BAD in ardisiposilioside III-induced apoptosis, confirmed the intrinsic BAD apoptotic pathway underlying the ardisiposilioside III-induced apoptosis.

Caspase-8 and -3 are activated in ardisiposilioside III-induced apoptosis

To investigate the contribution of the extrinsic apoptotic pathway in ardisiposilioside III-induced apoptosis, we

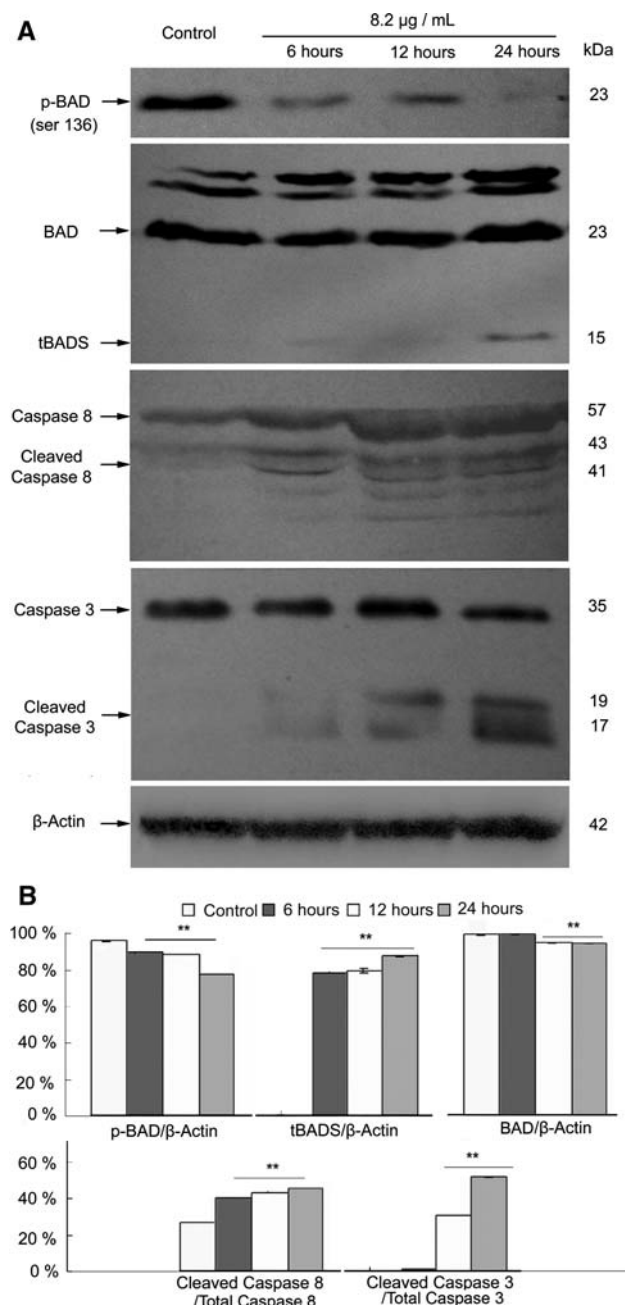


Fig. 4 Activation of BAD and caspases involved in ardisipusilioside III-induced apoptosis of human glioblastoma U251MG cells: (a) BAD was dephosphorylated at serine136 and cleaved to generate tBADS (15 kDa), as indicated by the arrow. Phospho-BAD expression decreased, while tBADS expression increased, during the indicated time course. The expression of total BAD was slightly decreased after ardisipusilioside III treatment. Rapid and substantial caspase-8 and -3 cleavage was detected after treatment with 8.2 μ g/ml ardisipusilioside III for the indicated time periods; (b) Statistical analysis of the Western blots. Bars, \pm SE; * $P < 0.05$; ** $P < 0.01$

assessed the activation of caspase-8 and caspase-3. Figure 4a shows that, in U251MG cells treated with 8.2 μ g/ml ardisipusilioside III, both caspase-8 and caspase-3 were

rapidly and substantially cleaved. There is some idiopathic activation of caspase-8 in normal glioblastoma cells. However, greatly increased caspase-8 activation was observed after 6 h of treatment ($41.7\% \pm 0.6\%$, $n = 3$, $P < 0.01$) and was maintained ($43.6\% \pm 0.6\%$, $n = 9$). The first cleavage products of caspase-3 were found after 6 h of exposure to ardisipusilioside III ($2.6\% \pm 1.0\%$, $n = 3$, $P < 0.05$), and increased significantly up to 12 h after exposure ($43.5\% \pm 5.2\%$, $n = 6$, $P < 0.01$). Data suggested the extrinsic apoptotic pathway mediated by caspase-8 and -3 also involved in the ardisipusilioside III-induced apoptosis.

Discussion

BAD is regulated primarily by phosphorylation, in response to survival factors, at several sites, including serine 136, which of BAD is vital for 14-3-3 interaction in mammalian cells [27]. BAD phosphorylation protects cancer cells from apoptosis by raising the threshold at which the mitochondria release cytochrome c to induce cell death [28]. Extensive researches have shown that carcinogenesis and cancer progression are highly correlated with BAD phosphorylation [29–31]. Over-expressed phospho-BAD is frequently detected in cancer cells such as rat glioma C6 cells [29], glioblastoma cells [30], and human neuroblastoma SH-SY5Y cells [31]. BAD dephosphorylation and redistribution from the cytosol to the mitochondria are early events in chemotherapy-induced apoptosis, and are regarded as prominent links between the inhibition of survival pathways and the onset of execution phase of apoptosis. Dephosphorylation at serine 136 decreases the accessibility to serine 155 by survival-promoting kinases, disrupting the associations between BAD and prosurvival proteins [32]. Compared to standard treatments, combined usage of anti-cancer agents to prevent BAD phosphorylation could be a promising therapeutic strategy. Recent experiments suggest that many anti-cancer agents, like doxorubicin [16], cannabinoids [30], peroxisome-proliferator-activated receptor-c (PPARc) agonists [33], paclitaxel [34], cyanide [35], and ammonia [36], induce apoptosis by dephosphorylating BAD. We found BAD dephosphorylation in ardisipusilioside III-induced U251MG apoptosis. BAD dephosphorylation may trigger the various signaling pathways by which apoptosis is induced in glioblastoma cells.

The final apoptotic pathway examined herein involved the caspase cascades that cleave key cellular components, including cytoskeletal proteins and nuclear proteins, to produce the altered morphology of the treated U251MG cells. Caspases can also activate other catabolic enzymes such as DNases, which cleave DNA into the characteristic

ladder [37] we have observed. Caspases also cleave Bcl-2 family members at specific sites during apoptosis induced by growth factor withdrawal or other apoptotic stimuli [38]. The BAD-caspase positive feedback cascade reaction can amplify the apoptotic signal to some extent. As a response to the cell death signals produced by anticancer agents, BAD is cleaved by caspase-3 to generate two truncated products: tBADL (truncated BAD large, 26 kDa) and tBADS (truncated BAD small, 15 kDa). tBADL inhibits apoptosis in impaired neurons [39]. tBADS is poorly phosphorylated due to alterations in its secondary structure as the result of cleavage, so it interacts inefficiently with 14-3-3 proteins but efficiently with Bcl-xL, thereby altering the Bcl-xL-Bax balance and activating the apoptotic program [39, 40]. Genistein [17], raloxifene [41], and isoliquiritigenin [42] induce apoptosis by cleaving BAD into small fragments in many types of cancer cell; thereby proving that tBADS is a more potent inducer of apoptosis than uncleaved BAD. We have detected increased levels of tBADS, which could reinforce the other mechanisms of ardisiposilioside III-induced U251MG apoptosis.

Since caspase-3 can be activated directly by caspase-8 in Fas-mediated apoptosis, and glioblastoma cells are Fas-sensitive, tBADS generated from caspase-3-dependent cleavage induces apoptosis more efficiently. Our study also showed that both caspase-8 and -3 were activated immediately after exposure. The activation of caspase-3 was amplified by caspase-8 activation in ardisiposilioside III-induced U251MG apoptosis. Therefore, the mechanism of caspase-dependent BAD cleavage may be critical to improving the chemotherapeutic treatment of glioblastoma. However, BAD activation and the caspase cascade occurred at almost the same time, so whether BAD was activated by the caspase cascade or not has not been clearly proven. Herein we have not excluded an independent effect of the caspase cascade on ardisiposilioside III-induced apoptosis; however, we consider that both BAD dephosphorylation and the cleavage-mediated intrinsic pathways, as well as the caspase-8 and -3 activation-mediated extrinsic pathways, play an important role in ardisiposilioside III-induced apoptosis. In addition, the fact that ardisiposilioside III did not significantly affect the viability of astrocytes, together with the neuroprotective effect of tBADL (data not shown), suggests that ardisiposilioside III may be safer in a clinical setting than other anti-cancer agents.

Because of the cross regulation necessary for intracellular homeostasis, alteration of one physiological protein may affect more than a single physiological process [15]. Impairment of cell cycle progression may activate the mitochondrial apoptotic pathway [30, 43]. BAD is interesting, both for its apoptosis-regulating capacity and for its effect on cell cycle progression [15]. Our previous study showed that cyclin D2 was overexpressed in glioblastoma

cells but hardly expressed in normal brain tissue [44], and was responsible for the progression of glioblastoma cells past the G₁/S transition of the cell cycle. It is possible that, following ardisiposilioside III treatment, BAD is dephosphorylated and thereby activated, altering the expression of cell cyclins such as cyclin D2, thus leading to cell cycle changes. However, these processes are not yet clear. The time course study of DNA histogram analysis revealed that ardisiposilioside III at a low concentration provoked cell cycle arrest at the G₂/M phase transition prior to the induction of apoptosis, a higher concentration induced apoptosis without any detectable cell cycle inhibition. These results prove that the apoptotic response of glioblastoma U251MG cells to ardisiposilioside III at a low concentration was different from that at a high concentration. Whether U251MG cells undergo cell cycle arrest or apoptosis after exposure to ardisiposilioside III at different concentrations is the result of initiation of different pathways. However, whether BAD was involved in the cell cycle arrest needs further study.

In conclusion, ardisiposilioside III is capable of inhibiting growth and inducing apoptosis of U251MG cells in a dose- and time-dependent manner. The anti-cancer effects of ardisiposilioside III may result from multiple mechanisms, such as interfering with cell cycle progression and inducing apoptosis. There are also multiple possible mechanisms by which ardisiposilioside III induces apoptosis, through both the intrinsic, BAD dephosphorylation and cleavage pathway, and the extrinsic caspase-8/caspase-3 pathway. We postulate that this new agent may hold a promise as an anti-cancer agent. Based on the observations that ardisiposilioside III has significant anti-glioblastoma activity *in vitro*, further study of its effects and related mechanisms *in vivo* are needed. This study provides important data to evaluate the possibility of clinical application for ardisiposilioside III as an anti-cancer agent.

Conclusion

We identified the novel saponin ardisiposilioside III as an anti-cancer and apoptosis-inducing agent using the glioblastoma cell line U251MG cells. Both the BAD-mediated intrinsic apoptotic signaling pathway and the caspase-8-mediated extrinsic apoptotic signaling pathway are involved in ardisiposilioside III-induced apoptosis of U251MG cells. These results suggest that ardisiposilioside III could be a potential novel agent for use in glioblastoma chemotherapy.

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