Abstract. The β-carboline alkaloids are plant substances that exhibit a wide spectrum of neuropharmacological, psychopharmacological and antitumor effects. In the present study, we found that harmol, a β-carboline alkaloid, induced autophagy and suppression of survivin expression, and subsequently induced apoptotic cell death in U251MG human glioma cells. Autophagy was induced within 12 h by treatment with harmol. When treated for over 36 h, however, apoptotic cell death was induced. Harmol treatment also reduced survivin protein expression. Small interfering RNA (siRNA)-mediated knockdown of survivin enhanced the harmol-induced apoptosis. Knockdown of survivin by siRNA also induced autophagy. Therefore, harmol-induced apoptosis is a result of the reduction in survivin protein expression. Treatment with 3-methyladenine (3-MA) in the presence of harmol did not affect the expression of survivin and diminished harmol-induced cell death. Treatment with chloroquine in the presence of harmol did not suppress the reduction of survivin expression and increased harmol-induced cell death. From these results, harmol-induced reduction of survivin expression was closely related to autophagy. It is assumed that when isolation membrane formation is inhibited by treatment with 3-MA, reduction of survivin protein expression and apoptotic cell death were not induced. However, when isolation membrane formation is started and an autophagosome is formed, survivin expression is suppressed and apoptosis is executed. Harmol treatment reduced phosphorylation of Akt, mammalian target of rapamycin (mTOR) and its downstream targets p70-ribosomal protein S6 kinase and 4E-binding protein 1, resulting in induction of autophagy. Conversely, activation of the Akt/mTOR pathway inhibited harmol-induced autophagy and cell death. These findings indicate that harmol-induced autophagy involves the Akt/mTOR pathway. Taken together, autophagy induced by harmol represented a pro-apoptotic mechanism, and harmol suppressed the expression of survivin and subsequently induced apoptosis.

Introduction

Gliomas are the most common type of malignant brain tumors and are resistant to many types of treatments, including chemotherapy, radiation and other adjuvant therapies. Patients with the most malignant histopathologic subtype, glioblastoma, present with the worst prognosis, with a median survival length of less than one year, despite aggressive surgery with adjuvant radiotherapy and chemotherapy (1,2). In addition, glioma cells are prone to acquire drug resistance (3). Currently, there is still a need to identify chemotherapeutic agents with cytotoxic effects exclusively targeted against malignant glioma cells. Improved chemotherapeutic regimens and other strategies are urgently needed.

‘Apoptosis’, signifying genetically controlled programmed cell death (PCD), not only plays a crucial role during tissue development and homeostasis, but is also involved in a wide range of pathologies (4). Since the 1960s, various morphological forms of PCD have been recognized. Clarke classified cell death into 4 types, including type I PCD (apoptosis) and type II PCD (autophagy) (5). Evasion of apoptosis is a hallmark of most malignant cells and contributes to the insensitivity to various current cancer therapies (6). Numerous studies have demonstrated that most chemotherapeutic agents and certain naturally occurring compounds induce cell death by activating the apoptotic pathways. It is thought that apoptosis induction in tumor cells, with either drugs or natural products, is an effective therapy for cancer and immune system diseases.

Autophagy is one of the major regulatory mechanisms in the degradation of intracellular proteins and organelles (7,8). It is a highly conserved evolutionary process which occurs in the cells of all eukaryotic organisms, from yeasts to humans. During autophagy, the cytosol and entire organelles become encased in double-membrane-bound vacuoles (autophagosomes), and subsequently fuse with lysosomes to form autolysosomes and are eventually degraded by lysosomal hydrolases (7). Although originally characterized as a survival response to nutrient deficiency, autophagy is now recognized as being frequently induced in response to a variety of stressors to maintain cellular homeostasis (9-11). The importance of
Autophagy has been emphasized in various biological fields, including cancer (10,12,14). Furthermore, cancer cells undergo less autophagy than normal cells (15,16). These findings indicate that autophagy induction is an attractive modality of anticancer therapy.

β-carboline alkaloids are present in several medicinal plants, including *Peganum harmala*, *Passiflora incarnata*, and *Banisteriopsis caapi* (17,18). These plants have been used in traditional medicine to treat asthma, jaundice, lumbago, and other human ailments (18-20). The β-carboline alkaloids are also found in common plants (e.g., wheat, rice, soybeans, grapes) and plant-derived drinks (e.g., wine, beer, whisky, brandy) (21). It is known that they also exist in mammalian tissues (22,23). It has been reported that certain β-carboline alkaloids and their related compounds exhibit cytotoxic effects on cancer cells (24-26). We also previously reported that harmol, a β-carboline alkaloid, induced apoptosis (27) and autophagic cell death in human non-small cell lung cancer cells (28).

In the present study, we investigated the anticancer effect of harmol on the U251MG human glioma cell line. Furthermore, we examined the types of cell death which were induced by harmol and the possible mechanisms.

**Materials and methods**

**Chemicals.** Harmol (purity, minimum 98%) (Fig. 1), dimethyl sulfoxide (DMSO), 3-methyladenine (3-MA) and chloroquine (CQ) were purchased from Wako Pure Chemical Industries, Ltd. (Osaka, Japan). Harmol was dissolved in DMSO at a concentration of 200 mM, and stored at -20°C until use. Monodansylcadaverine (MDC) was obtained from Sigma-Aldrich (St. Louis, MO, USA). Caspase substrates Ac-DEVD-7-amino-4-trifluoromethyl coumarin (AFC) (caspase-3), Ac-IETD-AFC (caspase-8), and Ac-LEHD-AFC (caspase-9) were obtained from MBL (Nagoya, Japan). A recombinant full-length human active Akt1 protein (rAkt1) was purchased from R&D Systems, Inc. (Minneapolis, MN, USA).

**Cell lines and culture conditions.** Human glioma cell line U251MG was obtained from the Human Science Research Resources Bank (Osaka, Japan). This cell line was cultured in Dulbecco's modified Eagle's medium (Wako) with 80 mg/l kanamycin sulfate (Wako) and heat-inactivated fetal bovine serum (Biowest, Miami, FL, USA) and maintained at 37°C in an incubator containing 95% air and 5% CO₂.

**Cell viability.** Cell viability was assessed with the CellTiter-Blue cell viability assay kit (Promega Corp., Madison, WI, USA). In viable cells, resazurin, which is contained in the CellTiter-Blue cell viability assay reagent, is metabolized to the fluorescent product resorufin. The tumor cells (4.5x10⁵ cells/well) were precultured in a 48-well flat-bottom microtiter plate overnight at 37°C in a 5% CO₂ humidified chamber. Then, various concentrations of harmol were added and the cells were incubated for 24-48 h. After incubation, 100 µl of CellTiter-Blue assay reagent (Promega Corp) was added to each well and the cells were further incubated for 1.5 h. After incubation, the medium of each well was analyzed using a Powerscan HT microplate reader (DS Pharma Biomedical Co., Osaka, Japan) at an excitation wavelength of 560 nm and an emission wavelength of 590 nm. Cell viability was determined based on the fluorescence intensity of non-treated cells.

**Assay for caspase activity.** Caspase activity was measured using fluorogenic peptide substrates. Both untreated control cells and cells which had been treated with 75 µM harmol were washed with ice-cold phosphate-buffered saline (PBS) and suspended in lysis buffer [100 mM HEPES (pH 7.5), 150 mM NaCl, 1% Nonidet P-40, 1 mM EDTA, 1 mM dithiothreitol (DTT), 1 mM phenylmethylsulfonfluoride, 10 µM leupeptin and 1 µM pepstatin] for 20 min on ice, followed by centrifugation at 12000 x g for 10 min at 4°C. In total, 50-70 µg of protein in 40 µl of buffer solution was mixed with 10 µl of 5 mM fluorogenic report substrate, for each individual caspase: Ac-DEVD-AFC for caspase-3, Ac-IETD-AFC for caspase-8 and Ac-LEHD-AFC for caspase-9. The reaction mixture was added to 50 µl of assay buffer [20 mM HEPES (pH 7.4), 0.1 M NaCl, 5 mM DTT, 0.1% Nonidet P-40], and incubated at 37°C for 1 h. The enzymatic product, AFC, which was released from the substrate, was excited at 400 nm to measure its emission at 505 nm. Untreated cells were used as a control.

**Visualization of MDC-labeled vacuoles in harmol-treated A549 cells.** U251MG cells were treated with 100 µM harmol for 0-12 h at 37°C, and then further treated with 30 µM MDC for 10 min. After treatment, cells were washed 3 times with PBS and immediately observed by fluorescence microscopy (BZ-8100, Keyence Co., Osaka, Japan).

**Electron microscopy.** Cultured U251MG cells with and without 100 µM harmol treatment were rinsed with phosphate buffer (PB) (0.1 M, pH 7.4). Cells were then fixed with 2.5% glutaraldehyde in PB for 3 h at 4°C, rinsed with PBS for 5 min at least 3 times, and post-fixed with 1% osmium tetroxide for 1 h at 4°C, followed by dehydration in a graded ethanol series for 5 min at room temperature (RT). After dehydration with absolute ethanol, cells were treated with a mixture of absolute ethanol and Spurr resin (1:1, v/v) for 1 h at RT. Then, the mixture was replaced with pure Spurr resin and cells were treated for 1 h at RT, twice. The pellets were transferred to a BEEM capsule, overlaid with new resin and subjected to resin polymerization at 70°C for 16 h. Ultrathin sections were cut with an Ultracut E microtome (Reichert-Jung Co., Vienna, Austria), stained with uranyl acetate and lead citrate, and examined using a JEM-1230 transmission electron microscope (JEOL, Tokyo, Japan).
Western blot analysis. Whole cell proteins were isolated from untreated and harmol-treated U251MG cells. After treatment, the cells were centrifuged at 300 x g for 5 min, and the pellet was lysed in a buffer containing 25 mM HEPES (pH 7.4), 150 mM NaCl, 0.5% Triton X-100, 10% glycerol, 1 mM DTT, 1 mM sodium orthovanadate, 25 mM β-glycerophosphate, 1 mM NaF and 5 µl/ml protease inhibitor cocktail (Wako). The protein content of each lysate was determined using a BCA protein assay kit (Pierce Biotechnology Inc., Rockford, IL, USA). Protein lysates were then mixed with an equal volume of gel loading buffer [20% glycerol, 4% sodium dodecyl sulfate (SDS), 100 mM Tris, 5% β-mercaptoethanol and 0.01% bromophenol blue] before being boiled for 5 min. After boiling, 40 µg of protein was subjected to SDS-polyacrylamide gel electrophoresis. Proteins were then transferred onto a polyvinylidene fluoride membrane (PVDF; GE Healthcare UK Ltd., Buckinghamshire, UK). Blots were then blocked for 1 h at RT in 5% non-fat dry milk diluted in Tris-buffered saline supplemented with 0.1% Tween-20 (TBS-T) (ICN Biomedicals Inc., Aurora, OH, USA). The following primary antibodies were incubated overnight at RT in TBS-T as follows: rabbit anti-poly-(ADP-ribose)-polymerase (PARP) (Cell Signaling Technology Inc., Danvers, MA, USA; 1:1000), rabbit anti-Akt (Cell Signaling Technology; 1:1000), rabbit anti-phospho-Akt (p-Akt) (Cell Signaling Technology; 1:1000), rabbit anti-mammalian target of rapamycin (mTOR) (Cell Signaling Technology; 1:1000), rabbit anti-phospho-mTOR (p-mTOR) (Cell Signaling Technology; 1:1000), rabbit anti-70 kDa ribosomal protein S6 kinase (P70S6K) (Cell Signaling Technology; 1:1000), rabbit anti-phospho-P70S6K (p-P70S6K) (Cell Signaling Technology; 1:1000), rabbit anti-eukaryotic translation initiation factor 4E-binding protein 1 (4E-BP1) (Cell Signaling Technology; 1:1000), mouse anti-microtubule-associated protein 1 light chain 3 (LC3) (MBL; 1:1000), and mouse anti-β-actin (Sigma-Aldrich; 1:3000). Peroxidase-conjugated secondary antibodies were incubated for 1 h at RT as follows: goat anti-mouse IgG (Jackson Immunoresearch Laboratories Inc., West Grove, PA, USA; 1:5000) or goat anti-rabbit IgG (Cell Signaling Technology; 1:2000). Blots were then washed with TBS-T and developed using Immobilon chemiluminescent substrate (Millipore Corp., Billerica, MA, USA).

Survivin siRNA transfection. Knockdown of survivin expression in U251MG cells was achieved by transfection of siRNA. The control and survivin siRNAs (Cell Signaling Technology) were diluted to a final concentration of 20 nM in Opti-Mem I (Invitrogen Corp., Carlsbad, CA, USA), and transfection was performed with cells at 40-50% confluency using Lipofectamine 2000 transfection reagent (Invitrogen) according to the manufacturer's protocol.

Statistical analysis. Results are presented as means ± SD. Statistical comparisons of the results were carried out using analysis of variance. A P-value of <0.05 was considered to indicate a statistically significant difference. Differences between the means of control and treated cells were analyzed by Dunnett's test.

Results

Cytotoxic effects of harmol on U251MG cells. Continuous treatment of U251MG human glioma cells with harmol for 24 and 48 h induced growth inhibition and cell death in a time- and dose-dependent manner (Fig. 2). The cell viabilities of U251MG cells exposed to harmol at various concentrations of 12.5, 25, 50, 75, and 100 µM in triplicate for 24 h were 102.3, 99.8, 93.7, 82.3, and 66.7%, respectively. After treatment with harmol for 48 h, the cell viabilities decreased to 98.0, 93.6, 72.5, 38.2, and 17.3%, respectively (Fig. 2).

Changes in caspase activities and poly-(ADP-ribose)-polymerase (PARP) cleavage. To confirm whether the cell death induced by harmol treatment was due to apoptosis, the activities of 3 types of caspases were examined (Fig. 3A). Harmol showed no noticeable effect on the activities of these caspases within the 24-h treatment. However, in the 36-h treatment, activities of caspase-3 and -9 were elevated to 486 and 346%, respectively, while caspase-8 activity showed no significant change. We also examined the effects of harmol on PARP cleavage. PARP is a key participant in DNA base excision repair and in maintaining genome integrity. PARP is well known as an endogenous substrate for caspase-3 and an early marker of apoptosis (29). After treatment with harmol for 6-48 h, the cleaved form of PARP (89 kDa) was detected at 36 and 48 h following treatment of the cell lysate (Fig. 3B). These data indicated that treatment with harmol for over 36 h induced apoptosis in U251MG cells.

Harmol treatment suppresses survivin protein expression. While harmol was shown to induce apoptosis in U251MG cells, the mechanisms of inducing this phenomenon were unclear. Afterwards, we investigated the effect of harmol treatment on the expression of survivin protein, which is a known anti-apoptotic protein. From the results, harmol treatment was shown to suppress the expression of survivin protein in a dose- and time-dependent manner (Fig. 4).
Figure 3. Effects of harmol treatment on caspase activity and cleavage of PARP in U251MG cells. (A) The activities of caspase-3, -8, and -9 in U251MG cells were measured as described in Materials and methods. Each value is expressed as mean ± SD, based on 3 independent experiments. *Significant difference between non-treated control and harmol-treated cells (P<0.001). (B) After treatment with 100 µM harmol for the indicated times, PARP cleavage was examined by western blot analysis.

Figure 4. Harmol treatment decreases survivin protein expression. (A) U251MG cells were treated with different concentrations of harmol for 24 h and subjected to western blot analysis. (B) Time course effect of harmol on survivin protein expression. U251MG cells were treated with 100 µM harmol for the indicated times. The amount of protein loaded in each lane was assessed by stripping and reprobing with a β-actin antibody.

Figure 5. Harmol-induced vacuolization and formation of MDC-labeled vesicles in U251MG cells. (Left panels) U251MG cells were treated or not treated with 100 µM harmol for 0-12 h before phase contrast microscopic observation. (Right panels) After treatment with 100 µM harmol for the indicated times, the cells were further incubated with 25 µM MDC for 10 min. Then, the cells were washed twice with PBS and immediately analyzed by fluorescence microscopy (scale bar, 20 µm).
Formation of autophagic vacuoles following treatment with harmol.

Phase contrast microscopy of harmol-treated U251MG cells showed numerous, high-density vacuoles (Fig. 5, left lower panel). Based on the above findings, we would have expected these vacuoles to be formed upon harmol treatment; they were not detected in untreated cell autophagosomes. The fluorescent compound MDC is a specific marker for autophagosomes (30), and is commonly used to stain autophagic vesicles. We studied the incorporation of MDC in cells where autophagy was stimulated by harmol treatment. U251MG cells were treated with 100 µM harmol for 12 h and then analyzed by fluorescence microscopy. As shown in Fig. 5 (right upper panel), in control cells (0 h treatment), MDC-labeled vacuoles were scarcely detected. On the other hand, in cells which were treated with harmol for 12 h, numerous MDC-labeled fluorescent dots were clearly detected (Fig. 5, right lower panel). Although MDC-labeled fluorescent dots were observed in harmol-treated U251MG cells, in order to identify them more precisely, we performed electron microscopy to obtain ultrastructural information regarding the morphology of harmol-induced autophagy in U251MG cells. Fig. 6 shows an electron micrograph of a non-treated U251MG cell (left panel) and a U251MG cell treated with harmol (right panel). Although the electron micrograph of the harmol-treated U251MG cell showed numerous cytoplasmic phagolysosomes that contained highly electron-dense materials (Fig. 6, right panel), few phagolysosomes were observed in the control cell (Fig. 6, left panel).

Quantitative detection of microtubule-associated protein 1 light chain 3 (LC3). To quantify the incidence of harmol-induced autophagy, we examined the expression of LC3-I and LC3-II proteins using western blot analysis, as these proteins are essential to the formation of autophagosomes (especially LC3-II) and have been widely used for estimating the number of autophagosomes or the incidence of autophagy (31). The expression of LC3-II protein in harmol-treated cells was detected within 12 h, and increased during treatment in a time-dependent manner (Fig. 7). Taken together, these results also indicated that harmol induced autophagy, and that harmol-induced autophagy preceded apoptosis in U251MG cells (Fig. 3 vs. Figs. 5-7).

Role of autophagy in harmol-mediated cell death.

Autophagy plays an important role in the regulation of cell survival or death (7,8). To determine whether the induction of autophagy by harmol is related to survival or death mechanisms, we investigated the effect of harmol on cell survival in the presence of autophagy inhibitors in U251MG cells. As shown in Fig. 8A, 3-MA, a specific inhibitor of the early stage autophagic process (32), reduced the harmol-induced cell death in U251MG cells. On the other hand, the late stage autophagy inhibitor CQ (33) increased the harmol-induced cell death. On western blot analysis, although treatment with harmol alone induced the expression of LC3-II protein (known as an autophagy marker), co-treatment with harmol and 3-MA notably suppressed LC3-II protein expression (Fig. 8B). In contrast, co-treatment with harmol and CQ induced progressive accumulation of LC3-II throughout the 24 h duration assay (Fig. 8B). Following treatment with harmol alone for 24 h, although PARP cleavage was not induced (Figs. 3 and 8B), CQ treatment in the presence of harmol induced PARP cleavage. Regarding the anti-apoptotic protein survivin, although >24 h of treatment with 100 µM harmol strongly suppressed the expression of survivin protein (Fig. 6), the combination treatment of harmol with 3-MA did not suppress the expression of survivin protein (Fig. 8B). However, co-treatment with harmol
Genetic suppression of survivin protein expression increases harmol-induced apoptosis. To demonstrate whether harmol-induced apoptosis is related to the suppression of survivin protein expression, we performed siRNA-mediated knockdown of survivin in U251MG cells. As a result, although in control-siRNA transfected U251MG cells, PARP cleavage was not detected within the 24 h treatment of harmol, in cells where the expression of survivin protein was supressed by transfec

tion with survivin siRNA, PARP cleavage was detected within 24 h following treatment with harmol (Fig. 9B). In addition, siRNA depletion of survivin protein induced autophagy, which was confirmed by LC3-II protein expression (Fig. 9B, the non-treated control of the survivin-siRNA transfected cell sample was compared with the non-treated control of the control-siRNA transfected cell sample). Furthermore, in the cytotoxicity experiment, the suppression of survivin expression by siRNA increased the harmol-induced cytotoxicity (Fig. 9A).

Akt/mTOR signaling pathway is involved in harmol-induced autophagy in U251MG cells. Our next objective was to identify and characterize the molecular pathways involved in harmol-induced autophagy. The classical pathway that regulates autophagy involves the serine/threonine kinase, mTOR (34). The Akt/mTOR/P70S6K, 4E-BP1 pathway is the main regulatory pathway that negatively regulates autophagy. However, several pathways appear to regulate autophagy in mammalian cells. Thus, we investigated whether the autophagy induced by harmol is mediated by an mTOR-dependent or -independent pathway using western blot analysis (Fig. 10). Harmol treatment caused a significant time-dependent decrease in the phosphorylation of Akt protein in U251MG cells. Equally, harmol treatment diminished the levels of the phosphorylated (activated) form of mTOR, a downstream target of Akt which may inhibit cell growth and induce autophagy. Furthermore,
examination of the phosphorylation of 2 downstream effectors of mTOR signaling, P70S6K and 4E-BP1, showed that both p-P70S6K and p-4E-BP1 decreased in a time-dependent manner, revealing a potent inhibitory effect of harmol treatment on the Akt/mTOR signaling pathway.

Activation of the Akt pathway inhibits harmol-induced autophagy and cytotoxicity in U251MG cells. It has been reported that the Akt/mTOR pathway mediates autophagy induced by various anticancer therapies and natural products (35-37). Thus, we used rAkt1 to activate the Akt pathway as described previously (38). In Akt pathway-activated U251MG cells treated with rAkt1, we examined the effect on harmol-induced autophagy and cell death. U251MG cells were treated with 100 µM harmol, 500 ng/ml rAkt1, or both, for 12 h for western blotting. The addition of rAkt1 inhibited the harmol-induced decrease in phosphorylated Akt, mTOR and P70S6K in U251MG cells (Fig. 11A). Furthermore, the addition of rAkt1 significantly decreased harmol-induced autophagy, confirmed by LC3-II protein expression (Fig. 11A). In the cytotoxicity experiment, the addition of rAkt1 significantly suppressed harmol-induced cytotoxicity in U251MG cells (Fig. 11B).

Discussion

Previous studies have demonstrated that harmol induced apoptosis in human lung H596 cells through the caspase-8-dependent pathway, but independently of Fas/Fas ligand interaction (27). Furthermore, harmol induced autophagic cell death in human lung A549 cells (28). In the present study, we found that harmol induced autophagy and apoptotic cell death in U251MG cells.

Defects in the regulation of apoptosis are common phenomena in many types of cancer and are also the critical steps in tumorigenesis and resistance to therapy (39). There are two key signaling mechanisms through which cells may be triggered to undergo apoptosis. In the intrinsic...
pathway, diverse stimuli, such as reactive oxygen species, DNA-damaging reagents, and Ca^{2+} mobilizing agents provoke cell stress or damage, typically by activating one or more members of the BH3-only protein family, leading to the release of cytochrome c. Cytochrome c then binds to apoptosis-activating factor 1 (ApaF-1) and procaspase-9, resulting in the activation of caspase-9 by proteolytic cleavage. The extrinsic pathway starts with death receptor ligation or Fas/Fas ligand interaction, followed by oligomerization of the receptor, use of Fas-associated death domain (FADD) protein, and activation of caspase-8 (40). Caspase-8 and -9 have been generally defined as initiator caspas, and can in turn activate caspase-3, the executor of apoptosis (41,42). As shown in Fig. 3, over 36 h of treatment with harmol increased the activity of both caspase-9 and -3, and induced PARP cleavage. However, no noticeable change in the activity of caspase-8 was observed. From these results, it is assumed that harmol-induced apoptosis was associated with an intrinsic pathway instead of an extrinsic pathway, such as Fas/Fas ligand interaction or the TNF receptor pathway. Although the reasons for the differences in the responses to harmol treatment in U251MG, H596 and A549 cells are unclear, these phenomena in each of these cells might be based on the characteristics of each cell line and the involved stimuli.

Survivin is the smallest member of the inhibitor of apoptosis (IAP) family that is selectively overexpressed in most common types of human cancers, but not in normal adult tissues, and has been implicated in the control of cell division, inhibition of apoptosis, and tumor cell resistance to certain anticancer agents and ionizing radiation (43-48). Therefore, these findings make survivin an attractive anticancer drug target. In U251MG cells, we found that harmol suppressed the expression of survivin protein and induced apoptosis associated with activation of caspases and cleavage of PARP, consistent with the role of survivin in antagonizing caspases, thereby blocking apoptosis. Although treatment with harmol alone induced cell death following a 36-h treatment (Fig. 3), the knockdown of survivin expression by specific siRNA sensitized U251MG cells to harmol-induced cell death. Cleaved PARP, a hallmark of apoptosis, was detected within 24 h (Fig. 9B). From these results, the possibility remains that harmol-induced apoptosis is due to the downregulation of survivin protein expression.

Of note, although harmol treatment induced not only apoptosis, but also autophagy (Figs. 5-7), the suppression of survivin protein expression by siRNA also induced autophagy (Fig. 9B, comparing the LC3-II band of the non-treated control sample of control siRNA with the LC3-II band of the non-treated control sample of survivin siRNA). Furthermore, survivin siRNA treatment increased harmol-induced autophagy, as evidenced by conversion of the autophagosomal marker LC3 protein from the cytosolic form (LC3-I) to the membrane-bound, phosphatidylethanolamine-conjugated form (LC3-II), associated with a size shift detected by western blotting (Fig. 9B). In brief, it appears that at first, harmol induces autophagy (induced within 12 h by treatment with harmol) (Fig. 7) and subsequently reduces the expression of survivin protein. Next, the reduction of survivin protein expression increases autophagy, and finally, apoptosis is induced. Autophagy plays a critical role in the regulation of cell survival or death (12). Therefore, to investigate whether the induction of autophagy by harmol has a cytoprotective function or leads to cell death (i.e., autophagic cell death), we treated U251MG cells with harmol and measured the effects on cell survival or apoptosis in the presence or absence of pharmacological autophagy inhibitors. Although 3-MA treatment significantly reduced harmol-induced cell death, CQ treatment highly enhanced harmol-induced cell death (Fig. 8). In addition, although harmol treatment decreased the expression of survivin, 3-MA treatment in the presence of harmol suppressed the reduction of survivin expression (Fig. 8B). On the other hand, CQ treatment in the presence of harmol did not suppress the reduction of survivin expression (Fig. 8B). The 3-MA molecule, a specific inhibitor of the early-stage autophagic process, is known as a PI3K inhibitor. Class III phosphatidylinositol-3-kinase (class III PI3K) is closely associated with autophagosome membrane formation (10). Therefore, treatment with 3-MA inhibits the formation of autophagic vacuoles. On the other hand, the late-stage autophagy inhibitor CQ blocks the fusion of autophagosomes and lysosomes, thereby suppressing the degradation of the contents within the autophagosome. Considering this information, the possibility remains that when harmol-induced autophagosome formation is inhibited, suppression of survivin protein expression is not induced. In contrast, if an autophagosome is formed, suppression of survivin protein expression occurs, and subsequently apoptosis is induced. In brief, it is assumed that harmol-induced autophagy plays pro-apoptotic roles. However, the detailed mechanisms of these phenomena are unclear. Additional experiments are required to clarify these mechanisms.

Autophagy is regulated by multiple signaling pathways as diverse as class III PI3K (10), protein kinase mTOR (7,49,50), mitogen activated protein kinase (MAPK) (51,52), AMP-activated protein kinase (AMPK) (53,54), Beclin1 (55) and bcl-2 (56). In particular, Akt/mTOR signaling is the major pathway and plays a variety of physiologic roles, including cell growth, cell cycle regulation, migration and survival (57-59). Furthermore, this pathway is also frequently associated with oncogenesis (60). It is known that the Akt/mTOR pathway negatively regulates autophagy (10). Several studies have indicated that inhibition of the Akt/mTOR pathway is associated with the triggering of autophagy in cancer cells (60-63). In U251MG cells, harmol treatment inhibited the phosphorylation of Akt (both Ser473 and Thr308), and moreover, exposure of cells to harmol also inactivated mTOR and reduced phosphorylation of its downstream targets p70S6K and 4E-BP1 (Fig. 10). In contrast, the activation of the Akt pathway by rAkt1 treatment increased the phosphorylation of mTOR and its downstream targets p70S6K and 4E-BP1 (Fig. 11A). Treatment with rAkt1 also improved the viability of harmol-treated U251MG cells (Fig. 11B). These results are consistent with many studies indicating that inhibition of the Akt/mTOR pathway is associated with induction of autophagy in cancer cells (35,36,50,61).

β-carbolines which occur naturally as harmala alkaloids in Peganum harmala Linné, are also widely found in other medicinal plants (17,18), and are present endogenously in mammalian tissues (22,23). Harmala alkaloids have been used in certain hallucinogenic preparations by some South American and African tribes (62). Furthermore, plants containing harmala alkaloids have long been used in traditional medicine
to treat asthma, jaundice, lumbago, and other ailments (18-20). Moreover certain β-carboline alkaloids have a wide spectrum of neuropharmacological and psychopharmacological actions on the central nervous system, such as tremorgenesis (63,64), hypothermia (65), hallucinogenesis (66,67), monoamine oxidase inhibition (68,69), and convulsive or anticonvulsive actions (70). The β-carboline-induced central nervous effects mainly occur with β-carbolines which have a methoxyl group at the C-7 position in their structures (71). Therefore, β-carbolines which have a methoxyl group at the C-7 position are not suitable for chemotherapeutic agents. On the other hand, it is reported that harmol, which has a hydroxyl group at the C-7 position (Fig. 1), showed only slight central depression and convulsive effects (72). Furthermore, harmol has either slight or no tremor-inducing action (73,74).

With regard to its metabolic pathway, harmol is conjugated with sulfate and excreted in the urine and bile; it has a rapid rate of urinary excretion in humans after intravenous administration (75). Therefore, harmol is considered to have low toxicity in humans and animals.

In conclusion, we demonstrated that harmol induces autophagy and apoptosis in U251MG human glioma cells. As these effects of harmol were investigated only in glioma cells, harmol's effects on other types of cancer should also be examined. Furthermore, although additional investigation is needed to precisely clarify the autophagy and apoptosis induction pathway, it is thought to be a candidate pathway that can be targeted by a chemotherapeutic agent in cancer treatment.

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