Ginkgolic acids induce HepG2 cell death via a combination of apoptosis, autophagy and the mitochondrial pathway

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Abstract. Ginkgolic acids may induce malignant cell death via the B-cell lymphoma 2 (Bcl-2)-associated X protein (Bax)/Bcl-2 apoptosis pathway. Concurrently, apoptosis, autophagy and mitochondrial dysfunction may also be involved in bringing about this endpoint. The anticancer effect of Ginkgolic acids (GAs) was investigated using the HepG2 cell line. The median lethal dose of the GAs of the HepG2 was measured via an MTT assay, the dose-response curves were evaluated and changes in cell morphology were monitored by microscopy. Autophagy in HepG2 cells was down regulated using 3-methyladenine (3-MA) or Beclin-1-specific small interfering RNA (siRNA) and the expression of apoptosis associated proteins caspase-3, Bax/Bcl-2, and the autophagy-associated protein 5 and microtubule-associated protein 1A/1B-light chain 3 in the GA-treated HepG2 cells were all measured by western blot analysis. The level of apoptosis in the GA-treated cells was also assessed using terminal deoxynucleotidyl-transferase-mediated dUTP nick-end labeling (TUNEL) assay, and the mitochondrial membrane potential $(\Delta \psi_m)$ was detected by immunofluorescence. The results of the MTT and TUNEL assays indicated that the proliferation of HepG2 cells treated with GAs was significantly reduced compared with the control group, and the rate of the inhibition was dose-dependent. Western blot analysis indicated that treatment with the Gas induced apoptosis and autophagy in the HepG2 cells. The $\Delta \psi_m$ of the GA-treated HepG2 cells was decreased compared with the control, as monitored by immunofluorescence. However, upon the administration of 3-MA or Beclin-1-specific siRNAs (inhibitors of the autophagy), the expression levels of the apoptosis- and autophagy-associated proteins were decreased. In conclusion, the results of the present study indicated that GAs are potent anticancer agents that function through a combination of the apoptosis, autophagy and mitochondrial pathways.

Introduction

In previous years, novel strategies of management for malignant cells in an integrated therapeutic manner, including surgery and radiotherapy (1,2), immune therapy (3,4) and chemotherapy (5,6), have been investigated. Concurrently, increasing rates of chemoresistance in cancer cells against classic chemotherapy agents, including cisplatin (7), cyclophosphamide (8) and hydroxyurea (9), have been observed. Consequently, the cytotoxicity exhibited by ginkgolic acids (GAs) against malignant cells suggests that they have the potential to become a novel antitumor drug (10,11).

Gas are natural components of the leaves, nuts and episperm of the *Ginkgo biloba* plant, and are identified in small concentrations in *G. biloba* cells (12). Natural GAs extracted from *G. biloba* are derivatives of 6-alkyl or 6-alkyl salicylic acid; the number of side chain carbon atoms in the 6th position may range from 13 to 19, and the number of side chain double bonds may be between 0 and 3. Therefore, the group is a mixture of homologues. Gas have been previously demonstrated to exhibit various toxicities, including allergenic (13), cytotoxic (14) and immunotoxic (13) effects. In previous decades, it has also been identified that the Gas exert antibacterial (15), insecticidal (16) and antitumor (17) effects, although the mechanism of their antitumor activity remains unclear.

Autophagy is a conserved process in cells; it supplies basic energy, and amino acids, lipids and glucose molecules for cell survival under stressful conditions including hypoxia and starvation, via the recycling of cellular components (18). Previous studies have demonstrated that autophagy can also induce apoptosis in cells (19,20), yet whether autophagy is a cytotoxic (19,20) or cytoprotective (21) process remains unclear (22). However, a previous study indicated that autophagy is involved in the cell death process, concomitant with other well-known cell death pathways, including apoptosis and mitochondrial dysfunction (19,20). A number of classical chemotherapy agents, including cisplatin (23), cyclophosphamide (24) and hydroxyurea (25), induce mitochondrial

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dysfunction and autophagy. The present study examined a potential autophagy-based anticancer strategy. GAs as novel anti-cancer agents have been indicated to induce cell apoptosis (26,27), yet whether Gas induce autophagy in cells remains unclear. The present study aimed to identify whether GA-induced autophagy and mitochondrial dysfunction could also contribute to the death of cancer cells, and whether there was a cross-talk with apoptosis.

Materials and methods

Materials. The human hepatoblastoma HepG2 cell line (28) was purchased from the Cell Bank of Type Culture Collection of Chinese Academy of Sciences (Shanghai, China). MTT and dimethyl sulfoxide (DMSO) were purchased from Amresco LLC (Solon, OH, USA). 3-methyladenine (3-MA) was obtained from Sigma-Aldrich (Merck KGaA, Darmstadt, Germany). The small interfering RNA (siRNA) targeting the human Beclin-1 gene was purchased from Shanghai GenePharma Co., Ltd. (Shanghai, China). Lipofectamine® 2000 was obtained from Invitrogen (Thermo Fisher Scientific, Inc., Waltham, MA, USA). The rabbit polyclonal anti-caspase-3 (cat. no. PB0183), anti-B-cell lymphoma 2 (Bcl-2; cat. no. A00040-2), anti-Bcl-2-associated X protein (Bax; cat. no. A00183) and anti-Beclin-1 antibodies (cat. no. PB0014) were purchased from Wuhan Boster Biological Technology, Ltd. (Wuhan, China). The rabbit monoclonal anti-microtubule-associated protein 1A/1B-light chain 3 (LC3; cat. no. 2057-1) antibody was obtained from Epitomics, Abcam (Cambridge, MA, USA), horseradish peroxidase (HRP)-conjugated goat anti-rabbit antibody (cat. no. CW0234S) was purchased from CWBio (http://www.cwbiotech.com/; Shanghai, China). The HRP-conjugated mouse β -actin antibody (cat. no. sc-4778) was purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). The polyvinylidene difluoride (PVDF) membrane was purchased from EMD Millipore (Billerica, MA, USA). Dulbecco's modified Eagle's medium (DMEM), trypsin, and EDTA-disodium salt 2 and fetal bovine serum (FBS) were purchased from Gibco; Thermo Fisher Scientific, Inc. A Mitochondrial Transmembrane Potential $(\Delta \psi_m)$ Analysis kit (cat. no. KGA602) was purchased from Nanjing KeyGen Biotech Co., Ltd. (Nanjing, China). All other supplies for cell culture were purchased from Corning Costar (Corning Incorporated, Corning, NY, USA).

The Gas was provided by Dr Xiaoming Yang from the School of Chemistry of Jiangsu University (Zhenjiang, China), with the purity >98% as monitored by a high-performance liquid chromatography. The parameters of HPLC were as following, the Gas were detected by ProStar 240 (Varian Medical Systems, Inc., Palo Alto, CA, USA.) with ODS-2 (4.6*250 mm, 5 μ m; Dalian Elite Analytical Instruments Co., Ltd., Dalian, China) at 40°C. The sample quantity was 10 μ l, and the composition of mobile phase was methanol-3% HAc (92:8 v/v), the flow rate was 1.0 ml/min, and the detection was by 325 type UV-detector (Varian Medical Systems, Inc.) and UV-2450 ultraviolet-visible spectrophotometer (Shimadzu Corporation, Kyoto, Japan.). Reference materials, Gas>99%, were supplied by from Dr Jaggy H of Dr Willmar Schwabe GmbH & Co. KG, Karlsruhe Germany.

Cell culture. HepG2 cells were cultured in DMEM with 10% (v/v) FBS, 100 U/ml penicillin and 100 U/ml streptomycin at 37°C with 5% CO₂ and 100% humidity. When the cells reached 50-70% confluence, Gas was administered to the cells.

GA administration and MTT analysis. When the HepG2 cells were in the exponential growth phase at 50-70% confluence, the cells were harvested and then suspended in DMEM with 5% FBS at a certain concentration $(1x10^4)$. The prepared cells were cultured in 96-well plates for 24 h with 10% FBS at 37°C and 5% CO₂. The next day, cells were incubated with various concentrations of GAs: 0, 1.0625, 2.125, 4.25, 8.5, 17 and 34 μ g/ml. Next, the cells were cultured for an additional 48 h at 37°C. Following this, 20 µl MTT (5 mg/µl) was administered to each well, and incubated for an additional 4 h at 37°C. Subsequently, 150 µl DMSO was added into each well and the cells were shaken for 10 min. Finally, the absorbance was measured in triplicate using a standard spectrophotometer at 490 nm. The dose-depended curve for cell inhibition was calculated based on the data obtained. Simultaneously, the cells in 10 random fields of view were observed by a light microscope (magnification, x100), and the half-maximal inhibitory concentration (IC₅₀) of the Gas in the HepG2 cells was calculated with the formula: IC_{50} =lg-1[Xm-i(Σ P-0.5)], where Xm represents the log of the max concentration of the experiment, i represents the log of each concentration measured in the experiment, ΣP denotes the sum of the rate of inhibition of each group and 0.5 is the empirical constant.

3-MA administration. When the HepG2 cells (~1x10⁴) were at 50-70% confluence in 6-well plates, the cells were treated with 2.5 mM 3-MA. The next day, the Gas were added to the 3-MA-treated or negative control cells at a concentration of 20 μ g/ml for an additional 24 h. Next, the cells were harvested and analyzed by western blot analysis.

siRNA administration. When HepG2 cells in 6-well plates were at 50-70% confluence, the cells were treated with 50 nM Beclin-1-specific siRNA (Shanghai GenePharma Co., Ltd.) using Lipofectamine 2000, according to manufacturer's protocol. The sequences were as follows: Pair 1: Forward, 5'-GGAGCCAUUUAUUGAAACUTT-3' and reverse, 5'-AGU UUCAAUAAAUGGCUCCTT-3'; Pair 2: Forward, 5'-GUG GAAUGGAAUGAGAUUATT-3' and reverse, 5'-UAAUCU CAUUCCAUUCCACTT-3'; Pair 3: Forward, 5'-GCUGCC GUUAUACUGUUCUTT-3' and reverse, 5'-AGAACAGUA UAACGGCAGCTT-3'. After 24 h, Gas were administrated to the siRNA-treated or negative control cells at a concentration of 20 μ g/ml for an additional 24 h. Following this, the cells were harvested and analyzed by western blot analysis as described subsequently.

Western blot analysis. HepG2 cells (~1x10⁴) with or without Gas treatment were lysed in radioimmunoprecipitation assay lysis buffer (Beijing Solarbio Science and Technology Co., Ltd.), with a protease inhibitor cocktail, [1% Trypsin inhibitor (Beijing Solarbio Science and Technology Co., Ltd., Beijing, China; cat. no. A8260), 1% phosphatase inhibitor (Beijing Solarbio Science and Technology Co., Ltd.; cat. no. P1260), and

	1.053	2.125	4.25	8.5	17	34
1.053	-	_	_	P<0.05	P<0.05	P<0.05
2.125	-	-	-	P<0.05	P<0.05	P<0.05
4.25	-	-	-	P<0.05	P<0.05	P<0.05
8.5	P<0.05	P<0.05	P<0.05	-	P<0.05	P<0.05
17	P<0.05	P<0.05	P<0.05	P<0.05	-	P<0.05
34	P<0.05	P<0.05	P<0.05	P<0.05	P<0.05	-
-, no significa	nt difference.					

Table I. The P-value of every two groups for Fig. 1A.

1% PMSF (Beijing Solarbio Science and Technology Co., Ltd.; cat. no. P8340)], according to the protocol of the manufacturer. The protein concentration was detected by a BCA kit (Thermo Fisher Scientific, Inc.). Equivalent amounts of protein $(100 \,\mu g)$ from every sample were separated by 10% SDS-PAGE, then separated by electrophoresis, and then transferred to a PVDF membrane. Next, the membrane was blocked with 5% bovine serum albumin in TBS (Gibco; Thermo Fisher Scientific, Inc.) for 2 h at 37°C. The membrane was incubated at 37°C for 2 h with primary antibodies (dilutions, 1:200) against the specific proteins, and then incubated with HRP-conjugated secondary antibodies (dilutions, 1:5,000) at 37°C for 2 h. The membranes were then washed 3 times for 10 min each. Then, the protein bands were scanned using a Typhoon 9400 Variable Mode Imager (Amersham; GE Healthcare, Chicago, IL, USA) and detected using Pierce Electrochemiluminescence Plus Substrate (Thermo Fisher Scientific, Inc.). The software used for densitometric analysis was Lane 1D (version 4.0.00.001; Beijing Sage Creation Science Co., Ltd).

 $\Delta \psi_m$ analysis. The $\Delta \psi_m$ of HepG2 cells was detected using JC-1 dye. HepG2 cells (~1x10⁴) were incubated in 24-well plates, and treated with Gas at a concentration of 20 μ g/ml for 24 h. Next, following the JC-1 manufacturer's protocol (2.5 μ g/ml at 37°C for 30 min; Nanjing KeyGen Biotech Co., Ltd.), the cells were stained. The cells were monitored by fluorescence microscopy (magnification, x200).

Terminal deoxynucleotidyl-transferase-mediated dUTP nick-end labeling (TUNEL) assay. The HepG2 cells (~1x10⁴) were treated with Gas at a concentration of 20 μ g/ml in the logarithmic growth phase for 24 h, and then fixed with 4% paraformaldehyde at 37°C for 24 h. Next, following to the protocol of the manufacturer of the TUNEL kit (cat. no. KGA702; Nanjing KeyGen Biotech Co., Ltd.), these cells were stained and monitored by a microscope. The mounting medium used was glycerol, and the cells in 10 random fields of view were observed with a light microscope (magnification, x200).

Statistical analysis. Data are presented as the mean ± standard deviation. Data analyses were performed using one-way analysis of variance with Student Newman-Keuls post-hoc test (>3 groups) or Student's t-test (two groups) in SPSS v.23 software (IBM Corp., Armonk, NY, USA). P<0.05 was

considered to indicate a statistically significant difference. All experiments were repeated at least 3 times.

Results

GAs inhibit cells proliferation. The dose-dependent curves for GAs in HepG2 cells were measured by MTT assay following 24 h Gas treatment. HepG2 cell viability decreased with the increases in GAs concentration and incubation time. When the cells were treated with GAs, the optical density (OD), increased in the GAs treatment groups as the GAs concentration decreased, and at the 4.25 μ g/ml, the dose-depended curve reached the threshold (4.25 μ g/ml) compared with the control group. Next, the inhibition ratio of the GAs increased significantly (P<0.05, Table I). Simultaneously, the inhibition rate also increased over time following Gas treatment. Therefore, the inhibition rate caused by GAs was time- and dose-dependent. The OD values of the HepG2 cells treated with GAsfor 24 hare depicted in Fig. 1A. Using the formula IC_{50} =lg-1[Xm-i(Σ P-0.5)], where Xm denotes the log of the max concentration of the experiment, i represents the log of each concentration measured in the experiment, ΣP represents the sum of the rate of inhibition of each group and the 0.5 is an empirical constant, the IC_{50} of Gas was calculated to be 25 μ g/ml. Additionally, the morphological changes in the treatment cells were monitored by microscopy. As depicted in Fig. 1B, an increased number of cells in the GAs treatment group exhibited an increased rate greater number of characteristic vacuolation compared with the control group.

GAs induce apoptosis in HepG2 cells. The TUNEL assay was used to monitor the apoptosis in HepG2 cells (Fig. 2A). The level of apoptosis in the GA groups was markedly increased compared with the control group (P<0.05). Apoptosis of the HepG2 cells was also monitored by western blotting analysis. The expression levels of the apoptosis-associated proteins caspase-3, Bcl-2 and Bax in the HepG2 cells were detected at 24 h after GA treatment. The expression levels of caspase-3, Bax and Bcl-2 were increased in the GA-treated groups compared with the control group (P<0.05); whereas the Bax/Bcl-2 ratio exhibited a peak when the concentration of GAs was 20 μ g/ml (Fig. 2B and C). And the number of apoptotic cells in the GAs groups was increased, compared with that in the control group (P<0.05), as demonstrated in Fig. 2C



Figure 1. GAs inhibit HepG2 cells proliferation. (A) Dose-depended curve of HepG2 cells following treatment with Gas for 24 h. The cells treated with Gas for 24 h exhibited a significantly increased level of inhibition of HepG2 cells compared with the control group (P<0.05 vs. control group). (B) Images of HepG2 cells treated with GAs at 24 h captured by light microscopy (magnification, x200). (Ba) control group and (Bb) HepG2 cells treated with GAs. The cells treated with GAs were decreased in size compared with those in the control group. GAs, ginkgolic acids (the P-value of each two groups depicted in Table I).



Figure 2. Apoptosis and its associated proteins expression levels in HepG2 cells at 24 h following treatment with GAs. (A) Apoptosis of GAs-treated cells, as measured by terminal deoxynucleotidyl-transferase-mediated dUTP nick-end labeling staining. The proportion of apoptotic GA-treated cells was markedly higher than the control-treated cells. (B) The expression of apoptosis-associated proteins in HepG2 cells was detected by western blot analysis. The expression of caspase-3, Bax, and Bcl-2 in the GA-treated groups (various concentrations) was increased compared with that in the control group, and the caspase-3, Bax and Bcl-2 protein expression levels increased as the concentration of Gas increased. The Bax/Bcl-2 ratio reached its peak in the20 μ g/ml group, which was close to the median lethal dose of GAs. (C) Quantification of the apoptosis-associated proteins bands in (B) (*P<0.05). GAs, ginkgolic acids; Bcl-2, B-cell lymphoma 2; Bax, Bcl-2-associated X protein.





Figure 3. Gas induce autophagy inHepG2 cells. The expression of the autophagy-associated proteins Beclin-1 and LC3 in the GAs group significantly differed compared with that of the control group (*P<0.05). GAs, ginkgolic acids; LC3, microtubule-associated proteins 1A/1B light chain 3.



Figure 4. GAs induced a decrease in the mitochondrial membrane potential of HepG2 cells. An increased number of dysfunctional mitochondria were present in the group treated with Gas compared with the control group. GAs, ginkgolic acids.

GAs induce HepG2 cell autophagy. As the GA-treated HepG2 cells exhibited morphological changes such as vacuolization, it may be that autophagy is involved in this process. The autophagy activity of cells with or without administration of Gas was detected. The ratio of LC3-II/ LC3-I is a good indicator of autophagy activity (29), and in the preliminary experiments of the present study, it was identified that it took ~24 h for significant evidence of vacuolization following treatment of the HepG2 cells with 20 μ g/ml GAs (data not shown). Therefore, 12 h was the time-point used for the ratio

of LC3-II to LC3-I in the cells at 12 h following GAs administration, the Beclin-1 protein level was also analyzed as an additional autophagy marker. As depicted in Fig. 3, the expression of the autophagy-associated proteins Beclin-1 and LC3 in the HepG2 cells of the GA-treated group was significantly higher than in the control groups (P<0.05).

GAs induce mitochondrial dysfunction in HepG2 cells. Mitochondria are involved in apoptosis (30); therefore, the $\Delta \psi_m$ of the 20 µg/ml GA-treated HepG2 cells and the control HepG2 cells was monitored using JC-1 staining. If the cells



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Figure 5. Down regulation of autophagy contributes to alterations in the expression of apoptosis-associated proteins in HepG2 cells following decreasing treatment with GAs. (A) Expression of autophagy-associated proteinsBeclin-1 and LC3 following3-MA treatment. The expression levels of the autophagy-associated proteins Beclin-1 and LC3 were decreased in the cells that underwent3-MA treatment compared with the control group (*P<0.05). (B) The expression of the apoptosis-associated proteins caspase-3 and-9 in the GAs group was increased compared with the GAs+3-MA group and the control group (*P<0.05; the P-value of each two groups are depicted in Table II). (C) Expression of autophagy-associated protein Beclin-1 and LC3 following siRNA treatment. The expression levels of the autophagy-associated proteins Beclin-1 and LC3 were decreased in cells following siRNA treatment, compared with the control group (*P<0.05). (D) The expression of apoptosis-associated proteins caspase-3, -8 and -9 in the GAs group increased compared with those in the GAs+siRNA group or the control group (*P<0.05; the P-value of each two groups are depicted in Table III). GAs, ginkgolic acids; LC3, microtubule-associated protein 1A/1B-light chain 3; 3-MA, 3-Methyladenine; siRNA, small interfering RNA.

exhibited an increase in the intensity of green staining compared with red, this indicated an increased number of dysfunctional mitochondria. It was observed that the green fluorescence intensity increased and the red fluorescence intensity decreased in the cells treated with GAs. The ratio of red to green was significantly decreased in the GAs-treated group compared with the control group (P<0.05), as indicated in Fig. 4.

Autophagy contributes to HepG2 cells apoptosis introduced by GAs. Numerous studies have demonstrated that autophagy can be cytoprotective (18,21) and cytotoxic (19,20) activities. To investigate whether autophagy was involved in the apoptosis process induced by GAs in HepG2 cells, and whether its activity was cytoprotective or cytotoxic, the autophagy inhibitor 3-MA was administered to the GA-treated HepG2 cells. Then, the autophagy-associated protein Beclin-1 was detected, and it was demonstrated that the level of autophagy in the GA-treated HepG2 cells was inhibited (P<0.05) (31,32), as indicated in Fig. 5A. The expression levels of the apoptosis-associated proteins caspases-3, -8 and 9 were monitored, and the results revealed that they were all decreased in the 3-MA-treated GAs HepG2 group compared with the GAs HepG2 group without 3-MA treatment (P<0.05, Table II), as indicated Fig. 5B. As 3-MA is a non-specific autophagy suppressor (33), the Class III phosphoinositide 3-kinase (PI3K-III), an autophagy inducer, and PI3K-I, an autophagy suppressor, pathways were inhibited by 3-MA. The autophagy regulator Beclin-1 was silenced to additionally confirm the results of the 3-MA treatment assay. The efficacy of the siRNA in the HepG2 cells was initially detected by western blot analysis (Fig. 5C). The Beclin-1 protein expression level was decreased significantly in HepG2 cells upon Beclin-1 siRNA treatment (P<0.05). Subsequently, the levels of caspase-3, -8 and -9 in the Beclin-1-silenced HepG2 cells treated with GAs was monitored, and the results confirmed those of the 3-MA treatment experiment, which were that when the rate of autophagy was decreased, the level of apoptosis was decreased (P<0.05, Table III; Fig. 5D).

Discussion

In previous decades, increasing levels of chemotherapy resistance in cancer have been described (34,35). Therefore, an integrated management system for malignant tumors, which includes surgery and radiotherapy (1,2), gene therapy (36) and immune therapy (3), has become a reasonable therapeutic strategy. One potential strategy to overcome resistance is the use of novel antitumor drugs that interact with traditional or novel targets; Gas represent one such class of these potential therapeutics.



Table II. The P-val	ue of every tw	o groups for	Fig. 5B.
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	GAs+3-MA	GAs	Control group
Caspase3			
GAs+3-MA	-	P<0.05	P<0.05
GAs	P<0.05	-	P<0.05
Control group	P<0.05	P<0.05	-
Caspase-8	GAs+3-MA	GAs	Control group
GAs+3-MA	-	-	-
GAs	-	-	-
Control group	-	-	-
Caspase-9	GAs+3-MA	GAs	Control group
GAs+3-MA	-	P<0.05	-
GAs	P<0.05	-	P<0.05
Control group	-	P<0.05	-

Table III. The P-value of every two groups for Fig. 5D.

	GAs+siRNA	GAs	Control group
Caspase3			
GAs+siRNA	-	P<0.05	P<0.05
GAs	P<0.05	-	-
Control group	P<0.05	-	-
Caspase-8	GAs+siRNA	GAs	Control group
GAs+siRNA	-	P<0.05	P<0.05
GAs	P<0.05	-	P<0.05
Control group	P<0.05		-
Caspase-9	GAs+siRNA	GAs	Control group
GAs+siRNA	-	P<0.05	-
GAs	P<0.05	-	P<0.05
Control group	-	P<0.05	-

-, no significant difference; GAs, ginkgolic acids; siRNA, small interfering RNA.

GAs have been used previously as antitumor agents in preclinical studies (14,37). Its potential application as an anti-cancer drug is due to its effects on tumor cell apoptosis (38), and previous studies have indicated that Gas exhibit marked antitumor potency. In the present study, multiple mechanisms of GAs-induced cytotoxicity were elucidated, including mitochondria land apoptotic pathways.

To the best of our knowledge, the present study demonstrated for the first time that autophagy is activated in the Gas-treated HepG2 cells, and that potential crosstalk exists between apoptosis and autophagy, in gastric carcinoma cells exposed to GAs. This result is supported by the following observations: Firstly, morphological changes in theHepG2 cells treated with Gas indicated vacuolization (39); secondly, LC3-II/LC3-I are molecular chaperones in autophagy that are involved with autophagy membrane elongation and closure (40), and when HepG2 cells were infected with GAs in the present study, the expression level of LC3-II/LC3-I was significantly increased; thirdly, suppression of autophagy by the chemical 3-MA reduced the levels of GAs-induced autophagy and apoptosis; fourthly, knockdown of Beclin-1 with a specific siRNA decreased the GA-induced LC3-II conversion and the expression of caspase-3, -8 and-9, indicating that the induction of autophagy may result in crosstalk with apoptosis in the GAs-treated HepG2 cells. In addition, it was observed that the mitochondrial pathway was involved in the GA-induced HepG2 cell death, as the $\Delta \psi_m$ decreased in the Gas-treated cells in comparison with the control group These experimental data indicate that the induction of HepG2 cell death was a net effect of the autophagy, mitochondrial and apoptosis pathways. Autophagy triggers apoptosis through the Beclin-1-mediated down regulation of Bcl-2 (41). In previous decades, a number of studies have demonstrated that Bcl-2 is involved in apoptosis and autophagy (30,42-44). Beclin-1 is considered to be a member of the Bcl-2 family as it also contains an N-terminal BH3 domain, and contributes to the activity of Bcl-2 and Bcl-extra large, and inhibits the formation of the autophagosome (44,45). In the present study, western blot analysis revealed that the expression of Beclin-1 increased in GA-treated HepG2 cells in comparison with the control group, and therefore the interaction between Beclin-1 and Bcl-2 decreased, promoting autophagy and apoptosis. In the present study, knockdown of Beclin-1 using the Beclin-1-specific siRNA suppressed the conversion of LC3-I to LC-II, but also decreased caspase-3 protein expression. Consequently, crosstalk between autophagy and apoptosis may have occurred.

It has been demonstrated that the calmodulin proteins, calpains, degrade autophagy protein 5 (ATG-5), which is involved in the autophagy pathway (46). The cleavage of ATG-5 may anchor an amino-terminal production of ATG-5 cleaved by calpain to the mitochondria, causing the release of cytochrome c and followed by mitochondrial dysfunction. In the present study, treatment with GAs caused a reduction

in HepG2 cell mitochondrial trans-membrane potential, demonstrating that mitochondrial dysfunction contributed to GA-induced HepG2 cell death by increasing mitochondrial permeability transition and generating reactive oxygen species. It is also known that mitochondria may introduce apoptosis via the intrinsic pathway (47). Therefore, we hypothesized that treatment with GAs induced HepG2 cell apoptosis via the mitochondrial, apoptosis and autophagy and pathways.

In conclusion, the results of the present study indicated that treatment with Gas induced HepG2 cell death via apoptosis, autophagy and/or mitochondrial pathways. Although the association between autophagy, apoptosis and mitochondrial dysfunction remains unclear, Gas may represent a powerful drug candidate for antitumor therapy.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

QMQ cultured the cells and interpreted the effects of 3-MA and siRNA administration, and was a major contributor in writing the manuscript. YCX, JL, DS and JXD performed the western blotting, and monitored apoptosis and autophagy. SQC, YHL and TCG performed the JC-1 experiments. MBW analyzed and interpreted all the data, and designed the study. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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