Abstract. Overexpression of apoptosis-stimulating of p53 protein 2 (ASPP2) can induce apoptotic cell death in hepatoma cells, which contributes to a killing effect of ASPP2 on treating hepatocellular carcinoma (HCC). In the present study, ASPP2 overexpression failed to induce apoptotic cell death in the HCC Huh7.5 cell line, but promoted autophagy development by inhibiting AKT/mTOR pathway. Inhibition of autophagy using 3-methyladenosine recovered the function of ASPP2 on inducing apoptotic cell death, indicating that ASPP2-induced autophagy has an anti-apoptotic role in Huh7.5 cells. A previous study demonstrated that ASPP2-induced autophagy could induce apoptosis in a CHOP- and DRAM-dependent manner, in which CHOP is involved in the initiation of autophagy and DRAM allows autophagy to induce apoptosis. In the present study, CHOP and DRAM were not involved in ASPP2-induced autophagy; however, the induction of DRAM overexpression recovered the apoptosis-inducing function of ASPP2, indicating that DRAM overexpression switches the role of ASPP2-induced autophagy from anti-apoptotic to pro-apoptotic in Huh7.5 cells. Thus, in combination with DRAM, ASPP2 may better perform its pro-apoptotic role by preventing the occurrence of anti-apoptotic autophagy.

Introduction

Hepatocellular carcinoma (HCC) is one of the most difficult types of cancer to treat. The current therapeutic strategies designed to induce apoptosis are not effective enough to completely eliminate HCC (1). Apoptosis-stimulating of p53 protein 2 (ASPP2) is composed of ankyrin repeats, an SH3 domain, and a proline-rich region (1). ASPP2 binds to p53 through its C-terminus to stimulate the transactivation function of p53 on the promoters of pro-apoptotic genes (1). Other studies also demonstrated that ASPP2 could induce apoptosis in a p53-independent manner (2-4). Previous results have indicated that induction of ASPP2 overexpression can promote apoptotic cell death in hepatoma cells (such as HepG2 or Hep3B), which emphasizes the value of ASPP2 in treating HCC (3,4).

DNA damage regulated autophagy modulator 1 recovers the function of apoptosis-stimulating of p53 protein 2 on inducing apoptotic cell death in Huh7.5 cells

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Materials and methods

Cell culture and treatment. Huh7.5 cells were grown in Dulbecco’s Modified Eagle’s Medium (DMEM; Invitrogen; Thermo Fisher Scientific, Inc., Waltham, MA, USA). DMEM
was supplemented with 10% fetal bovine serum (FBS; Invitrogen; Thermo Fisher Scientific, Inc.). Cells were transplanted with plasmids (5 µg) encoding ASPP2, DRAM and green fluorescent protein-microtubule associated protein 1 light chain 3 (GFP-LC3) by using Lipofectamine 2000 (Invitrogen; Thermo Fisher Scientific, Inc.) for 48 h. Small interfering RNAs (siRNAs; Santa Cruz Biotechnology, Inc., Dallas, TX, USA) were used to decrease the expression of CHOP (cat no. sc-35437) or DRAM (cat no. sc-96209). The control siRNA (cat no. sc-37007) was also purchased from Santa Cruz Biotechnology, Inc. siRNA transfections were performed at 80 nM by reverse transfection with Lipofectamine RNAiMax (Invitrogen; Thermo Fisher Scientific, Inc.) for 48 h, according to the manufacturer’s protocol; 3-methyladenine (3-MA; 10 mM, Santa Cruz Biotechnology, Inc.) was added into the medium of Huh7.5 cells for 12 h at 37°C to inhibit autophagy. Cells were grown on glass cover slips for the TUNEL assay.

Western blot analysis. Cell lysates were subjected to western blot analysis, as described previously (3). Briefly, total cell lysates were separated by 10% SDS-PAGE, and the separated proteins were transferred to polyvinylidene difluoride membranes. The protein blots were blocked with 5% non-fat milk for 1 h at room temperature and sequentially probed with specific primary antibodies and horseradish peroxidase-conjugated secondary antibodies. Primary antibodies and secondary antibodies were used at a dilution of 1:1,000 and 1:5,000, respectively. The detection of specific proteins on the blots was achieved using a Pierce™ ECL Western Blotting Substrate (Thermo Fisher Scientific, Inc.), and the results were captured on ImageQuant LAS 4000 (GE Healthcare, Chicago, IL, USA) using ImageQuant™ TL 7.0 software (GE Healthcare). The antibodies for the detection of LC3-I/II, p62, RAC serine/threonine-protein kinase (AKT), phosphorylated (p)-AKT, mechanistic target of rapamycin (mTOR), p-mTOR were purchased from Cell Signaling Technology, Inc. (Danvers, MA, USA); the antibodies for detection of β-actin, CHOP, DRAM were purchased from Abcam (Cambridge, CA, USA). Anti-ASPP2 antibody was purchased from Sigma-Aldrich (Merck KGaA). Goat antiMouse IgG and goat anti-Rabbit IgG secondary antibodies were purchased from Thermo Fisher Scientific Inc.

Cell death and apoptosis analysis. Cell death was quantified with the calcein acetoxymethyl ester (calcein-AM)/propidium iodide (PI) assay (Thermo Fisher Scientific, Inc.) as previously described (7). Briefly, calcein-AM (1 µg/ml) and PI (1 µg/ml) were added into the supernatant, and the cells were incubated with calcein-AM/PI for 15 min. Apoptosis was detected using the terminal deoxynucleotidyl transferase dUTP nick-end labeling (TUNEL) assay (Promega Corporation, Madison, WI, USA) and the slides were mounted with 50% glycerol, as described previously (7). Briefly, at room temperature, the cells were fixed with 10% paraformaldehyde in PBS for 15 min and then incubated in 1% Triton X-100/PBS for 5 min. TUNEL detection solution was dropped onto the glass cover slips, and the cells were incubated at 37°C in the dark for 60 min. Nuclear staining with DAPI was conducted for 5 min at room temperature following rinsing with PBS. Finally, the slips were mounted with 50% glycerol following rinsing with PBS. The images of the calcein-AM/PI and TUNEL assays were obtained with an Olympus IX71 inverted fluorescence microscope (magnification, x20; Olympus Corporation, Tokyo, Japan). A quantitative cell death/apoptosis analysis was performed by counting >1,000 cells in each examination.

Statistical analysis. Data in the present study are the results of at least three independent experiments and are expressed as the mean ± standard deviation. One-way analysis of variance followed by Tukey’s multiple comparison test was used to evaluate the differences between the groups. P<0.05 was considered to indicate a statistically significant difference.

Results

Overexpression of ASPP2 induces autophagy in Huh7.5 cells. Huh7.5 cells were co-transfected with plasmids encoding GFP-LC3 and ASPP2 (ASPP2-p) for 48 h. A vector plasmid was used as a control for ASPP2 (Ctrl-p). An immunofluorescence assay determined that ASPP2 overexpression induced the development of GFP-LC3-II puncta-positive cells, indicating that ASPP2 overexpression induces autophagy in Huh7.5 cells (Fig. 1A and B). Huh7.5 cells were transfected with ASPP2-p or Ctrl-p for 48 h and then western blotting determined that ASPP2 overexpression also induced the development of LC3-II and reduced the expression of p62 (Fig. 1C). These data indicated that ASPP2 overexpression induced autophagy in Huh7.5 cells.

ASPP2 overexpression induces autophagy by inhibiting AKT/mTOR pathway, but not by inducing the expression of CHOP or DRAM, in Huh7.5 cells. ASPP2 overexpression can induce autophagy by inducing the expression of CHOP and DRAM (4). In the present study, ASPP2 overexpression did not increase the expression of CHOP and DRAM in Huh7.5 cells (Fig. 2A). Additionally, knockdown of CHOP or DRAM using siRNAs did not significantly affect the level of GFP-LC3-II puncta-positive Huh7.5 cells following transfection with ASPP2-p for 48 h (Fig. 2B and C). Thus, in the present study, CHOP and DRAM were not involved in ASPP2-induced autophagy. Mechanistic target of rapamycin (mTOR) has been identified to be a negative regulator of autophagy development (8). When mTOR is phosphorylated by AKT, it is activated and then inhibits autophagy development. Here, western blot analysis determined that ASPP2 overexpression significantly reduced the levels of p-AKT and p-mTOR, indicating that ASPP2 overexpression induced autophagy development by inhibiting the activation of the AKT-mTOR pathway in Huh7.5 cells (Fig. 2A).

Autophagy impairs the function of ASPP2 overexpression on inducing apoptotic cell death in Huh7.5 cells. Huh7.5 cells were transfected with ASPP2-p for 48 h and then TUNEL and Calcin AM/PI assays were used to detect apoptosis and cell death, respectively. Notably, the TUNEL and Calcin AM/PI assays determined that ASPP2 overexpression could not induce apoptosis and cell death in Huh7.5 cells (Fig. 3A and B). Autophagy has been identified to be an anti-apoptotic factor in cells, so whether ASPP2-induced autophagy could impair the pro-apoptotic function of ASPP2 was investigated. An autophagy inhibitor, 3-MA, was used to inhibit autophagy development in
Huh7.5 cells with or without transfection with ASPP2-p. It was determined that 3-MA treatment successfully inhibited basal and ASPP2-induced autophagy in Huh7.5 cells as demonstrated by a significant decrease of GFP-LC3-II puncta-positive cells (Fig. 4A). Next, TUNEL and Calcein AM/PI assays determined that ASPP2 overexpression significantly induced apoptosis and cell death when autophagy was inhibited by 3-MA treatment in Huh7.5 cells (Fig. 4B). These data indicated that, at least in Huh7.5 cells, autophagy impaired the function of ASPP2 overexpression on inducing apoptotic cell death.

**Overexpression of DRAM recovers the function of ASPP2 on inducing apoptotic cell death in Huh7.5 cells.** It was previously demonstrated that the expression of DRAM could give autophagy the ability to induce apoptosis (4). In the present study, Huh7.5 cells were co-transfected with plasmids encoding ASPP2 and DRAM for 48 h to induce the overexpression of ASPP2 and DRAM (Fig. 5A). The results of the TUNEL and Calcein AM/PI assays revealed that overexpression of ASPP2 and DRAM significantly induced apoptotic cell death in Huh7.5 cells, indicating that overexpression of DRAM recovered the pro-apoptotic function of ASPP2 (Fig. 5B). These data indicated that induction of DRAM expression could improve the function of ASPP2 on inducing apoptosis, which may aid the treatment of HCC.

**Discussion**

ASPP2, an inducer of apoptosis, has been reported to inhibit the growth of hepatoma cells in vitro and in vivo (9). The present study demonstrated that ASPP2 overexpression failed to induce apoptosis in Huh7.5 cells, as ASPP2-induced autophagy impaired the function of ASPP2 on inducing apoptosis. These results are different from previous results, which demonstrated that ASPP2 overexpression-induced autophagy could induce apoptosis in HCC Hep1-6, HepG2 and Hep3B cell lines in a CHOP- and DRAM-dependent manner. We hypothesize that the different background of HCC cell lines is a critical factor that affects the function of ASPP2-induced autophagy on inducing or inhibiting apoptosis.

Previous studies demonstrated that ASPP2 is an autophagy inhibitor that impairs the formation of the autophagosome membrane by interacting with Atg5 (10,11). However, a previous study also demonstrated that ASPP2 could induce autophagy development by increasing CHOP expression. CHOP reduces the level of B-cell lymphoma-2 (Bcl-2) and then reduces the formation of Bcl-2-Beclin-1 complexes, which contributes to the increase of free Beclin-1 in the cytoplasm and the initiation of autophagy (4). In the present study, although CHOP is not involved in ASPP2-induced autophagy,
Figure 2. ASPP2 overexpression inactivates the AKT/mTOR pathway and CHOP/DRAM has no effect on ASPP2-induced autophagy. (A) Hun7.5 cells were transfected with ASPP2-p or Ctrl-p for 48 h. Western blotting detection with indicated antibodies. (B) Huh7.5 cells were co-transfected with siRNAs to knock down expression of CHOP/DRAM (upper panel, CHOP-si; lower panel, DRAM-si) and ASPP2-p for 48 h. Western blotting was used to detect the effect of siRNAs treatment on knocking down of CHOP and DRAM. (C) Huh7.5 cells were co-transfected with siRNAs to knock down CHOP/DRAM (left panel, CHOP-si; right panel, DRAM-si) and ASPP2-p for 48 h. Immunofluorescence assay was used to detect the level of GFP-LC3-II-positive cells. The values represent the mean ± standard deviation of three independent experiments. ASPP2, apoptosis-stimulating protein of p53; AKT, RAC serine/threonine-protein kinase; mTOR, mechanistic target of rapamycin; Ctrl-p, control plasmid; siRNA, small interfering RNA; CHOP, C/EBP homologous protein; DRAM, DNA damage regulated autophagy modulator 1; CHOP-si, siRNA targeting CHOP; GFP-LC3-II, green fluorescent protein-tagged microtubule-associated protein 1A/1B light chain 3B.

Figure 3. ASPP2 overexpression fails to induce apoptotic cell death in Huh7.5 cells. (A) TUNEL and (B) Calcein AM/PI were used to detect the levels of apoptosis and cell death in Huh7.5 cells transfected with ASPP2-p for 48 h, respectively. The values represent the mean ± standard deviation of three independent experiments. ASPP2, apoptosis-stimulating protein of p53; TUNEL, terminal deoxynucleotidyl transferase dUTP nick-end labeling; AM, acetoxymethyl ester; PI, propidium iodide.
it was identified that inactivation of the AKT/mTOR pathway also contributes to ASPP2-induced autophagy, indicating that in different situations ASPP2 can induce autophagy through activation of different mechanisms.

DRAM has been identified to induce apoptosis in an autophagy dependent or independent manner (4). In the present study, although DRAM is not involved in ASPP2-induced autophagy, overexpression of DRAM recovers the function of ASPP2 on inducing apoptotic cell death in Huh7.5 cells. To the best of our knowledge, the mechanism by which DRAM induces apoptosis or autophagic apoptosis remains unclear; however, the data in the present study strongly indicated that elucidation of the mechanisms by which DRAM induces apoptosis is critical for treating tumors (4,12).

Autophagy is regarded to serve dual roles on cell death. Autophagy is reported to prevent cells from cell death signals, including nutrition depletion or organelle damage (5). Autophagy could degrade certain cytoplasmic redundant organelle or proteins to provide nutrition for cells (5). The autophagy-mediated degradation of impaired organelles, including uncoupled mitochondria, eliminates the large production of pro-apoptotic inducers, preventing pro-apoptotic factor-initiated cell death (13). However, other studies have demonstrated that autophagy can also be a pro-apoptotic factor, since the inhibition of autophagy reduces the level of apoptosis and the promotion of autophagy has an opposite result (4,12,14). In the present study, ASPP2-induced autophagy had an anti-apoptotic role and the induction of autophagy was associated with the inactivation of the AKT/mTOR pathway.

In fact, although autophagy and apoptosis are involved in maintaining cellular homeostasis and the two physiological functions are regarded to be closely associated with each other, the mechanism by which autophagy induces apoptosis remains, to the best of our knowledge, unclear. The data generated in the present study indicated that the different autophagy-inducing signals may determine the different roles of autophagy on apoptosis; for example, CHOP/DRAM-induced autophagy induces apoptosis, but inactivation of AKT/mTOR pathway induces an anti-apoptotic autophagy.

mTOR is a major target of AKT. When mTOR is activated by AKT via phosphorylation, activated mTOR will inhibit autophagy (15). Previous studies indicate that inhibition of mTOR function by its inhibitors promotes autophagy development and reduces the sensitivity of cells to cell death signals (16,17). Up to now, the mechanism by which ASPP2 inactivates the AKT/mTOR pathway remains unclear, because, to the best of our knowledge, few studies have investigated the associated between the AKT/mTOR pathway and ASPP2. A previous study demonstrated that ASPP2 could bind to AKT in nucleus (3). In the present study, although it is not known whether the interaction between of ASPP2 and AKT could affect the activation of AKT, the data indicated that ASPP2 had the ability to suppress the function of AKT, and overexpression of ASPP2 might be benefit for treating certain tumors with high AKT activation levels.

Taken together, the results of the present study revealed a mechanism by which hepatoma cells could escape from ASPP2-induced apoptosis. These data may prove valuable for the future study of tumor therapy.
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Availability of data and materials

All data generated or analyzed during this study are included in this published article.

Authors' contributions

DL and RL contributed equally to this work, both completed the majority of the laboratory work. XG and LP assisted DL and RL to complete the rest of the laboratory work. DC and KL designed the project. YZ assisted DC and KL to design the experimental work. KL completed the writing of this article.

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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