

# Saikosaponin D inhibits autophagosome-lysosome fusion and induces autophagy-independent apoptosis in MDA-MB-231 breast cancer cells

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**Abstract.** The present study aimed to explore the effect of Saikosaponin D (SSD) and its underlying mechanism on apoptosis and autophagy in human breast cancer MDA-MB-231 cells. MTT assay, flow cytometry, western blotting and confocal fluorescence microscopy detection were employed. SSD, a kind of triterpenoid saponins extracted from *Radix bupleuri*, has been demonstrated to have the effects of anti-inflammatory, antioxidative and anticancer effects and can regulate autophagy. The present study revealed that SSD induced apoptosis through the activation of the p38 mitogen-activated protein kinase (MAPK) signaling pathway in human breast cancer MDA-MB-231 cells. The administration of SSD promoted the phosphorylation/activation of p38 MAPK in MDA-MB-231 cells, whereas pretreatment with SB203580, an effective p38 MAPK inhibitor, attenuated SSD-mediated apoptosis, the cleavage of PARP and the activation of caspase-3. In addition, SSD blocked autophagic degradation by inhibiting autolysosome formation, resulting in the accumulation of autophagosomes. Mechanistically, the results of the present study revealed that SSD inhibited the formation of autophagosomes by inhibiting autophagosome-lysosome fusion, rather than by damaging lysosome function. Furthermore, blocking autophagy degradation was not associated with SSD-mediated apoptosis. The genetic knockdown of autophagy-related protein 5 markedly reduced SSD-mediated LC3B-II accumulation; however, it did not affect the SSD-mediated phosphorylation/activation

of p38, cleavage of PARP, activation of caspase-3 or apoptosis. In conclusion, the findings of the present study suggest that SSD may induce apoptosis and block autophagic degradation, which provides further evidence of the association between the inhibition of autophagic degradation and cell death.

## Introduction

Autophagy is important for the removal of damaged, degenerative, non-functional and aging proteins and organelles, as well as for cellular homeostasis (1). Autophagy dysfunction is associated with various types of disease, such as cardiovascular disease, neurological disease and cancer (2). Autophagy is a continuous process known as the autophagic flux, which includes three stages: Autophagosome formation, autophagosome-lysosome fusion and autolysosome degradation (3). The association between autophagy and cancer is complex and remains poorly understood. Autophagy has been found to lead to the autophagic death of cancer cells, but it also can protect cancer cells from nutritional deficiency, chemotherapy or radiotherapy (4). A previous study also reported that the inhibition of autophagy contributes to anticancer therapy, in particular to the inhibition of autophagosome-lysosome fusion or autolysosome degradation (5). Autophagic degradation inhibitors, such as hydroxychloroquine, liensinine and andrographolide, have also been found to enhance the anticancer activity of multiple anticancer drugs (6-8). However, there are few studies on whether the inhibition of autophagy degradation itself contributes to cell death. Notably, a recent study has reported that classic autophagy degradation inhibitors, such as ammonium chloride chloroquine and bafilomycin, may cause autophagy-independent cell death (9). Saikosaponin D (SSD), a kind of triterpenoid saponin, is extracted from the Chinese traditional herbal, *Radix bupleuri* (10). SSD has been found to inhibit a variety of tumor cells, including ovarian cancer, lung cancer, hepatoma and glioblastoma (11-14). Previously, SSD was also discovered to induce apoptotic cell death through the mitogen-activated protein kinase (MAPK) signaling pathway (14). A previous study also reported that SSD regulated autophagy; SSD induced autophagic cell death through inhibiting sarco/endoplasmic reticulum  $Ca^{2+}$  ATPase

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in apoptosis-defective cells (15). Additionally, SSD suppressed enterovirus A71 infection through inhibiting autophagy (16).

The present study demonstrated that SSD induced apoptosis by activating the p38 MAPK signaling pathway in MDA-MB-231 cells and revealed that SSD potentially blocked autophagic degradation through inhibiting autophagosome-lysosome fusion. Further research identified that blocking autophagy degradation was not associated with SSD-mediated apoptosis. These findings provide additional evidence for the association between inhibition of autophagy degradation and cell death.

## Materials and methods

**Antibodies and reagents.** Antibodies against phosphorylated (p)-mTOR (cat. no. 2971; dilution 1:500), m-TOR (cat. no. 2972; dilution 1:1,000), serine/threonine protein kinase ULK1 (ULK1; cat. no. 8054S; dilution 1:1,000), p-ULK1 (cat. no. 6888; dilution 1:500), p-p38 (cat. no. 9216, dilution 1:500), p-ERK (cat. no. 4376; dilution 1:1,000), ERK (cat. no. 9102; dilution 1:1,000), autophagy-related protein (ATG)5 (cat. no. 12994S; dilution 1:1,000), ATG7 (cat. no. 8558; dilution 1:1,000), p62 (cat. no. 5114; dilution 1:500), JNK (cat. no. 9252; dilution 1:1,000), c-caspase-3 (cat. no. 9661S; dilution 1:200), cathepsin D (cat. no. 2284; dilution 1:200), p-JNK (cat. no. 9251; dilution 1:500) and p38 (cat. no. 9212; dilution 1:1,000) were purchased from Cell Signaling Technology, Inc.; antibodies against lysosome-associated membrane glycoprotein (LAMP)1 (cat. no. sc-20011; dilution 1:1,000), LAMP2 (cat. no. sc-18822; dilution 1:1,000) and cathepsin B (cat. no. sc-13985; dilution 1:500) were obtained from Santa Cruz Biotechnology, Inc.; anti-cleaved (c)-poly (ADP-ribose) polymerase (c-PARP; cat. no. 380374; dilution 1:500) was from Zen-Bio, Inc.; anti-GAPDH (cat. no. AG019; dilution 1:1,000) was purchased from Beyotime Institute of Biotechnology; anti-microtubule-associated protein 1 light chain 3 beta (LC3B; cat. no. L7543; dilution 1:1,000) and anti-Becn1 (cat. no. B6186; dilution 1:500) were obtained from Sigma-Aldrich; Merck KGaA; and horseradish peroxidase-conjugated secondary antibodies (cat. nos. 074-1802, 074-1516 or 074-1802) were purchased from KPL, Inc. The following reagents were used in the present study: SSD (cat. no. A0259; Chengdu Must Bio-Technology Co. Ltd.), bafilomycin A1 (Baf; cat. no. 11038; Cayman Chemical Company), rapamycin (Rapa; cat. no. S1039; Selleck Chemicals), SB230580 (p38 MAPK inhibitor; cat. no. S1076; Selleck Chemicals), LysoTracker™ Red DND-99 (cat. no. L7528; Thermo Fisher Scientific, Inc.) DAPI (cat. no. C1006; Beyotime Institute of Biotechnology) and DMSO (as solvent and placebo; cat. no. D2650; Sigma-Aldrich; Merck KGaA).

**Cell culture.** MDA-MB-231 cells were purchased from the American Type Culture Collection. The MDA-MB-231 cells were cultured in DMEM (cat. no. C11995500bt; Gibco, Thermo Fisher Scientific, Inc.), supplemented with 10% FBS (cat. no. 6140071; Gibco, Thermo Fisher Scientific, Inc.), and maintained in a humidified atmosphere at 37°C and 5% CO<sub>2</sub>.

**MTT cell viability assay.** A total of 5x10<sup>3</sup> cells/well were cultured at 37°C and 5% CO<sub>2</sub> for 24 h in 96-well plates. Following incubation with SSD (0, 2, 4, 6, 10, 12 or 15 μM) for 24 h, 20 μl 5% MTT solution was added to each well and

incubated at 37°C and 5% CO<sub>2</sub> for 4 h. Subsequently, the medium was discarded and the purple formazan was dissolved in 150 μl DMSO/well. The absorbance was detected using a microplate reader (iMark™, Bio-Rad Laboratories, Inc.) at 495 nm. The half maximal inhibitory concentration (IC<sub>50</sub>) was calculated using SPSS 17.0 software (SPSS, Inc.).

**Flow cytometric analysis of apoptosis.** A total of 1x10<sup>5</sup> cells/well were cultured in 6-well plates at 37°C and 5% CO<sub>2</sub> for 24 h. Following incubation with SSD (8 or 10 μM) for 24 h or pretreated with 10 μM SB203580 for 2 h and then exposed to 10 μM SSD at 37°C for 24 h, the cells were collected by centrifuging at 600 x g for 5 min at 4°C, then the cells pellet were resuspended in 1 ml PBS and centrifuged at 600 x g for 5 min at 4°C again. Subsequently the cells pellet were stained with 10 μl propidium iodide (PI; cat. no. P4170; Sigma-Aldrich; Merck KGaA; 5.0 μg/ml) and 5 μl Annexin V-FITC (cat. no. 556419; BD Pharmingen). Apoptotic cells were subsequently detected by flow cytometry (Accuri C6; BD Diagnostics; Becton, Dickinson & Company) and analyzed by FlowJo 7.6.1 software (FlowJo LLC).

**Plasmids and establishment of stable cell lines.** A total of 5x10<sup>3</sup> cells/well were cultured in 24-well plates at 37°C and 5% CO<sub>2</sub> for 24 h, then transfected with 500 ng plasmids for 24 h before the subsequent experimentation using Lipofectamine® 3000 reagent (cat. no. L3000015; Invitrogen; Thermo Fisher Scientific, Inc.) according to the manufacturer's protocol. The plasmids include LAMP1-mGFP plasmid (cat. no. 34831; Addgene, Inc.), tflc3 plasmid (a tandem reporter construct carrying both EGFP-LC3 and mRFP-LC3; cat. no. 21074; Addgene, Inc. and mRFP-LC3 plasmid (cat. no. 21075; Addgene, Inc.). The short hairpin (sh)RNA plasmids of target ATG5 (shATG5; 5'-TTTCATTCAGAAGCTGTTT-3') and non-specific control (shCon; 5'-TTCTCCGAACGTGTCACG T-3') were purchased from Gene Chem Co. Ltd. (Shanghai, China). A total of 1x10<sup>6</sup> 293FT cells cultured in 60 mm plate were co-transfected with lentiviral packaging vectors (cat. no. A43237, Invitrogen; Thermo Fisher Scientific, Inc.) along with 1.5 μg shATG5 plasmid or 1.5 μg shCon plasmid using Lipofectamine® 3000 (cat. no. L3000015; Invitrogen; Thermo Fisher Scientific, Inc.) according to the manufacturer's protocols. Supernatant containing the lentivirus was harvested after 48 h. A total of 1.5x10<sup>5</sup> MDA-MB-231 cells cultured in 6-well were infected with 1 ml lentivirus for 24 h, subsequently knockdown stable cell lines were selected using 4 μg/ml puromycin (cat. no. P9620; Sigma-Aldrich; Merck KGaA).

**Confocal fluorescence microscopy detection.** Cells (5x10<sup>3</sup> cells/well) transfected with LAMP1-mGFP, tflc3 or mRFP-LC3 were treated with 8 μM SSD, 20 nM Baf or 0.25 μM Rapa for 24 h. Lysosomes were stained with 75 nM LysoTracker™ Red DND-99 at 37°C for 2 h. The fluorescence in the cells was detected using a Carl Zeiss LSM 780NLO confocal microscopy with a 63x oil objective.

**Western blotting.** Total protein was extracted from cells using RIPA lysis buffer (cat. no. P0013; Beyotime Institute of Biotechnology). The concentration of protein was determined by the BCA method and 15-60 μg proteins were separated

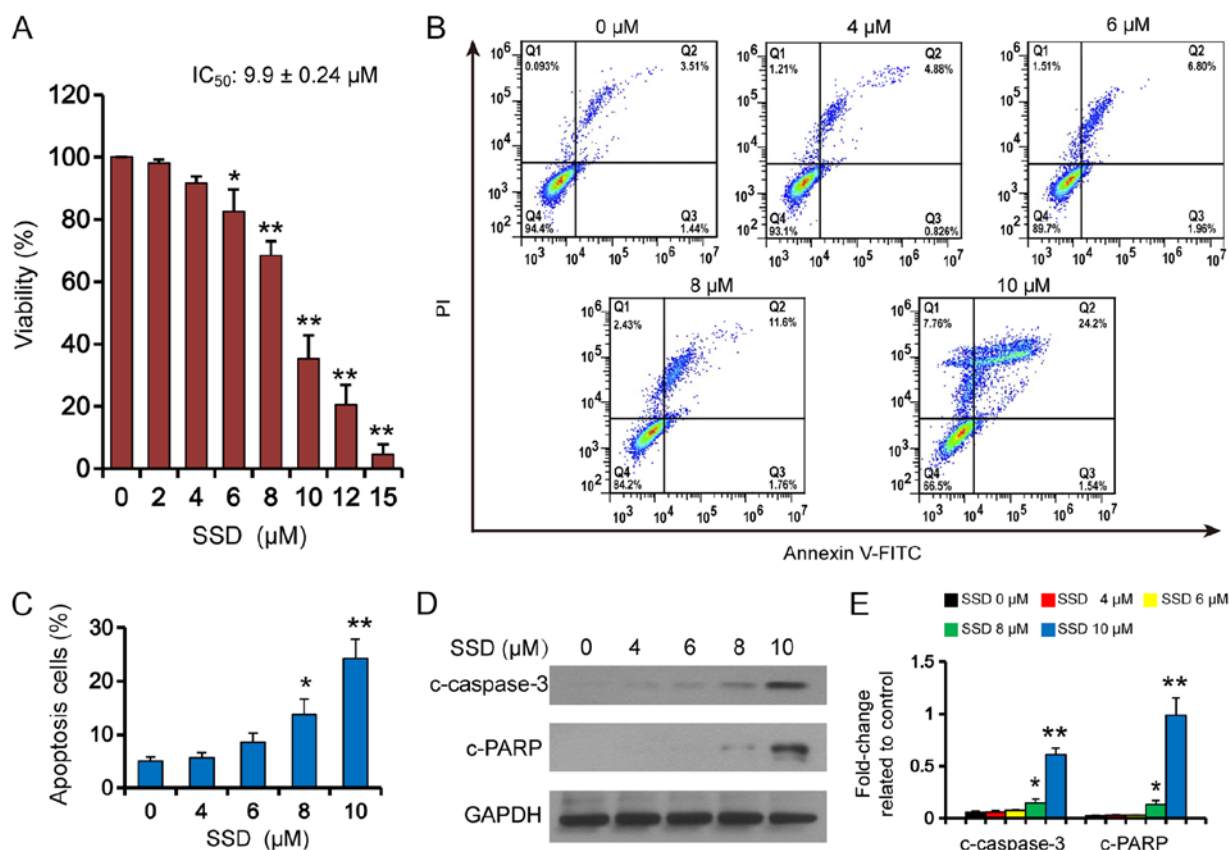


Figure 1. SSD induces apoptosis. MDA-MB-231 cells were exposed to SSD for 24 h and (A) cell viability was examined using an MTT assay (n=3) and (B) the percentage of apoptotic cells was determined by flow cytometry (n=3). (C) Quantitative analysis of the apoptotic rate from data presented in part B. (D) MDA-MB-231 cells were exposed to SSD for 24 h and the expression levels of c-caspase-3 and c-PARP were detected by western blotting (n=3). (E) Semi-quantitative analysis of the expression levels presented in part D. All data are presented as the mean ± SD. \*P<0.05, \*\*P<0.01 vs. 0 μM. SSD, saikosaponin D; c-, cleaved; IC<sub>50</sub>, half maximal inhibitory concentration; PI, propidium iodide; PARP, poly (ADP-ribose) polymerase.

by 10-12% SDS-PAGE. The separated proteins were subsequently transferred onto PVDF membranes and blocked with 5% milk at room temperature for 30 min. The membranes were incubated with the aforementioned primary antibodies at 4°C overnight. Following the primary antibody incubation, the membranes were incubated with the secondary antibodies (1:10,000) at room temperature for 2 h. The membrane was washed three times for 30 min each and total protein was visualized using an ECL substrate (cat. no. 170-5060; Bio-Rad Laboratories, Inc.). The densitometric analysis was performed by ImageJ 1.37C software (National Institutes of Health).

**Statistical analysis.** Data were expressed as the means ± standard deviation from three independent experiments. Statistical analysis was performed using SPSS 17.0 software (SPSS, Inc.). Statistical differences among multiple groups were compared using a one-way ANOVA and Scheffe post hoc test. The statistical differences among two groups were determined using a Student's t-test if the data followed a normal distribution or with a U-Mann Whitney test otherwise. P<0.05 was considered to indicate a statistically significant difference.

## Results

**SSD induces apoptosis in human breast cancer MDA-MB-231 cells.** The cytotoxic effects of SSD were tested in MDA-MB-231

cells using an MTT assay. 6-15 μM SSD significantly inhibited the viability of MDA-MB-231 cells in a dose dependent manner (Fig. 1A); IC<sub>50</sub> was found to be 9.9±0.24 μM. Flow cytometric analysis of apoptosis also demonstrated that 8-10 μM SSD significantly increased the rate of apoptosis in MDA-MB-231 cells (Fig. 1B and C). Consistent with these findings, 8-10 μM SSD was also revealed to promote the cleavage of caspase-3 and PARP (Fig. 1D and E).

**SSD activates the p38 MAPK signaling pathway.** The MAPK cascade is an important signaling pathway involved in the apoptotic process (17). JNK, ERK and p38 MAPK all belong to MAPK signaling pathways (18). To understand whether these MAPKs were involved in SSD-mediated apoptosis, the protein expression levels of p-JNK, JNK, p-ERK1/2, ERK1/2, p-p38 and p38 were detected using western blotting. SSD promoted a significant increase in p-p38 expression levels; however, it did not affect the expression levels of p-JNK or p-ERK (Fig. 2A and B). These results indicated that p38 MAPK may be involved in SSD-mediated apoptosis.

The co-administration of SB203580, a p38-MAPK inhibitor, was found to significantly attenuate the apoptotic rate induced by SSD (Fig. 2C and D), reduce the upregulation of c-caspase-3 and c-PARP expression levels mediated by SSD (Fig. 2E and F), and weaken the inhibitory effect on cell viability mediated by SSD in MDA-MB-231 cells (Fig. 2G).

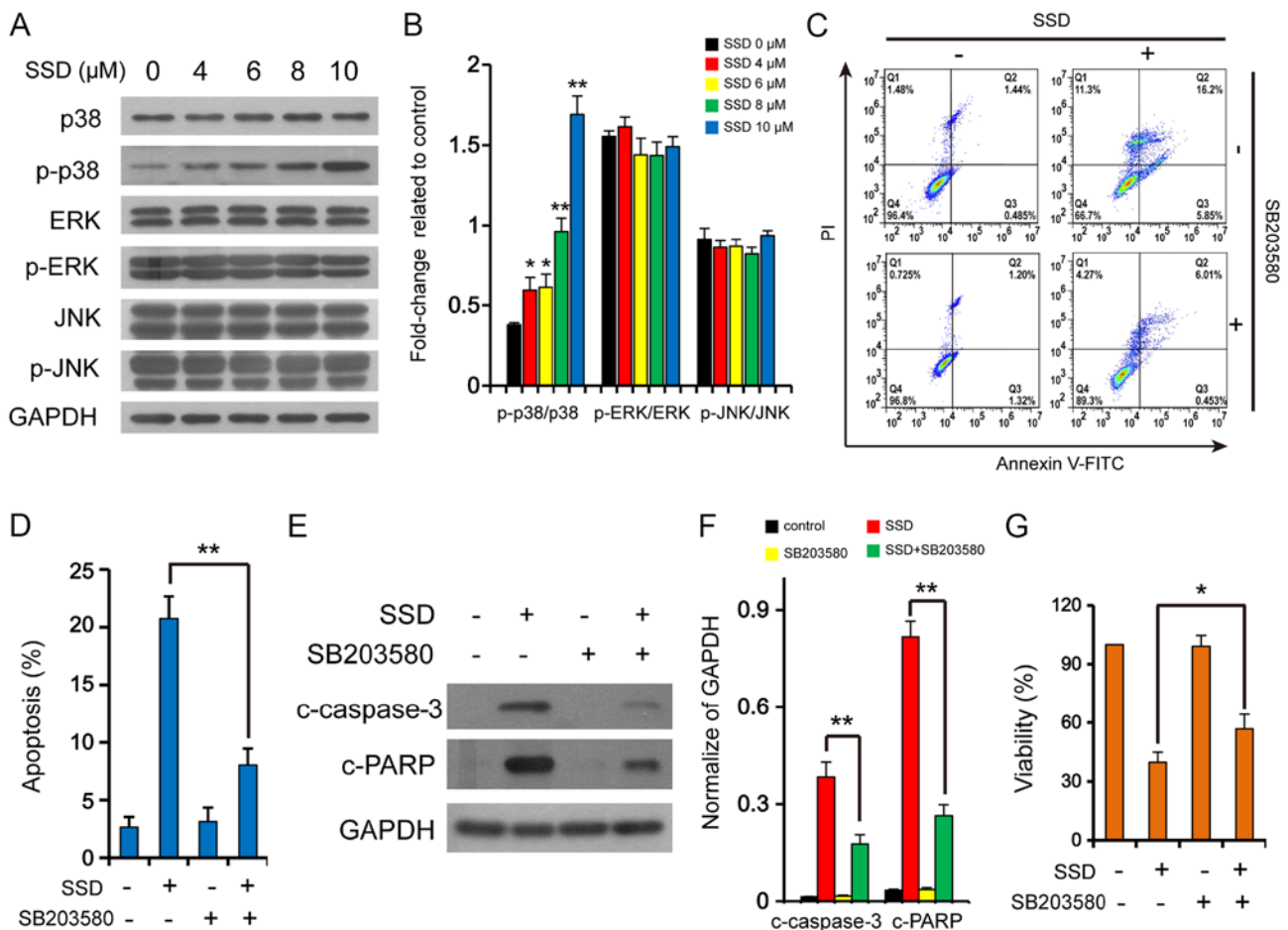


Figure 2. SSD activates the p38 mitogen-activated protein kinase signaling pathway. (A) MDA-MB-231 cells were exposed to SSD for 24 h and the expression levels of JNK, p-JNK, ERK, p-ERK, p38 and p-p38 were detected by western blotting (n=3). (B) Semi-quantitative analysis of the expression levels from part A. \*P<0.05, \*\*P<0.01 vs. 0 μM. MDA-MB-231 cells were pretreated with 10 μM SB203580 for 2 h and then exposed to 10 μM SSD for 24 h. (C) Apoptotic cells were determined using flow cytometry (n=3). (D) Quantitative analysis of the apoptotic rate. \*\*P<0.01. (E) Expression levels of c-caspase-3 and c-PARP were detected using western blotting. (F) Semi-quantitative analysis of the expression levels in part E. \*\*P<0.01 vs. control. (G) Cell viability was determined using MTT assays (n=3). \*P<0.05. Data are presented as the mean ± SD. SSD, saikosaponin D; c-, cleaved; p-, phosphorylated; PI, propidium iodide; PARP, poly (ADP-ribose) polymerase.

These results further suggested that SSD may induce apoptosis through the p38 MAPK signaling pathway.

**SSD inhibits autophagic degradation in MDA-MB-231 cells.** ATGs are important for autophagy (19). The treatment of cells with SSD promoted the accumulation of LC3-II and p62; however, it did not affect the expression levels of Beclin 1, ATG5, ATG7, p-ULK1, ULK1, p-mTOR or mTOR (Fig. 3A and B). These findings indicated that SSD may not promote autophagosome formation but inhibit autophagic degradation (20). To further confirm these observations, tFLC3 plasmid (a tandem reporter construct carrying both EGFP-LC3 and mRFP-LC3) was applied (21). Cells transfected with tFLC3 were treated with Baf, Rapa and SSD and the fluorescence of EGFP-LC3 and mRFP-LC3 puncta were detected using confocal laser microscopy. Both Baf and SSD treatment induced the formation of a large number of both red and green puncta in cells, which displayed a yellow overlay (Fig. 3C). However, only red puncta were detected in cells treated with Rapa. These findings indicated that SSD blocked the process of autophagic degradation.

**SSD blocks autophagic degradation by inhibiting autophagosome-lysosome fusion.** LAMP1 and LAMP2, major lysosomal membrane proteins, are important for maintaining lysosomal function and assisting in autophagosome-lysosome fusion (22). Cathepsin B and D, types of lysosomal cathepsins, are also important for maintaining lysosomal function and the loss of lysosomal function is often accompanied by a decrease in cathepsin expression (23). The expression levels of LAMP1 and LAMP2 were significantly upregulated by SSD treatment compared with untreated cells; however, the expression levels of cathepsin B and D were not affected, suggesting that SSD may not impair lysosomal function (Fig. 4A and B).

The intralysosomal acidic environment is another important factor for lysosomal functions (24). The intralysosomal acidic environment can be detected using LysoTracker™ Red DND-99, an acidic organelle tracer that can emit deep red fluorescence (25). The red fluorescence was observed in cells treated with SSD, Rapa or the placebo, but not in cells treated with Baf. These results indicated that SSD may not affect the intralysosomal acidic environment (Fig. 4C).

Detecting the colocalization of LAMP1 and LC3 can be used to confirm autophagosome-lysosome fusion (26).

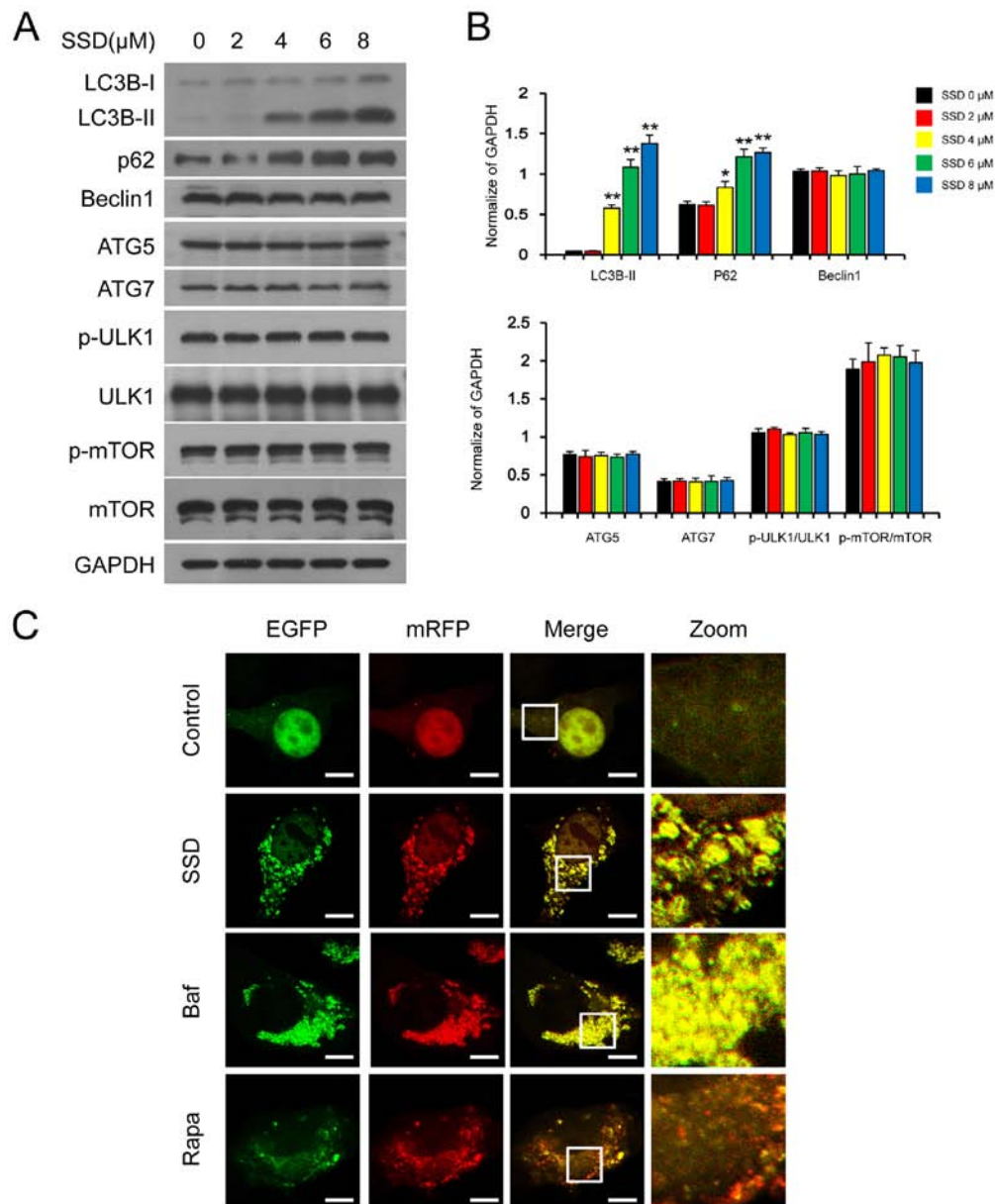


Figure 3. SSD inhibits autophagic degradation. (A) MDA-MB-231 cells were exposed to SSD for 24 h and the expression levels of LC3, p62, Beclin 1, p-mTOR, mTOR, ATG7, ATG5, p-ULK1 and ULK1 were analyzed using western blotting. (B) Semi-quantitative analysis of the expression levels of part A. n=3; \*P<0.05, \*\*P<0.01 vs. 0  $\mu$ M. (C) Cells were transfected with tLC3 and exposed to 8  $\mu$ M SSD, 0.25  $\mu$ M Rapa or 20 nM Baf for 24 h. The EGFP and mRFP puncta were observed using confocal laser microscopy (scale bar=10  $\mu$ m). SSD, saikosaponin D; p-, phosphorylated; ATG, autophagy-related protein; Baf, Bafilomycin A1; Rapa, rapamycin; LC3B, microtubule-associated protein 1 light chain 3  $\beta$ ; ULK1, serine/threonine-protein kinase ULK1.

There was no overlap between the green fluorescence of LAMP1-mGFP and the red fluorescence of mRFP-LC3 in cells treated with SSD or Baf; however, a yellow overlap could be observed in cells treated with Rapa (Fig. 4D), suggesting that SSD may impair the autophagosome-lysosome fusion, resulting in the inhibition of autolysosome formation, which eventually leads to the blockage of autophagic degradation.

*Excessive accumulation of autophagosomes does not contribute to SSD-mediated apoptosis.* To further understand whether the excessive accumulation of autophagosomes contributed to the SSD-mediated apoptosis, the expression levels of LC3B-II were reduced following the stable knock-down of ATG5 expression levels and the treatment with SSD (Fig. 5A and B). Knocking down ATG5 expression levels did not

affect the apoptotic rate of SSD-treated cells (Fig. 5C and D) or the inhibitory effect of SSD on cell proliferation (Fig. 5E) in MDA-MB-231 cells. In addition, the knockdown of ATG5 did not alter the SSD-mediated activation of c-caspase-3, c-PARP or p-p38 (Fig. 5A and B). These results indicated that the excessive accumulation of autophagosomes may not contribute to the SSD-mediated apoptosis in MDA-MB-231 cells.

## Discussion

The present study provided evidence to suggest that SSD may induce apoptosis in MDA-MB-231 cells through the activation of p38 MAPK signaling. In addition, the current results revealed that SSD blocked autophagic degradation through the inhibition of autophagosome-lysosome fusion and that

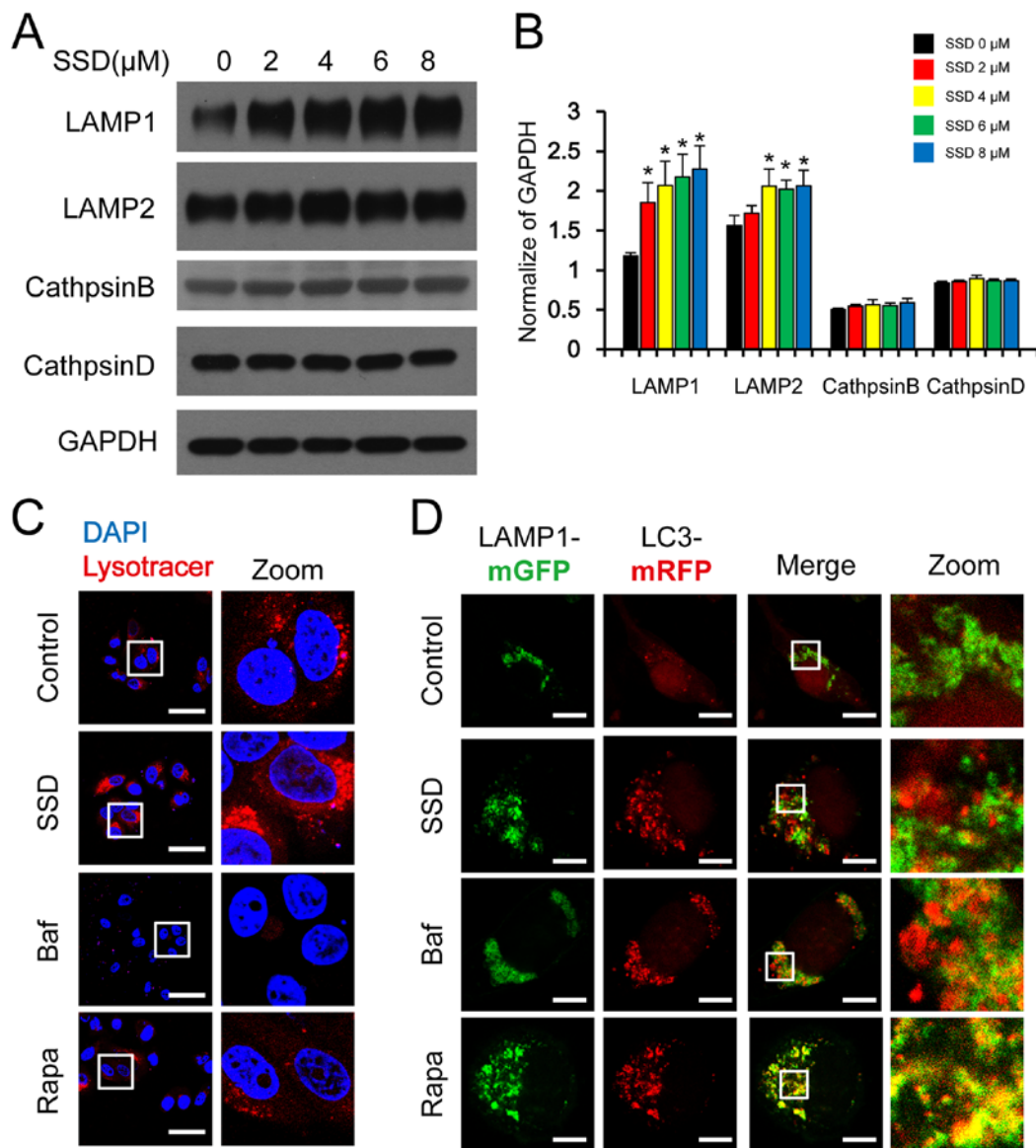


Figure 4. SSD blocks autophagosome-lysosome fusion. (A) MDA-MB-231 cells were exposed to SSD for 24 h. The expression levels of LAMP1, LAMP2, cathepsin B and cathepsin D were analyzed using western blotting. (B) Semi-quantitative analysis of part A. Data are presented as the mean  $\pm$  SD; n=3; \*P<0.05 vs. SSD 0  $\mu\text{M}$ . (C) MDA-MB-231 cells were exposed to 8  $\mu\text{M}$  SSD, 20 nM Baf or 0.25  $\mu\text{M}$  Rapa for 24 h and then stained with LysoTracker™ Red. Fluorescence was observed using a confocal laser microscope (scale bar=50  $\mu\text{m}$ ) (D) MDA-MB-231 cells were co-transfected with mRFP-LC3 and LAMP1-mGFP plasmids and then exposed to 8  $\mu\text{M}$  SSD, 20 nM Baf or 0.25  $\mu\text{M}$  Rapa for 24 h. The fluorescence of mRFP-LC3 and LAMP1-mGFP puncta was observed by confocal laser microscopy (scale bar=10  $\mu\text{m}$ ). SSD, saikosaponin D; Baf, bafilomycin A1; Rapa, rapamycin; LAMP, lysosome-associated membrane glycoprotein; GFP, green fluorescent protein; RFP, red fluorescent protein.

the blockade of autophagic degradation was not involved in SSD-mediated apoptosis.

SSD, a kind of triterpenoid saponins extracted from *Radix bupleuri*, has been demonstrated to have an inhibitory effect on tumor cells. For example, SSD suppressed liver cancer proliferation by inhibiting cyclooxygenase 2 expression (27), as well as inhibiting the proliferation, invasion and migration of osteosarcoma cells (28). Previous studies also reported that SSD inhibited the proliferative activity of breast cancer cells, such as HCC1937 and MCF-7 (29,30). The results of the present study revealed that SSD induced apoptosis and inhibited the proliferative activity of MDA-MB-231 breast cancer cells.

p38 MAPK, one of the MAPK family members, serves an important role in differentiation, growth inhibition, cell cycle arrest and apoptosis (31). Thus, p38 demonstrates potential to be

developed into a cancer inhibitor, as the phosphorylation of p38 is important for the induction of apoptosis (32,33). The present results confirmed that SSD induced apoptosis through the phosphorylation/activation of the p38 MAPK signaling pathway. First, SSD induced the phosphorylation/activation of p38 MAPK. Second, the p38 MAPK inhibitor SB203580 reversed the SSD-mediated increase of apoptosis, the inhibitory effect over cell viability, and the activation of caspase-3 and PARP.

Numerous studies have reported that autophagy can be blocked at different stages of the autophagic flux (34-37). For example, 3-MA inhibited the formation of autophagosomes by inhibiting phosphatidylinositol 3-kinase catalytic subunit type 3 (38). In addition, Baf was found to disrupt the autophagic flux by changing the lysosomal acidity through inhibiting H<sup>+</sup> V-ATPase (39). It is also important to maintain

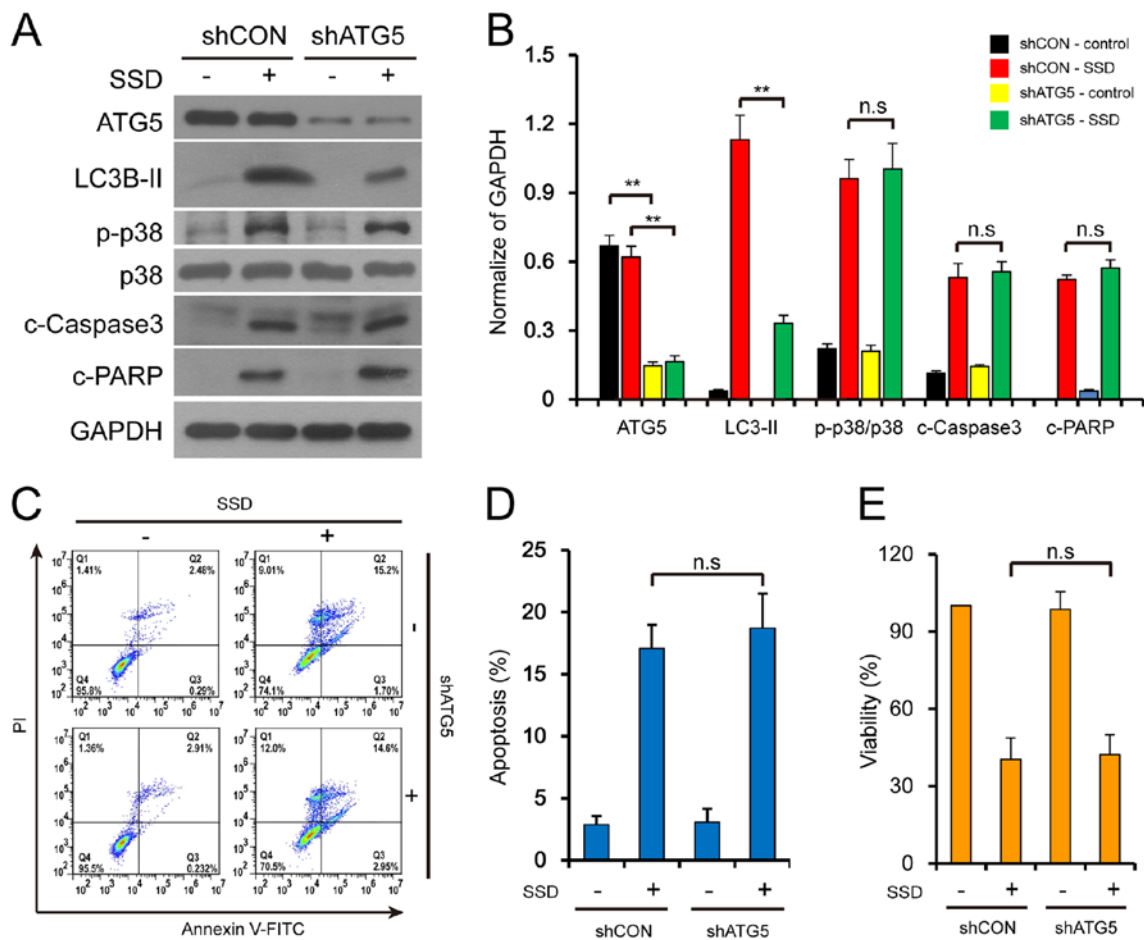


Figure 5. Excessive accumulation of autophagosomes does not contribute to SSD-mediated apoptosis. Stable knockdown MDA-MB-231 cells (shCON or shATG5) were exposed to 10  $\mu$ M SSD for 24 h. (A) Following treatment, the expression levels of ATG5, LC3B-II, p-p38, p38, c-caspase-3 and c-PARP were detected using western blotting. (B) Semi-quantitative analysis of the expression levels in part A, n=3; \*\*P<0.01. (C) Apoptotic cells were determined using flow cytometry. (D) Quantitative analysis of the apoptotic rate from part C. n=3. (E) Cell viability was determined using MTT assays. n=3. All data are presented as the mean  $\pm$  SD. sh, small hairpin RNA; CON, control; ATG, autophagy-related protein; c-, cleaved; n.s., not significant; p-, phosphorylated; LC3B, microtubule-associated protein 1 light chain 3  $\beta$ ; PARP, poly (ADP-ribose) polymerase; SSD, saikosaponin D; PI, propidium iodide.

the normal function of lysosomes during autophagy, since lysosome dysfunction resulted in the blockade of autophagic degradation (40). LAMP1 and LAMP2, two major lysosomal membrane proteins, are important for maintaining lysosomal function and assisting autophagosome-lysosome fusion, and the disruption of both LAMP1 and LAMP2 was found to result in the increased accumulation of autophagosomes (22). Lysosomal cathepsins (cathepsin B and cathepsin D) are also essential for maintaining normal lysosome function, and the inhibition of lysosomal cathepsins has been observed to result in the blockage of autophagic degradation (41). In addition, the blockade of autophagosome-lysosome fusion inhibited the formation of autolysosomes, which subsequently resulted in the inhibition of the autophagic flux (42).

The results of the present study confirmed that SSD was unable to impair the lysosomal function; however, it inhibited the fusion of autophagosomes and lysosomes, resulting in the inhibition of autolysosome formation. First, SSD treatment did not affect the acidic environment of the lysosomes; second, SSD treatment did not reduce the expression levels of LAMP1 or LAMP2; third, SSD treatment did not affect the expression levels of the lysosomal cathepsins; and finally, SSD treatment disrupted the co-localization of LC3 and LAMP1.

Cell death may be caused by either the induction or inhibition of autophagy (43,44). However, previous research has reported that classical autophagy degradation inhibitors, such as ammonium chloride chloroquine and Baf can cause autophagy-independent cell death (9). Consistent with these findings, results of the current study also showed that the accumulation of autophagosomes was not involved in SSD-mediated apoptosis. The genetic knockdown of ATG5 expression levels significantly decreased the SSD-mediated accumulation of LC3B-II; however, it did not affect the SSD-mediated apoptotic rate, or the activation of caspase-3, PARP or p38 MAPK. These results indicated that the blockade of autophagic degradation and apoptosis, both of which are mediated by SSD, may be two separate events. Thus, these findings provided additional evidence to support the association between the inhibition of autophagy degradation and cell death.

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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

RF and JC conceived and designed the study; RF, LZ and YL performed the experiments; RF, LZ and HX designed the experiments; YM, ZL and BL analyzed the data; BL and JC reviewed and edited the manuscript; RF and HX wrote the original manuscript; and JC and BL acquired the funding. All authors read and approved the final manuscript.

## Ethics approval and consent to participate

Not applicable.

## Patient consent for publication

Not applicable.

## Competing interests

The authors declare that they have no competing interests.

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