Cold PSM, but not TRAIL, triggers autophagic cell death: A therapeutic advantage of PSM over TRAIL

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Received November 2, 2017; Accepted March 26, 2018

DOI: 10.3892/ijo.2018.4413

Abstract. Tumor necrosis factor-related apoptosis-inducing ligand (TRAIL) and cold plasma-stimulated medium (PSM) are promising novel anticancer tools due to their strong anticancer activities and high tumor-selectivity. The present study demonstrated that PSM and TRAIL may trigger autophagy in human malignant melanoma and osteosarcoma cells. Live-cell imaging revealed that even under nutritional and stress-free conditions, these cells possessed a substantial level of autophagosomes, which were localized in the cytoplasm separately from tubular mitochondria. In response to cytotoxic levels of PSM, the mitochondria became highly fragmented, and aggregated and colocalized with the autophagosomes. The cytotoxic effects of PSM were suppressed in response to various pharmacological autophagy inhibitors, including 3-methyladenine (3-MA) and bafilomycin A1, thus indicating the induction of autophagic cell death (ACD). Lethal levels of PSM also resulted in non-apoptotic, non-autophagic cell death in a reactive oxygen species-dependent manner under certain circumstances. Furthermore, TRAIL exhibited only a modest cytotoxicity toward these tumor cells, and did not induce ACD and mitochondrial aberration. The combined use of TRAIL and subtoxic concentrations of 3-MA resulted in decreased basal autophagy, increased mitochondrial aberration, colocalization with autophagosomes and apoptosis. These results indicated that PSM may induce ACD, whereas TRAIL may trigger cytoprotective autophagy that compromises apoptosis. To the best of our knowledge, the present study is the first to demonstrate that PSM can induce ACD in human cancer cells. These findings provide a rationale for the advantage of PSM over TRAIL in the destruction of apoptosis-resistant melanoma and osteosarcoma cells.

Introduction

Tumor necrosis factor (TNF)-related apoptosis-inducing ligand (TRAIL) is a member of the TNF superfamily, which preferentially kills malignant cells over nontransformed cells (1-4). TRAIL can induce extrinsic and intrinsic death pathways by binding its specific receptors with death domain TRAIL receptor (TRAIL-R1)/death receptor (DR)4 and TRAIL-R2/DR5 (5,6). However, some cancer cell types are inherently resistant to TRAIL, despite expressing death-inducing receptors (7-11). Furthermore, some cancer cell types acquire considerable tolerance to TRAIL during prolonged treatment. Accordingly, current clinical trials have been disappointing, and the combined use of agents that overcome drug resistance is necessary for efficient TRAIL therapy. Non-thermal (cold) atmospheric plasma (CAP) has emerged as another promising cancer treatment, since like TRAIL, it kills various cancer cells while sparing nontransformed cells under optimal conditions (12-15). Cold plasma-stimulated medium (PSM) also exhibits vigorous and tumor-selective anticancer activities (16-19) and has emerged as an alternative method of direct CAP irradiation; PSM is better than direct CAP irradiation for systematic or local administration to deep tissues.

Cancer cells, including malignant melanoma (MM) and osteosarcoma (OS) cells, are characterized by their intrinsic resistance to apoptosis; in addition, they frequently become more tolerant to numerous apoptosis-inducing antitumor drugs. Nevertheless, the majority of conventional drugs primarily kill cells by apoptosis. Accordingly, current chemotherapy toward these cancers is severely compromised by intrinsic and acquired resistance; therefore, induction of another mode
of cell death may be a useful approach for the treatment of apoptosis-resistant cells (20,21). Autophagy is a primary catabolic process that degrades cellular components and damaged organelles via lysosomes; this process copes with cellular stressors, such as starvation, and supplies energy and metabolic precursors. Autophagy consists of numerous processes, including induction of cytoplasmic double-layered membranes, which are known as phagophores, phagophore elongation and autophagosome formation, a fusion of autophagosomes with lysosomes, and degradation and recycling. All processes, from the formation of autophagosomes to the degradation of cellular components, are strictly regulated by autophagy-related (Atg) proteins that are encoded by Atg genes (22). Autophagy is classified into three different types: Macroautophagy (subsequently referred to as autophagy), microautophagy and chaperone-mediated autophagy. Autophagy is negatively regulated by mammalian target of rapamycin complex I in response to insulin and amino acid signals, and is driven transiently via removal of its suppression through the depletion of these nutrients (23-25). Therefore, autophagy is of particular importance for the survival of constitutively proliferating cells, such as cancer cells, that are regularly imposed to energy demands (20,26). In addition, autophagy contributes to cancer cell survival by removing damaged organelles, including mitochondria and endoplasmic reticulum (ER) by microautophagy, which is also known as mitophagy and ERphagy, respectively. These damaged organelles are degraded via lysosomal enzymes following engulfment into autophagosomes; such quality control is crucial for cell survival. Conversely, autophagy is also characterized by a unique cell death pathway that acts as a tumor suppressor when it leads to autophagic cell death (ACD) (27-29).

Our previous study revealed that PSM prepared by CAP irradiation of Dulbecco’s modified Eagle’s medium (DMEM) could kill an array of MM, OS and lung cancer cells, while sparing nontransformed melanocytes and fibroblasts (30). In addition, PSM led to increased caspase-3/7 activity, and modest cleavage of caspase-9, caspase-3/7 and poly ADP-ribose polymerase; furthermore, caspase-3/7-specific inhibitors failed to suppress cell death. Therefore, the present study aimed to examine the possibility that PSM may induce another cell death modality. The results demonstrated that PSM can trigger ACD in MM and OS cells.

Materials and methods

Materials. Soluble recombinant human TRAIL was obtained from Enzo Life Sciences, Inc. (Farmingdale, NY, USA). 3-Methyladenine (3-MA), chloroquine (CQ) and bafilomycin A1 (Baf) were obtained from Sigma-Aldrich (Merck KGaA, Darmstadt, Germany). The pan-caspase-inhibitor z-VAD-fluoromethylketone (Z-VAD-FMK) was purchased from Merck Ltd. (Tokyo, Japan). All insoluble reagents were dissolved in dimethyl sulfoxide (DMSO) and diluted with high glucose-containing DMEM supplemented with 10% fetal bovine serum (FBS) (both from Sigma-Aldrich; Merck KGaA) or Hank’s balanced salt solution (HBSS; pH 7.4; Nissui Pharmaceutical Co., Ltd., Tokyo, Japan) (final DMSO concentration, <0.1%) prior to use. The manganese-porphyrin superoxide dismutase mimetic MnTBAp (Enzo Life Sciences, Inc.) was dissolved in 1 mM NaOH (pH 8.0) and HBSS was added to lower the pH to 7.4.

Cell culture. Human A375 MM cells (American Type Culture Collection (ATCC)® cell number CRL-1619) were obtained from ATCC (Manassas, VA, USA). A2058 MM cells (cell number IFO 50276) and human A549 lung adenocarcinoma cells (cell number JCRB0076) were purchased from the JCRB Cell Bank of National Institutes of Biomedical Innovation, Health, and Nutrition (Osaka, Japan). Human HOS (TE85) OS cells (cell number RCB0992), SAOS-2 OS cells (cell number RCB0428) and 143B cells (cell number RCB0701) were obtained from the Riken BioResource Center (Tsukuba, Japan). Human dermal fibroblast (HDF) cells from facial dermis were obtained from Cell Applications (San Diego, CA, USA). All cells were cultured in 10% FBS/DMEM supplemented with 100 U/ml penicillin and 100 µg streptomycin (Thermo Fisher Scientific, Inc., Waltham, MA, USA) in a humidified atmosphere containing 95% air/5% CO2 at 37°C. The cells were harvested by incubating with 0.25% trypsin-EDTA (Thermo Fisher Scientific, Inc.) for 5 min at 37°C. Throughout the study, various cell lines were used in each experiment to determine whether the reactions observed were specific for the cell line used or were general among the various cell lines. A2058 and HOS (or SAOS-2) cells were used as the main representatives of MM and OS cells, respectively. In some experiments, autophagy was compared between normally grown A375 (confluence, 70-80%) cells and overgrown (confluent, partially floating) cells.

PSM preparation. CAP was generated using an originally-developed low-frequency plasma jet device equipped with an asymmetrical dielectric barrier discharge, as previously described (30). The typical frequency was 20 kHz, with a peak voltage of 8 kV, a current of 20 mA and a helium flow rate of 300 ml/min. PSM was prepared by irradiating 1 ml FBS/DMEM with CAP for 5 min once. The original PSM (100%) was diluted to a final concentration of 25% with FBS/DMEM (for cell experiments) or HBSS (for biochemical experiments), and indicated as PSM (25%).

Cell viability assay. Cell viability was measured according to the WST-8 assay using Cell Counting Reagent SF (Nacalai Tesque, Inc., Kyoto, Japan) as previously described (31). This is a colorimetric assay that detects viability based on the formation of a water-soluble formazan product. Briefly, the cells were seeded at a density of 8x103 cells/well in 96-well plates (Corning Incorporated, Corning, NY, USA) and cultured with PSM (12.5, 25, 50, and 100%), 100 ng/ml TRAIL, 3-MA (1.3-5 mM), CQ (100 and 300 µM), and Baf (100 and 300 nM) alone or in combination for 72 h at 37°C, prior to the addition of 10 µl cell counting reagent SF for 1 h. For some experiments, the cells were cultured with the aforementioned reagents in the presence of 10 µM Z-VAD-FMK or 30 µM MnTBAp. Absorbance was measured at 450 nm using an ARVO MX microplate reader (PerkinElmer Japan Co., Ltd., Yokohama, Japan).

Live-cell imaging. The mitochondrial network was analyzed as previously described (32) with minor modifications. Briefly,
cells in FBS/DMEM (3x10^4/well) adherent on an 8-well chambered coverslip with a glass bottom (Imaging Chamber 8 CG; Zell-Kontakt GmbH, Nörten-Hardenberg, Germany) were treated with PSM (25 and 100%), 100 ng/ml TRAIL, 0.5 or 1 µM rapamycin and 5 mM 3-MA alone or in combination for 24 h in a humidified atmosphere containing 95% air/5% CO₂ at 37°C. Following media removal by aspiration, the cells were washed with fresh FBS/DMEM and stained with 20 nM MitoTracker Red CMXRos for 1 h at 37°C in the dark. The nuclei were counterstained with Hoechst 33342. The cells were then washed with and immersed in FluoroBrite™ DMEM (Thermo Fisher Scientific, Inc.). Images were obtained using a BX-Z-700 Fluorescence Microscope (Keyence Corporation, Osaka, Japan) equipped with a 100X, 1.40 n.a. UPLSAPO super-apochromat, coverslip-corrected oil objective (Olympus Corporation, Tokyo, Japan). Images were analyzed using BZ-H3A application software (Keyence Corporation) and National Institutes of Health (NIH) ImageJ software (bundled with 64-bit Java 1.8.0_112; NIH, Bethesda, MD, USA). The formation of autophagosomes was determined using the CYTO-ID® Autophagy Detection Kit (Enzo Life Sciences, Inc.) according to the manufacturer's protocol. Briefly, cells were stained with CYTO-ID for 1 h at 37°C in the dark and treated with the aforementioned agents. For monitoring colocalization of mitochondria and autophagosomes, cells were coincidently stained with MitoTracker Red CMXRos and the CYTO-ID. Images were obtained using EVOS FL Cell Imaging system (Thermo Fisher Scientific, Inc.) equipped with a 40X, 0.60 n.a. LUCPLFLN objective (Olympus Corporation) as previously described (33).

**Western blotting.** Following treatment with PSM (25 and 100%) for 12, 24, 36, 48 and 72 h, cells were washed with Ca²⁺-, Mg²⁺-free PBS, and were lysed during 30 min agitation with CellLytic™ Tisox buffer (Sigma-Aldrich; Merck KGaA) containing a protease inhibitor cocktail and a phosphatase inhibitor cocktail (both from Sigma-Aldrich; Merck KGaA). Cell debris was removed by centrifugation at 20,000 x g for 15 min at 4°C. The supernatants were collected and protein concentrations were analyzed using a bicinchoninic acid protein assay (Thermo Fisher Scientific, Inc.), according to the manufacturer's protocol. Equal amounts of protein (10 µg) were separated by 4-12% NuPage Bis-Tris acrylamide gels (Invitrogen; Thermo Fisher Scientific, Inc.) and were transferred to polyvinylidene difluoride membranes (Immobilon-P; EMD Millipore, Bedford, MA, USA). Blots were blocked for 30 min in Tris-buffered saline with 0.05% Tween-20 (TBST; Sigma-Aldrich; Merck KGaA) containing 2% nonfat dry milk. The blots were then incubated with primary antibodies against p62 (PM045, 1:1,000; Medical & Biological Laboratories Co., Ltd., Nagoya, Japan), microtubule-associated protein 1A/1B-light chain 3 (LC3; #12741, 1:1,000) and GAPDH (#5174, 1:1,000; Cell Signaling Technology, Inc., Danvers, MA, USA) overnight at 4°C in TBST containing 2% nonfat dry milk. After washing two times with TBST, membranes were incubated with horseradish peroxidase-conjugated goat anti-rabbit (#7074, 1:2,000; Cell Signaling Technology, Inc.) for 1 h at room temperature. Subsequently, blots were washed three times with TBST and immersed in enhanced chemiluminescence reagent (GE Healthcare, Chicago, IL, USA) to enhance the signals. The signals were then captured using an LAS-4000 Camera system (Fujifilm Corporation, Tokyo, Japan).

**Statistical analysis.** Data are presented as the means ± standard deviation. Data were analyzed by one-way analysis of variance followed by Tukey's post hoc test using add-in software for Excel 2016 for Windows (Microsoft Corporation, Redmond, WA, USA). P<0.05 was considered to indicate a statistically significant difference.

**Results**

**Human MM and OS cells exhibit substantial basal autophagy.** To gain insight into the possible role of autophagy in the antitumor activity of PSM, the present study evaluated its ability to modulate autophagic flux in MM and OS cells. CYTO-ID® Green is a cationic amphiphilic dye that can precisely monitor autophagic vacuoles without transfection. Live-cell imaging revealed a considerable number of CYTO-ID puncta in A375 and HOS cells in DMEM supplemented with 10% FBS. The CYTO-ID puncta markedly increased in response to 1 µM rapamycin, whereas these puncta were decreased almost entirely following treatment with 5 mM 3-MA (Fig. 1), thus validating that staining represents autophagosomes. Consistent with our previous study (30), treatment with PSM for 24 h resulted in robust cell damage. Accordingly, adherent spindle HOS and A549 cells became round, and some cells lost their adherence and integrity (Fig. 2A and B). Concomitantly, CYTO-ID puncta became clustered in these cells. Conversely, HDF cells possessed only modest basal CYTO-ID puncta and minimal changes in CYTO-ID puncta were observed following PSM treatment (Fig. 2C).

**PSM induces colocalization of aggregated mitochondria and autophagosomes.** Damaged mitochondria are removed by autophagic processes known as mitophagy. Accordingly, mitophagy controls the quality of mitochondria, thereby facilitating cell survival, whereas the excessive removal of mitochondria facilitates cell death (34, 35). Our previous study reported that PSM could induce excessive mitochondrial fragmentation and aggregation, alongside intensive mitochondrial damage (30). Therefore, the present study simultaneously monitored CYTO-ID puncta and mitochondrial morphology in live cells. Unstimulated A375 cells possessed tubular mitochondria and the CYTO-ID puncta that primarily diffused in the cytoplasm. In this case, the puncta and the mitochondria were separately located, as indicated by the minimal overlapping of green and red signals (Fig. 3A, upper panels). Following PSM treatment, the mitochondria became fragmented, punctate and aggregated, as previously described (30). Furthermore, some of the damaged round cells possessed clustered CYTO-ID puncta, the majority of which was colocalized with the aggregated mitochondria (Fig. 3A, lower panels). Even in the absence of insult, overgrown A375 cells became round and damaged (Fig. 3B, upper panels). Similarly, in PSM-treated cell images, damaged cells possessed numerous clustered CYTO-ID puncta, which were colocalized with aggregated mitochondria. To determine whether these effects were specific for PSM, the effects of TRAIL were determined.
The results demonstrated that TRAIL increased CYTO-ID puncta in TRAIL-resistant A375 cells; however, TRAIL led to modest mitochondrial fragmentation, and caused minimal mitochondrial aggregation and colocalization with autophagosomes (Fig. 3C, lower panels). These results indicated that PSM may induce colocalization of aggregated mitochondria and autophagosomes.

Toxic PSM specifically increases autophagic flux. The present results indicated that PSM specifically leads to colocalization of aggregated mitochondria and autophagosomes. To determine whether this intrinsic event is cytoprotective or cytotoxic, the study examined the association of PSM with cell damage. PSM (100%) displayed significant cytotoxic activity toward MM (A2058) and OS (HOS) cells, whereas it was ineffective at lower concentrations (≤50%) (Fig. 4A and B). Subsequently, the present study compared the effects of toxic and subtoxic PSM on autophagic flux. The cytosolic protein LC3-I forms a complex with phosphatidylethanolamine, thus resulting in the formation of LC3-II, which directly associates with the autophagosome membrane. LC3-II enables the binding of adaptor proteins, such as p62/sequestome 1 (36). Therefore, the present study analyzed the expression levels of p62 and LC3-II following PSM treatment. Subtoxic PSM had minimal effects on the expression levels of p62 and LC3-II up to 72 h post-stimulation (Fig. 4C, left panels). However, toxic PSM increased the expression levels of LC3-II, but not p62, 36-48 h post-stimulation (Fig. 4C, right panels). These results suggested that toxic PSM may specifically increase autophagic flux.

**Colocalization of mitochondria and autophagosomes is associated with autophagy induction and cell death.** To further explore the relationship between the colocalization of mitochondria and autophagosomes and autophagy, the present study examined the effects of autophagy modulators on mitochondrial morphology and autophagosome location. Treatment with the autophagy promoter rapamycin markedly increased CYTO-ID puncta in HOS cells (Fig. 5A). Concomitantly, some, but not all, of the puncta were colocalized with the mitochondria, as shown by the appearance of yellow signals in the overlay images. Conversely, 3-MA completely suppressed CYTO-ID puncta and increased mitochondrial hyperfusion (Fig. 5A). In addition, subtoxic PSM did not induce robust colocalization of the mitochondria and autophagosomes (Fig. 5A). All of these agents caused minimal mitochondrial fragmentation and aggregation. Furthermore, toxic PSM strongly induced mitochondrial fragmentation and aggregation, and colocalization of mitochondria and autophagosomes (Fig. 5A). Microscopic analysis with a higher resolution confirmed these observations and indicated the colocalization of clustered CYTO-ID puncta and aggregated mitochondria (Fig. 5B). These results suggested that the colocalization of mitochondria and autophagosomes may be associated with autophagy induction and cell damage.
PSM induces ACD during the initial 24 h post-stimulation.

To further explore the role of autophagy in the antitumor activity of PSM, the present study examined the effects of numerous pharmacological autophagy inhibitors with various mechanisms of action on PSM cytotoxicity. 3-MA inhibits autophagosome formation by inhibiting the type III phosphatidylinositol-3-kinase, whereas the antimalarial drug CQ inhibits the fusion of autophagosomes and lysosomes. The antibiotic Baf inhibits vacuolar-type H^+\text{-}ATPases and increases lysosomal pH, thus compromising the fusion of autophagosomes and lysosomes. HOS cells were relatively susceptible to PSM cytotoxicity. Accordingly, their viability often decreased considerably (maximum reduction, 80%) at 24 h following treatment with PSM (≥25%) (Fig. 6A). Baf inhibited the cytotoxic effects of 25% PSM entirely, and those of 100% PSM partially. In addition, 3-MA significantly reduced the effects of PSM (25%), but not those of PSM (100%). PSM (100%), but not PSM (25%), also significantly decreased the viability of SAOS-2 and MG63 cells (Fig. 6B and C), and 3-MA inhibited the cytotoxic effects of PSM in MG63 cells, but not in SAOS-2 cells, whereas Baf blocked the effects in both cell types. These results demonstrated that PSM may induce ACD during the initial 24 h post-stimulation.

PSM induces another reactive oxygen species (ROS)-dependent cell death modality upon prolonged treatment. PSM exerted higher degrees of cytotoxicity toward MM and OS cells following prolonged treatment (72 h). As a result, PSM (≥25%) markedly decreased the viability of HOS, 143B and A2058 cells (Fig. 7A-C). Notably, in all cell lines tested, these effects were not blocked by any of the autophagy inhibitors (3-MA, Baf and CQ) (Fig. 7A-C). In addition, 3-MA (5 mM) and CQ (100 µM) alone significantly reduced the viability of 143B cells. Previously, it was demonstrated that ROS serve a vital role in mediating PSM cytotoxicity (30,37). As a result, the superoxide oxidase mimetic antioxidant MnTBaP has been revealed to suppress the effects of PSM in HOS cells, whereas the effects were more pronounced in cells treated with PSM (100%) compared with in those treated with PSM (25%) (37). In agreement with our previous observations (30), MnTBaP significantly reduced the cytotoxicity of PSM in the present study, with a higher potency toward the lower concentration of PSM (Fig. 7D). Conversely, the pan-caspase-inhibitor z-VAD-FMK exhibited minimal effects on PSM cytotoxicity regardless of the PSM concentration and cell type examined (data not shown). These results indicated that PSM may also induce another ROS-dependent, non-apoptotic, non-autophagic cell death upon prolonged treatment.

TRAIL predominantly triggers cytoprotective autophagy, which prevents apoptotic and non-apoptotic cell death. Since TRAIL increased CYTO-ID puncta in cancer cells, the present study explored the possible role of autophagic flux in the antitumor activity of TRAIL. A 72-h-incubation protocol was employed, because cells were highly TRAIL-resistant and TRAIL treatment for 24 h had minimal cytotoxicity. The autophagy inhibitors 3-MA (≥1.3 mM) and CQ (≥100 µM) alone dose-dependently decreased the viability of A2058 and SAOS-2 cells. Furthermore, the low concentration of 3-MA (1.3 mM) significantly potentiated the cytotoxicity of TRAIL in the two cell lines (Fig. 8A and B). Conversely, ≤300 nM Baf neither decreased cell viability nor enhanced the effect of TRAIL, regardless of the cell type examined. The concentration of 3-MA suitable for TRAIL sensitization varied depending on the cell lines tested (i.e., HOS and SAOS-2 cells), z-VAD-FMK entirely suppressed the effect. Z-VAD-FMK significantly inhibited the effects of 3-MA (2.5 mM) on HOS.
In addition, z-VAD-FMK completely blocked the effects of 3-MA (1.3 mM) and minimally inhibited the effects of 3-MA (2.5 mM) on SAOS-2 cells (Fig. 8C and D). Furthermore, the present study analyzed the effects of cotreatment with TRAIL and 3-MA on autophagy and mitochondrial morphology. TRAIL alone increased CYTO-ID puncta, whereas 3-MA alone abolished them in HOS cells (Fig. 8E). Concomitantly, mitochondria became hyperfused in response to TRAIL or 3-MA, as shown by increased highly intra-connected mitochondria (Fig. 8E). The combined application of TRAIL and 3-MA led to the formation of large clusters of CYTO-ID puncta that colocalized with the mitochondria (Fig. 8E). Collectively, these results indicated that TRAIL predominantly triggers cytoprotective autophagy, the prevention of which may lead to colocalization of the mitochondria and autophagosomes, apoptosis and another form of non-apoptotic cell death.

**Discussion**

The present study aimed to determine whether PSM modulated autophagy in human MM and OS cells. Western blotting demonstrated that toxic, but not subtoxic, PSM increased LC3-II expression, which is required for autophagosome formation. Furthermore, the cytotoxic effects of PSM were
strongly inhibited following treatment with the pharmacological autophagy inhibitors 3-MA and Baf, thus indicating that autophagy is responsible for cell death. These findings suggested that PSM may trigger ACD in tumor cells. Notably, mitochondrial abnormalities accompanied the induction of ACD. In agreement with our previous study (30), PSM induced mitochondrial aberrations in a concentration-dependent manner. Toxic PSM led to excessive mitochondrial fragmentation and aggregation, whereas subtoxic PSM induced only modest mitochondrial fragmentation. In addition, live-cell imaging of CYTO-ID puncta revealed marked alterations in the status and location of autophagosomes in response to PSM. When unstimulated, autophagosomes existed as small diffuse particles, the majority of which were localized in the cytoplasm separate from the mitochondria. However, following PSM treatment, autophagosomes formed a cluster that colocalized with the aggregated mitochondria. In mitophagy, damaged mitochondria are engulfed by autophagosomes, so that mitochondria and autophagosomes eventually colocalize. Therefore, the specific effects of toxic PSM appear similar to mitophagy. Our previous study demonstrated that mitochondria were heavily damaged, resulting in the loss of mitochondrial membrane potential ($\Delta \Psi_m$) and integrity, when they became punctate and aggregated (32,33). Notably, dissipation of the $\Delta \Psi_m$ is a primary trigger of mitophagy (34). Diverse insults, such as carbonyl cyanide m-chlorophenylhydrazone, carbonyl cyanide p-trifluoromethoxyphenylhydrazone and salinomycin, commonly induce loss of the $\Delta \Psi_m$, thereby activating mitophagy (38,39). Loss of the $\Delta \Psi_m$ leads to the accumulation of phosphatase and tensin homolog-induced putative kinase 1 (PINK1) on the outer mitochondrial membrane. The localized PINK1 phosphorylates E3-ubiquitin ligase Parkin and activates Parkin-mediated ubiquitination, thus resulting in autophagic degradation of the damaged organelles (34). Therefore, it is possible that PSM injures the mitochondria, thereby leading to mitophagy in the present cell system. p62 has been reported to serve a vital role in mitophagy (40), whereas it is superfluous in some circumstances (41). The present study demonstrated that only marginal p62 accumulation occurred following PSM treatment; therefore, the role of p62 in autophagic flux appears to be minor in the present cell system. Several lines of evidence indicated that the mitophagy-like event observed promoted cell death and contributed to the antitumor activity of PSM: i) The event spontaneously occurred in damaged or dying cancer cells; ii) TRAIL, which had only a modest cytotoxic effect, did not induce the event; iii) toxic, but not subtoxic PSM, caused the event; (iv) TRAIL and 3-MA synergistically increased the event and cell killing. It is widely accepted that controlled mitophagy is cytoprotective, since it removes damaged mitochondria; however, excessive mitophagy leads to cell damage by compromising the energy supply, and Ca$^{2+}$ and metabolic homeostasis. Therefore, it is possible that PSM may induce excessive mitophagy, which is responsible for the antitumor activity of PSM. However, further studies are required to verify this hypothesis.

The present data indicated that a basal level of autophagy operates in MM and OS cells; these cells possessed substantial levels of autophagosomes even under nutritional and stress-free conditions. Compared with the mitophagy-like event, basal autophagy appeared to serve a cytoprotective role, since

Figure 4. Toxic PSM increases autophagic flux. (A) A2058 and (B) HOS cells were treated with various concentrations of PSM for 72 h. Subsequently, cell viability was analyzed using the WST-8 assay in triplicate. Data are presented as the means ± standard deviation of a representative experiment (n=3). (C) 143B cells were treated with PSM (25 and 100%) for 12, 24, 36, 48 and 72 h at 37°C. Western blotting was used to detect p62, LC3 and GAPDH expression. LC3, microtubule-associated protein 1A/1B-light chain 3; PSM, cold plasma-stimulated medium.
autophagy inhibitors, including 3-MA and CQ alone exerted significant cytotoxicity toward these cancer cells in the absence of insult. Notably, the effects were more pronounced following 72 h treatment compared with after 24 h. These observations indicated that basal autophagy may be essential for cell survival, particularly in response to hypo-nutritional conditions. The present study suggested that in these cancer cells, TRAIL primarily induced cytoprotective autophagy, since TRAIL increased autophagosome formation, whereas its cytotoxicity was amplified, rather than inhibited, following treatment with...
Figure 6. PSM induces ACD during the initial 24 h post-stimulation. (A) HOS, (B) SAOS-2 and (C) MG63 cells were treated with PSM (25 and 100%), 5 mM 3-MA and 100 nM Baf alone or in combination for 24 h, and cell viability was analyzed using WST-8 assay in triplicate. Data are presented as the means ± standard deviation of a representative experiment (n=3). Data were analyzed by one-way analysis of variance followed by Tukey’s post hoc test. #P<0.001 vs. control; NS, not significant vs. control; *P<0.05; ***P<0.001. 3-MA, 3-methyladenine; Baf, bafilomycin A; PSM, cold plasma-stimulated medium.

Figure 7. PSM induces another ROS-dependent cell death modality upon prolonged treatment. (A) HOS, (B and D) 143B and (C) A2058 cells were treated with (A-C) PSM (25 and 100%) and 5 mM 3-MA, 100 µM CQ and 100 nM Baf alone or in combination for 72 h. (D) 143B cells were treated with PSM (25 and 100%) and 10 µM ZVAD and 30 µM Mn alone or in combination for 72 h. Cell viability was analyzed using the WST-8 assay in triplicate. Data are presented as the means ± standard deviation of a representative experiment (n=3). Data were analyzed by analysis of variance followed by Tukey’s post hoc test. """P<0.001 vs. control; NS, not significant vs. control; """"P<0.001. NS (All), no statistical significance between PSM in combination with 3-MA, CQ and Baf compared with PSM (25 and 100%) alone. 3-MA, 3-methyladenine; Baf, bafilomycin A; CQ, chloroquine; Mn, MnTBaP; PSM, cold plasma-stimulated medium; ZVAD, z-VAD-fluoromethylketone.
Figure 8. TRAIL predominantly triggers cytoprotective autophagy. (A) A2058 and (B) SAOS-2 cells were treated with 100 ng/ml TRAIL, 3-MA, CQ and Baf at the indicated concentrations (3-MA, mM; CQ, µM; Baf, nM) alone or in combination for 72 h. (C) HOS and (D) SAOS-2 cells were treated with 100 ng/ml TRAIL and 3-MA at the indicated concentrations (mM) for 72 h, in the absence or presence of 10 µM ZVAD. Cell viability was analyzed using the WST-8 assay in triplicate. Data are presented as the means ± standard deviation of a representative experiment (n=3). Data were analyzed by analysis of variance followed by Tukey's post hoc test. ###P<0.001; #P<0.05 vs. control; NS, not significant. ***P<0.001; **P<0.01; *P<0.05. (E) HOS cells were treated with 100 ng/ml TRAIL and 5 mM 3-MA alone or in combination, and were stained with CYTO-ID and MitoTracker Red CMXRos. Panels labeled ‘Zoom’ represent the expansion of the small white boxes within ‘Overlay’ images. Scale bar, 100 µm. 3-MA, 3-methyladenine; Baf, bafilomycin A; CQ, chloroquine; Mn, MnTBaP; PSM, cold plasma-stimulated medium; ZVAD, z-VAD-fluoromethylketone.
autophagy inhibitors, including 3-MA. Notably, 3-MA/CQ and Baf had various effects on cell viability and TRAIL cytotoxicity. At present, the reason for the results are unclear, because they commonly serve as autophagy inhibitors. However, it was previously demonstrated that 3-MA, but not Baf, interferes with mitochondrial Ca$^{2+}$ loading, which is an essential process for energy production and cell survival. These observations are intriguing, since our recent study demonstrated that disturbance of mitochondrial Ca$^{2+}$ may lead to cancer cell death and sensitization to TRAIL cytotoxicity (31). Therefore, it is possible that the difference resides in autophagy-independent effects. Further investigations are required.

The results of the present study are consistent with those previously reported in various cancer cell types, including lung, bladder, prostate, colorectal cancer and hepatoma cells (42-44). Notably, despite its ability to trigger such cytoprotective autophagy, TRAIL minimally led to the putative cytotoxicity autophagy-like event. Therefore, TRAIL may be similar to various antitumor drugs, such as temozolomide, epirubicin and sorafenib, which have been reported to engage cytoprotective autophagy (45-48). Furthermore, the blockade of basal autophagy by 3-MA enabled TRAIL to increase the mitophagy-like event and kill TRAIL-resistant MM and OS cells, whereas it promoted cell death, but not ACD. In addition, the mitochondrial morphological alterations observed in response to TRAIL + 3-MA appeared to be somewhat different from those observed following toxic PSM treatment. Collectively, these results suggested that suppression of basal cytoprotective autophagy may promote the cytocidal mitophagy-like event, whereas another fundamental cellular event is required for induction of the full mitochondrial abnormalities leading to ACD induction. Further studies are required to identify such events.

Recent evidence has indicated that autophagy and apoptosis are closely associated, and that they may regulate one another (49-51). This connection has a significant impact on tumor cell sensitivity to antitumor chemotherapeutic drugs. In particular, negative regulation of apoptosis by autophagy is significant, since various antitumor drugs have been revealed to induce cytoprotective autophagy, which allows cancer cells to cope with apoptotic insults. Accordingly, the induction of autophagy in cancer cells represents a double-edged sword, leading to cell survival or ACD depending on the cell type and conditions. Therefore, it is critical to understand the pathways controlling context-dependent autophagy for the application of ACD in cancer treatment.

In conclusion, the present study demonstrated that PSM may trigger cytocidal autophagy, whereas TRAIL triggered cytoprotective autophagy. To the best of our knowledge, the present study is the first to demonstrate that PSM may induce ACD in human cancer cells. These findings provide a rationale for the development of PSM as a novel approach for the treatment of apoptosis-resistant tumors, including MM and OS. In addition, PSM may serve as a useful model for studying the mechanisms underlying ACD induction in cancer cells.

Acknowledgements

The authors would like to thank the JCRB Cell Bank of National Institutes of Biomedical Innovation, Health and Nutrition (Osaka, Japan) and Riken BioResource Center (Tsukuba, Japan) for providing cell lines.

Funding

This work was supported in part by JSPS KAKENHI Grant Number 15K09750 to YSK and 15K09792 to TO.

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

TI performed experiments, analyzed data and wrote the manuscript. TA performed experiments and wrote the manuscript. MSK and TT performed experiments and analyzed data. YY, TO and YT conceived and designed the study, performed the critical revision of the manuscript, and provided analytical tools. YSK conceived and designed the study, performed experiments and wrote the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Conflict of interest

The authors declare that they have no competing interests.

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