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Lysosomal Ca²⁺ as a mediator of palmitate-induced lipotoxicitySoo-Jin Oh^{1,2,5}, Yeseong Hwang^{1,3,5}, Kyu Yeon Hur⁴ and Myung-Shik Lee^{1,2,3}✉

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While the mechanism of lipotoxicity by palmitic acid (PA), an effector of metabolic stress in vitro and in vivo, has been extensively investigated, molecular details of lipotoxicity are still not fully characterized. Since recent studies reported that PA can exert lysosomal stress in addition to well-known ER and mitochondrial stress, we studied the role of lysosomal events in lipotoxicity by PA, focusing on lysosomal Ca²⁺. We found that PA induced accumulation of mitochondrial ROS and that mitochondrial ROS induced release of lysosomal Ca²⁺ due to lysosomal Ca²⁺ exit channel activation. Lysosomal Ca²⁺ release led to increased cytosolic Ca²⁺ which induced mitochondrial permeability transition (mPT). Chelation of cytoplasmic Ca²⁺ or blockade of mPT with olesoxime or decylubiquinone (DUB) suppressed lipotoxicity. Lysosomal Ca²⁺ release led to reduced lysosomal Ca²⁺ content which was replenished by ER Ca²⁺, the largest intracellular Ca²⁺ reservoir (ER → lysosome Ca²⁺ refilling), which in turn activated store-operated Ca²⁺ entry (SOCE). Inhibition of ER → lysosome Ca²⁺ refilling by blockade of ER Ca²⁺ exit channel using dantrolene or inhibition of SOCE using BTP2 inhibited lipotoxicity in vitro. Dantrolene or DUB also inhibited lipotoxic death of hepatocytes in vivo induced by administration of ethyl palmitate together with LPS. These results suggest a novel pathway of lipotoxicity characterized by mPT due to lysosomal Ca²⁺ release which was supplemented by ER → lysosome Ca²⁺ refilling and subsequent SOCE, and also suggest the potential role of modulation of ER → lysosome Ca²⁺ refilling by dantrolene or other blockers of ER Ca²⁺ exit channels in disease conditions characterized by lipotoxicity such as metabolic syndrome, diabetes, cardiomyopathy or nonalcoholic steatohepatitis.

Cell Death Discovery (2023)9:100; <https://doi.org/10.1038/s41420-023-01379-0>

INTRODUCTION

Palmitic acid (PA) is an important molecule acting as an effector of lipid injury and metabolic stress in vitro and in vivo. Mechanisms of PA-induced lipotoxicity have been ascribed to endoplasmic reticulum (ER) stress, mitochondrial stress, production of ceramide or lysophosphatidylcholine [1–4], leading to JNK activation, apoptosis or necrosis, depending on the experimental condition and cellular context [3, 5–7]. While ER and mitochondria are well-established target organelles of PA-induced injury [2, 4, 8, 9], lysosomal changes have recently been reported to occur such as decreased lysosomal acidity, altered lysosomal enzyme activity or impaired lysosomal integrity [10–13]. We recently reported that lysosomal Ca²⁺ is released after treatment with PA, while the cellular consequences of lysosomal Ca²⁺ release could be distinct depending on the cell types or context of treatment [11, 12]. Since release of lysosomal Ca²⁺ can also lead to increased cytosolic Ca²⁺ content ([Ca²⁺]_i) [11, 12] and Ca²⁺ is one of the most important inducers of mitochondrial permeability transition (mPT) and subsequent cell death [14, 15], we conducted this investigation based on our hypothesis that lysosomal Ca²⁺ release by PA can lead to lipotoxic cell death through mPT.

RESULTS

PA elicits lysosomal Ca²⁺ release

PA can induce reactive oxygen species (ROS) by compromising mitochondrial complex I and III [11, 16], which can activate lysosomal Ca²⁺ channel [12, 17]. Lysosomal Ca²⁺ release due to lysosomal Ca²⁺ exit channel activation can increase cytosolic Ca²⁺ content ([Ca²⁺]_i), which, in turn, can induce mitophagy, lysosomal stress response [11, 12] or diverse types of cell death [18–20]. Hence, we studied whether PA-induced lipotoxicity entails mitochondrial ROS-induced lysosomal Ca²⁺ release and increased [Ca²⁺]_i.

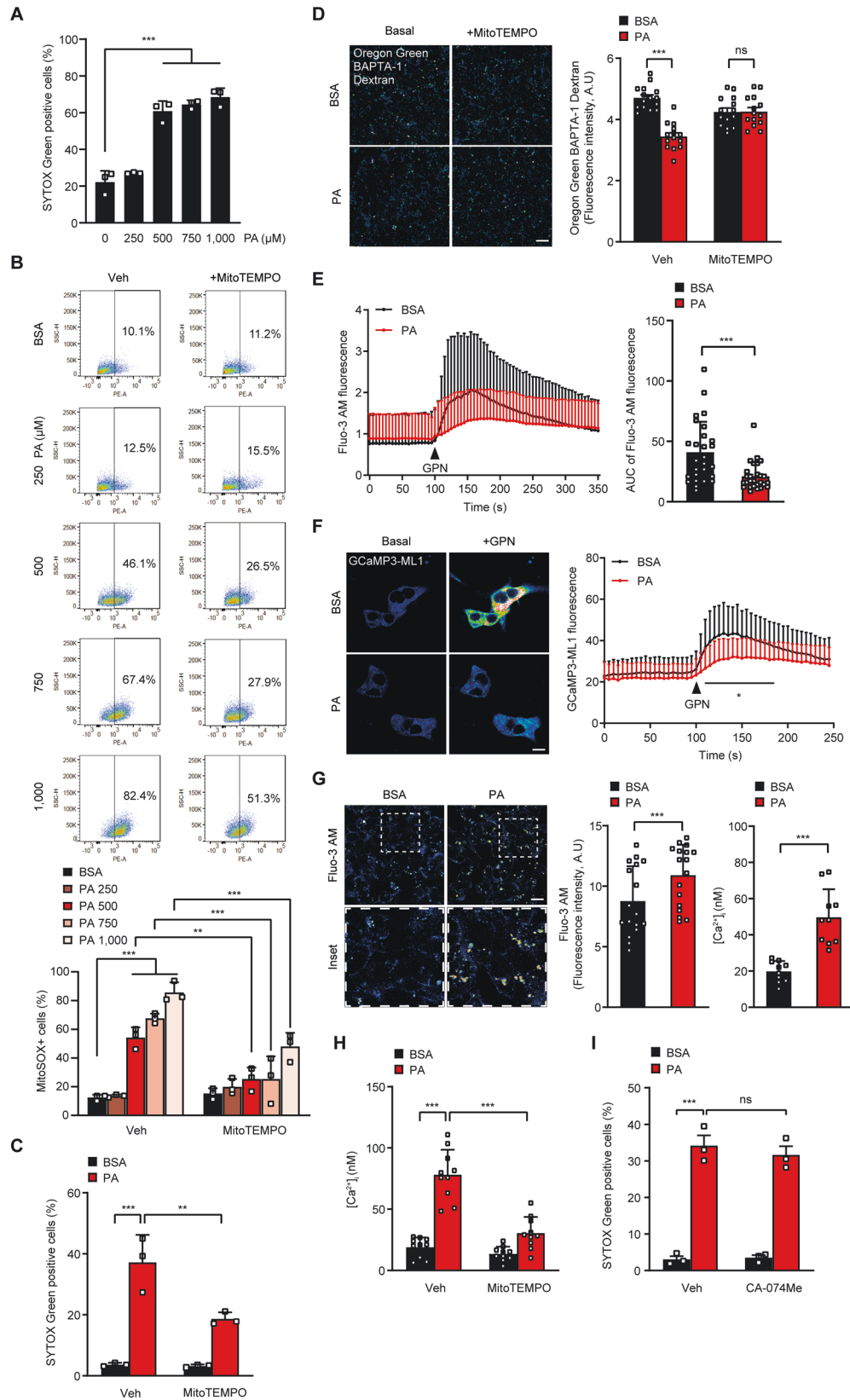
When we treated HepG2 cells with PA, significant lipotoxic cell death was observed in a dose range of 500 ~ 1,000 μM, as revealed by SYTOX Green staining (Fig. 1A). Furthermore, accumulation of mitochondrial ROS stained with MitoSOX was well visualized in the same dose range of PA, which was significantly reduced by MitoTEMPO, a mitochondrial ROS quencher (Fig. 1B), indicating mitochondrial ROS generation by PA. As PA doses higher than 500 μM could be unphysiologically high [21], we employed 500 μM of PA in the following experiments to minimize unphysiological effects of high dose of PA. In an experiment to find causal relationship between lipotoxic cell death and mitochondrial ROS, PA-induced death of HepG2

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Edited by Professor Myung-Shik Lee

Received: 4 September 2022 Revised: 22 February 2023 Accepted: 23 February 2023

Published online: 21 March 2023



cells was significantly reduced by MitoTEMPO (Fig. 1C), suggesting that mitochondrial ROS generation is a critical event in PA-induced lipotoxicity. When we studied possible changes of lysosomal Ca^{2+} after PA treatment, concentration of lysosomal Ca^{2+} ($[\text{Ca}^{2+}]_{\text{Lys}}$)

stained with Oregon Green-488 BAPTA-1 dextran (OGBD) was significantly decreased in HepG2 cells treated with PA (Fig. 1D), suggesting release of lysosomal Ca^{2+} by PA treatment, similar to the results using other types of cells [11, 12]. When we indirectly

Fig. 1 PA reduces lysosomal Ca^{2+} . **A** After treatment of HepG2 cells with 250–1000 μM PA for 24 h, cell death was determined using SYTOX Green. **B** After treatment of HepG2 cells with 250–1000 μM PA for 24 h in the presence or absence of 10 μM MitoTEMPO, PA-induced mitochondrial ROS was measured by MitoSOX followed by flow cytometry (lower panel). Representative scattergrams are shown (upper panel). **C** After treatment of HepG2 cells with 500 μM PA for 24 h in the presence or absence of 10 μM MitoTEMPO, cell death was determined using SYTOX Green. **D** Cells were loaded with Oregon Green-488 BAPTA-1 Dextran (OGBD) for 16 h and chased for 4 h. Cells were then treated with PA or BSA for 6 h in the presence or absence of 10 μM MitoTEMPO, followed by confocal microscopy to determine $[\text{Ca}^{2+}]_{\text{Lys}}$ (right). Representative fluorescence images are shown (left panel). Scale bar, 20 μm . **E** After staining with Fluo-3 AM for 30 min, cells were then treated in PA or BSA for 6 h, followed by confocal microscopy to monitor GPN-induced changes of lysosomal Ca^{2+} release and $[\text{Ca}^{2+}]_{\text{i}}$ increase (left). AUC of the curve was calculated as an estimate of lysosomal Ca^{2+} content (right). **F** Cells transfected with *GCaMP3-ML1* were treated with PA or BSA for 6 h, and then GPN-induced change of *GCaMP3-ML1* fluorescence was determined by confocal microscopy to visualize perilyosomal Ca^{2+} release (right). Representative fluorescence images are shown (left panel). Scale bar, 10 μm . **G** $[\text{Ca}^{2+}]_{\text{i}}$ in HepG2 cells treated with PA for 6 h was determined by staining with Fluo-3 AM (middle) or ratiometric analysis after Fura-2 loading (right). Representative Fluo-3 fluorescence images are shown (left panel). Scale bar, 20 μm . **H** $[\text{Ca}^{2+}]_{\text{i}}$ in HepG2 cells treated with PA for 6 h in the presence or absence of 10 μM MitoTEMPO was determined by ratiometric analysis after Fura-2 loading. **I** After treatment of HepG2 cells with PA in the presence or absence of 10 μM CA-074Me for 24 h, cell death was determined using SYTOX Green. Data are expressed as the means \pm SD of three independent experiments. Statistical comparisons were performed using one-way ANOVA with Tukey's multiple comparison test (**A–D**, **H**, **I**) or two-tailed unpaired Student's *t*-test (vehicle and PA comparisons in **E–G**). (* $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; ns not significant).

estimated lysosomal Ca^{2+} content by calculating area under the curve (AUC) of cytosolic Ca^{2+} concentration ($[\text{Ca}^{2+}]_{\text{i}}$) tracing using Fluo-3 AM staining after treatment with Gly-Phe β -naphthylamide (GPN), a lysosomotropic agent [22], lysosomal Ca^{2+} content was again significantly reduced by PA treatment (Fig. 1E). Decrease of $[\text{Ca}^{2+}]_{\text{Lys}}$ was abrogated by MitoTEMPO (Fig. 1D), suggesting that mitochondrial ROS produced by PA treatment induces lysosomal Ca^{2+} release. We also studied whether perilyosomal Ca^{2+} release can be observed after PA treatment employing *GCaMP3-ML1*, a probe detecting perilyosomal Ca^{2+} release [22]. In *GCaMP3-ML1*-transfected HepG2 cells, perilyosomal Ca^{2+} release was not directly visualized by PA treatment (Fig. 1F). However, perilyosomal Ca^{2+} release after treatment with GPN was markedly reduced (Fig. 1F), suggesting pre-emptying or release of lysosomal Ca^{2+} after PA treatment, similar to the results using other types of cells [12, 17].

When we determined $[\text{Ca}^{2+}]_{\text{i}}$ and studied whether release of lysosomal Ca^{2+} leads to increased cytosolic Ca^{2+} content ($[\text{Ca}^{2+}]_{\text{i}}$), $[\text{Ca}^{2+}]_{\text{i}}$ estimated by Fluo-3 AM staining was significantly increased by PA (Fig. 1G), supporting release of lysosomal Ca^{2+} to the cytosol. When ratiometric measurement of $[\text{Ca}^{2+}]_{\text{i}}$ after Fura-2 staining was conducted to avoid interference due to uneven loading or photobleaching [23], increased $[\text{Ca}^{2+}]_{\text{i}}$ by PA treatment was again well observed, which was reversed by MitoTEMPO (Fig. 1G, H), indicating that mitochondrial ROS-induced lysosomal Ca^{2+} release leads to increased $[\text{Ca}^{2+}]_{\text{i}}$. Since molecules other than Ca^{2+} might be released from lysosome and could affect cell viability, we studied effect of an inhibitor of cathepsin B that has been reported to be released from hepatocytes after PA treatment and to induce lipotoxic cell death [24]. CA-074Me, a cell-permeable inhibitor of cathepsin B, did not inhibit cell death by PA, suggesting no role of lysosomal cathepsin B release in lipotoxic cell death of HepG2 cells (Fig. 1I). Taken together, these data demonstrate that PA-induced mitochondrial ROS leads to release of lysosomal Ca^{2+} to cytoplasm and increased $[\text{Ca}^{2+}]_{\text{i}}$, contributing to lipotoxic HepG2 cell death.

mPT pore (mPTP) opening mediates lysosomal Ca^{2+} loss-driven cell death in response to PA

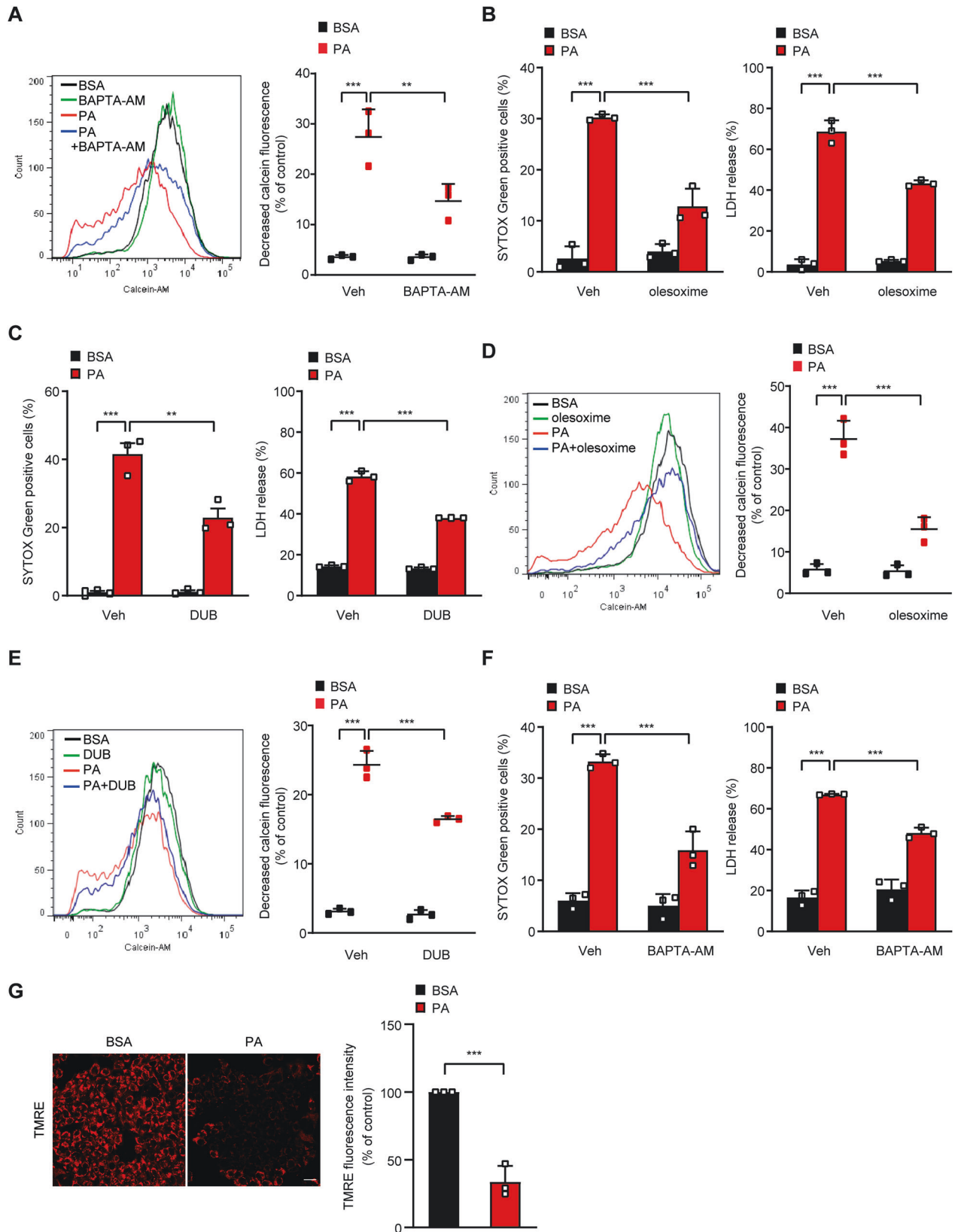
We next studied whether PA-induced release of Ca^{2+} from lysosome into cytosol can induce mPTP opening because Ca^{2+} is one of the most important inducers of mPT which can lead to mitochondrial catastrophe and cell death [2, 25]. When cells were stained by calcein-AM/CoCl₂ staining, mPTP opening visualized by Co²⁺ quenching of mitochondrial matrix calcein fluorescence [26] was well detected after PA treatment of HepG2 cells, showing occurrence of mPT by PA (Fig. 2A). PA-induced mPT was abrogated by BAPTA-AM chelating intracellular Ca^{2+} (Fig. 2A), indicating Ca^{2+} -dependent mPT after PA treatment. Since mPT

can lead to cell death such as necrosis or apoptosis depending on the severity of mPT and cellular context [15, 27, 28], we studied the role of mPT in HepG2 cell death by PA using inhibitors of mPT. Indeed, mPT inhibitors such as oleosoxime [29] or decylubiquinone (DUB) [14], significantly reduced PA-induced cell death assessed by SYTOX Green staining or LDH release assay (Fig. 2B, C). Reduced lipotoxic cell death by oleosoxime or DUB was accompanied by significantly reduced mPT as evidenced by decreased Co²⁺ quenching of calcein fluorescence (Fig. 2D, E), indicating that Ca^{2+} -mediated mPTP opening is an important mechanism of PA-induced lipotoxicity. PA-induced HepG2 cell death was also significantly reduced by BAPTA-AM, confirming the role of increased $[\text{Ca}^{2+}]_{\text{i}}$ and Ca^{2+} -mediated mPTP in lipotoxicity of HepG2 cells (Fig. 2F). As a consequence of mitochondrial damage associated with mPT, mitochondrial potential determined by TMRE staining was significantly lowered by PA treatment of HepG2 cells (Fig. 2G).

ER \rightarrow lysosome calcium flux and SOCE in lipotoxicity

While we showed the role of lysosomal Ca^{2+} release in lipotoxicity, lysosomal Ca^{2+} might not be a sufficient source of Ca^{2+} required for full execution of cellular process requiring Ca^{2+} , since lysosomal volume is 1% of cell volume and lysosomal Ca^{2+} pool is a relatively small Ca^{2+} reservoir [30]. We thus wondered whether release or emptying of lysosomal Ca^{2+} content in HepG2 cells treated with PA could be replenished from endoplasmic reticulum (ER), the largest intracellular Ca^{2+} reservoir [31] which has been observed when lysosomal Ca^{2+} emptying occurs (i.e., ER \rightarrow lysosome Ca^{2+} refilling) [12, 31]. When $[\text{Ca}^{2+}]_{\text{ER}}$ was determined after PA treatment of HepG2 cells using *GEM-CEPIA1er* [32] in a Ca^{2+} -free Krebs-Ringer bicarbonate (KRB) buffer to abolish possible store-operated Ca^{2+} entry (SOCE) [33], Ca^{2+} concentration in the ER ($[\text{Ca}^{2+}]_{\text{ER}}$) was significantly reduced (Fig. 3A), suggesting that ER Ca^{2+} is mobilized probably to replenish lysosomal Ca^{2+} loss. When $[\text{Ca}^{2+}]_{\text{ER}}$ was determined without removal of extracellular Ca^{2+} , $[\text{Ca}^{2+}]_{\text{ER}}$ was not significantly reduced likely due to activation of store-operated Ca^{2+} entry (SOCE) after ER Ca^{2+} emptying (Fig. 3A), which suggests that SERCA was not inhibited by PA since SERCA inhibition reduces $[\text{Ca}^{2+}]_{\text{ER}}$ regardless of extracellular Ca^{2+} [32] and eliminates the possibility of increased $[\text{Ca}^{2+}]_{\text{i}}$ after PA treatment due to SERCA inhibition.

Since these results suggested replenishment of lysosomal Ca^{2+} depletion by ER Ca^{2+} , we next studied whether ER to lysosomal Ca^{2+} movement indeed occurs after PA treatment inducing lysosomal Ca^{2+} release. When PA was removed after treatment for 6 h, a decrease of $[\text{Ca}^{2+}]_{\text{Lys}}$ by PA treatment was recovered (Fig. 3B). To study the role of ER Ca^{2+} in the recovery of reduced $[\text{Ca}^{2+}]_{\text{Lys}}$, we studied the effect of blockade of ER Ca^{2+} exit



channels that could be routes of ER to lysosomal Ca^{2+} movement during the recovery of $[\text{Ca}^{2+}]_{\text{Lys}}$. When we employed Xestospongin C, an IP3R antagonist, recovery of decreased $[\text{Ca}^{2+}]_{\text{Lys}}$ after removal of PA was not significantly affected (Fig. 3B). On the other

hand, dantrolene, an antagonist of ryanodine receptor (RyR), another ER Ca^{2+} exit channel, significantly suppressed the recovery of suppressed $[\text{Ca}^{2+}]_{\text{Lys}}$ after removal of PA (Fig. 3B), suggesting involvement of RyR channel in ER \rightarrow lysosome Ca^{2+}

Fig. 2 mPTP opening by PA due to lysosomal Ca^{2+} release. **A** After treatment of HepG2 cells with 500 μM PA in the presence or absence of 30 μM BAPTA-AM for 24 h, mPTP opening was assessed by flow cytometry after calcein-AM loading and Co^{2+} quenching (right). Representative histograms are shown (left). **B, C** After treatment of HepG2 cells with PA in the presence or absence of 100 μM olesoxime (**B**) or 200 μM DUB (**C**) for 24 h, cell death was determined using SYTOX Green (left) or LDH release assay (right). **D, E** After treatment of HepG2 cells with PA in the presence or absence of 100 μM olesoxime (**D**) or 200 μM DUB (**E**) for 24 h, mPTP opening was determined as in (**A**) (right). Representative histograms are shown (left). **F** After the same treatment as in (**A**), cell death was determined using SYTOX Green (left) or LDH release assay (right). **G** After treatment of HepG2 cells with PA as in (**A**), mitochondrial potential was determined by TMRE staining (right). Representative fluorescence images are shown (left). Scale bar, 20 μm . Data are expressed as the means \pm SD of three independent experiments. All statistical comparisons were performed using one-way ANOVA with Tukey's multiple comparison test. (* $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; ns not significant).

refilling after lysosome Ca^{2+} release by PA. When we chelated ER Ca^{2+} with a membrane-permeant metal chelator *N,N,N',N'*-tetrakis (2-pyridylmethyl) ethylene diamine (TPEN) that has a low Ca^{2+} affinity and can chelate ER Ca^{2+} but not cytosolic Ca^{2+} [34], recovery of $[\text{Ca}^{2+}]_{\text{Lys}}$ after removal of PA was significantly inhibited (Fig. 3B), again supporting the role of ER Ca^{2+} in the recovery of lysosomal Ca^{2+} .

We next studied the effect of blockade of ER Ca^{2+} exit channels on the increase of $[\text{Ca}^{2+}]_i$ after PA treatment. Again, dantrolene but not Xestospongine C, significantly suppressed the increase of $[\text{Ca}^{2+}]_i$ after PA treatment determined by ratiometric measurement following Fura-2 loading (Fig. 3C), suggesting that ER \rightarrow lysosome Ca^{2+} refilling through RyR channel contributes to the increase of $[\text{Ca}^{2+}]_i$ after PA treatment by replenishing decreased lysosomal Ca^{2+} content and sustaining lysosomal Ca^{2+} release. TPEN that reduced recovery of decreased $[\text{Ca}^{2+}]_{\text{Lys}}$ also attenuated increase of $[\text{Ca}^{2+}]_i$ after PA treatment (Fig. 3C). When cell death was determined, dantrolene but not Xestospongine C, significantly suppressed the cell death after PA treatment (Fig. 3D), suggesting that ER \rightarrow lysosome Ca^{2+} refilling through RyR channel contributes to the PA-induced cell death probably by supporting continuous increase of cytosolic Ca^{2+} and subsequent mPTP opening. TPEN also alleviated HepG2 cell death by PA (Fig. 3D), substantiating supportive role of ER Ca^{2+} in lipotoxicity. To study dynamic changes of $[\text{Ca}^{2+}]_{\text{ER}}$ and its temporal relationship with lysosomal Ca^{2+} refilling, we simultaneously traced $[\text{Ca}^{2+}]_{\text{ER}}$ and $[\text{Ca}^{2+}]_{\text{Lys}}$ in cells transfected with *GEM-CEPIA1er* and loaded with OGBD. When organelle $[\text{Ca}^{2+}]$ was monitored in cells that have reduced $[\text{Ca}^{2+}]_{\text{Lys}}$ after PA treatment for 6 h and subsequently were incubated in a Ca^{2+} -free medium, a decrease of $[\text{Ca}^{2+}]_{\text{ER}}$ occurred in parallel with an increase of $[\text{Ca}^{2+}]_{\text{Lys}}$ (Fig. 3E), which strongly supports that ER \rightarrow lysosome Ca^{2+} refilling occurs during HepG2 cell lipotoxicity.

Decrease of $[\text{Ca}^{2+}]_{\text{ER}}$ after PA treatment only in the absence of extracellular Ca^{2+} suggested SOCE without SERCA inhibition because SERCA inhibition would decrease $[\text{Ca}^{2+}]_{\text{ER}}$ regardless of extracellular Ca^{2+} [32]. Thus, we next studied possible occurrence and role of SOCE in lipotoxicity by PA. Since ER luminal protein STIM1 oligomerizes and recruits plasma membrane SOCE channel ORAI1 to activate Ca^{2+} entry when ER Ca^{2+} stores are reduced [33], we studied expression pattern of STIM1 after PA treatment that reduced ER Ca^{2+} store due to ER \rightarrow lysosome Ca^{2+} refilling. In control-treated cells, neither STIM1 oligomerization nor co-localization between STIM1 and ORAI1 was observed (Fig. 4A). In contrast, both STIM1 oligomerization and co-localization between STIM1 and ORAI1 were well observed after PA treatment (Fig. 4A), supporting that reduced ER Ca^{2+} content after PA treatment induced SOCE activation. To validate the role of SOCE in intracellular Ca^{2+} flux, we studied the effect of BTP2, a blocker of SOCE [35, 36] on $[\text{Ca}^{2+}]_{\text{ER}}$. Indeed, in the presence of BTP2, $[\text{Ca}^{2+}]_{\text{ER}}$ was significantly reduced by PA even without removal of extracellular Ca^{2+} . In contrast, without BTP2, PA-induced reduction of $[\text{Ca}^{2+}]_{\text{ER}}$ was not observed in the presence of extracellular Ca^{2+} (Fig. 4B), suggesting the occurrence of SOCE after PA treatment likely due to ER \rightarrow lysosome Ca^{2+} refilling and decreased $[\text{Ca}^{2+}]_{\text{ER}}$ unrelated to SERCA inhibition. Furthermore, the increase of $[\text{Ca}^{2+}]_i$

by PA determined by Fluo-3 AM staining or ratiometric measurement after Fura-2 loading was significantly attenuated by BTP2 (Fig. 4C, D), indicating the role of SOCE in the increase of $[\text{Ca}^{2+}]_i$ by PA. Furthermore, PA-induced death of HepG2 cells was significantly reduced by BTP2 (Fig. 4E), supporting the role of SOCE in Ca^{2+} -mediated lipotoxicity. EGTA chelating extracellular Ca^{2+} also significantly reduced increase of $[\text{Ca}^{2+}]_i$ and death of HepG2 cells after PA treatment (Fig. 4C, F), further supporting the role of extracellular Ca^{2+} flux in the increase of $[\text{Ca}^{2+}]_i$ and subsequent lipotoxicity.

In vivo lipotoxicity

Based on our in vitro results showing the role of lysosomal Ca^{2+} release coupled with ER \rightarrow lysosome Ca^{2+} refilling in mPT and cell death in vitro, we next studied the effect of blockade of lysosomal Ca^{2+} -mediated mPT on lipotoxicity in vivo. When mice were injected with ethyl palmitate followed by LPS administration, death of hepatocytes was observed by TUNEL staining which was accompanied by elevated serum alanine aminotransferase (ALT)/aspartate aminotransferase (AST) (Fig. 5A, B), similar to a previous paper [37]. When dantrolene that was able to inhibit PA-induced hepatocyte death through blockade of the increase of $[\text{Ca}^{2+}]_i$ was administered to mice before ethyl palmitate injection, death of hepatocytes identified by TUNEL staining was significantly ameliorated, suggesting blockade of in vivo lipotoxicity by dantrolene (Fig. 5A). Elevated serum ALT/AST levels after ethyl palmitate followed by LPS administration were also significantly reduced by dantrolene pretreatment (Fig. 5B). In addition, accumulation of mitochondrial ROS stained by MitoSOX was observed in the liver tissue of mice treated with LPS + ethyl palmitate (Fig. 5C), which is in line with mitochondrial ROS accumulation in HepG2 cells treated with PA in vitro. Such mitochondrial ROS accumulation in vivo was ameliorated by dantrolene administration (Fig. 5C). We also studied whether blockade of mPT could inhibit lipotoxicity in vivo based on our in vitro results showing the role of mPT in PA-induced lipotoxicity. When we pretreated mice with DUB, an inhibitor of mPT, hepatocyte death detected by TUNEL staining or elevated serum ALT/AST levels was significantly reduced (Fig. 5A, B), indicating the role of mPT in lipotoxicity in vivo. Mitochondrial ROS in the liver tissue of mice treated with LPS + ethyl palmitate was also reduced by DUB (Fig. 5C). These results showing amelioration of PA-induced hepatocyte death in vivo by dantrolene or DUB, suggest that lipotoxicity of hepatocytes in vivo can be mediated by Ca^{2+} -mediated mPT and indicate the role of lysosomal Ca^{2+} release supported by ER \rightarrow lysosome Ca^{2+} refilling and SOCE in mPT-induced lipotoxicity observed in vitro.

DISCUSSION

It is well established that PA, as an effector of metabolic stress in vitro, can induce stress in diverse organelles such as ER and mitochondria [2, 4, 8, 9]. However, recent investigation showed that PA can induce stress or dysfunction of lysosome as well [10–13], such as elevated pH or reduced Ca^{2+} content due to

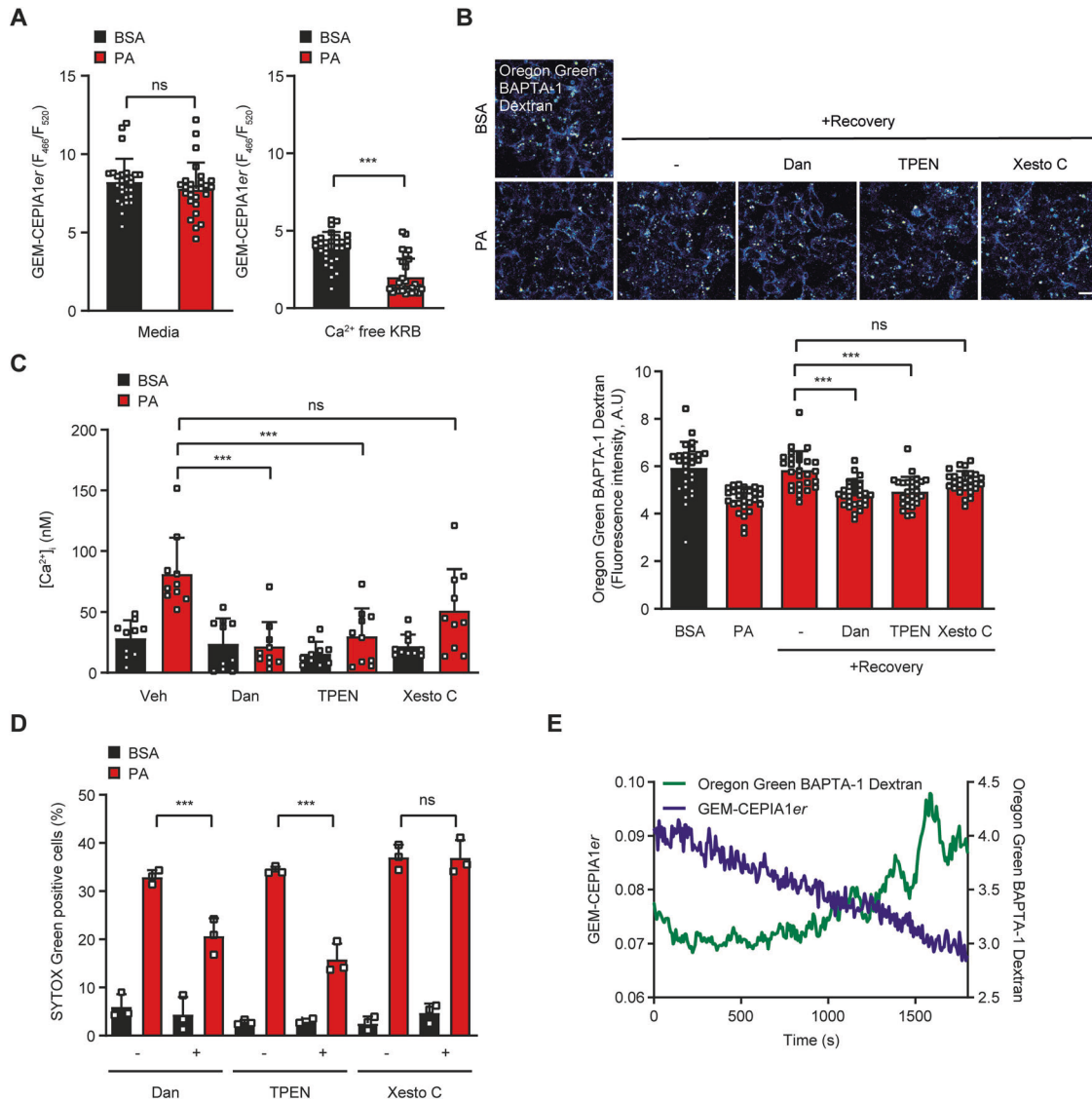


Fig. 3 Role of ER → lysosome Ca²⁺ refilling in lipotoxicity. **A** After treatment of *GEM-CEPIA1er*-transfected HepG2 cells with PA for 6 h in a Ca²⁺-free KRBB (right) or a full medium (left), [Ca²⁺]_{ER} was determined by confocal microscopy. **B** After treatment of OGBD-labeled cells with PA for 6 h, cells were incubated in a fresh medium without PA in the presence or absence of 10 μM dantrolene (Dan), 10 μM TPEN or 3 μM Xestospongine C (Xesto C). Recovery of [Ca²⁺]_{Lys} after removal of PA was determined by confocal microscopy (lower). Representative fluorescence images are shown (upper). Scale bar, 20 μm. **C** After the same treatment as in B, [Ca²⁺]_i was determined by radiometric analysis after Fura-2 loading. **D** After the same treatment as in (B), cell death was evaluated by SYTOX Green staining. **E** After treatment of *GEM-CEPIA1er* transfected cells loaded with OGBD with PA for 6 h and washout, recovery of [Ca²⁺]_{Lys} and changes of [Ca²⁺]_{ER} were monitored simultaneously in the absence of extracellular Ca²⁺. Data are expressed as the means ± SD of three independent experiments. Statistical comparisons were performed using two-tailed Student's *t*-test (A) or one-way ANOVA with Tukey's multiple comparison test (B–D). (**P* < 0.05; ***P* < 0.01; ****P* < 0.001; ns not significant).

release of Ca²⁺ from lysosome. Increased cytosolic Ca²⁺ can mediate diverse beneficial or harmful effects on cells. Since Ca²⁺ is a well-known inducer of mPT [14, 15], we hypothesized that lysosomal Ca²⁺ release might contribute to cell death after treatment with PA or lipotoxicity. Indeed, we observed that Ca²⁺ could be released from lysosome after PA treatment, which is likely due to activation of lysosomal Ca²⁺ exit channel by mitochondrial ROS. After release of lysosomal Ca²⁺, [Ca²⁺]_i was increased, which imposed mPT and subsequent death. Cell death due to mPT caused by PA-induced lysosomal Ca²⁺ release was inhibited by mPT inhibitors such as olesoxime or DUB, showing the role of lysosomal Ca²⁺ release and consequent mPT in lipotoxicity, which is consistent with previous papers reporting mPT induction by PA [2, 38]. Identity of lysosomal Ca²⁺ exit

channel responsible for PA-induced lysosomal Ca²⁺ release is not clear. Previous papers have reported the role of TRPML1 or TRPM2 channel in Ca²⁺- or Zn²⁺-mediated cell death [39–41]. We have also studied the effect of inhibitors of TRPML1 or TRPM2. However, ML-S11, ML-S13 or N-(p-aminocinnamoyl)anthranilic acid inhibiting TRPML1, TRPML1/2/3 and TRPM2, respectively [42–44], did not inhibit PA-induced lipotoxic death of HepG2 cells (Oh S-J et al., unpublished results). Furthermore, Ned-19, an inhibitor of TPC1/2 which has been reported to be associated with ischemia-reperfusion injury [45], was also without effect (Oh S-J et al., unpublished results). Besides such lysosomal Ca²⁺ exit channels, other members of the TRPM family or those belonging to different families might play a role in Ca²⁺-mediated lipotoxic cell death, which could be a subject for further studies. It remains to be

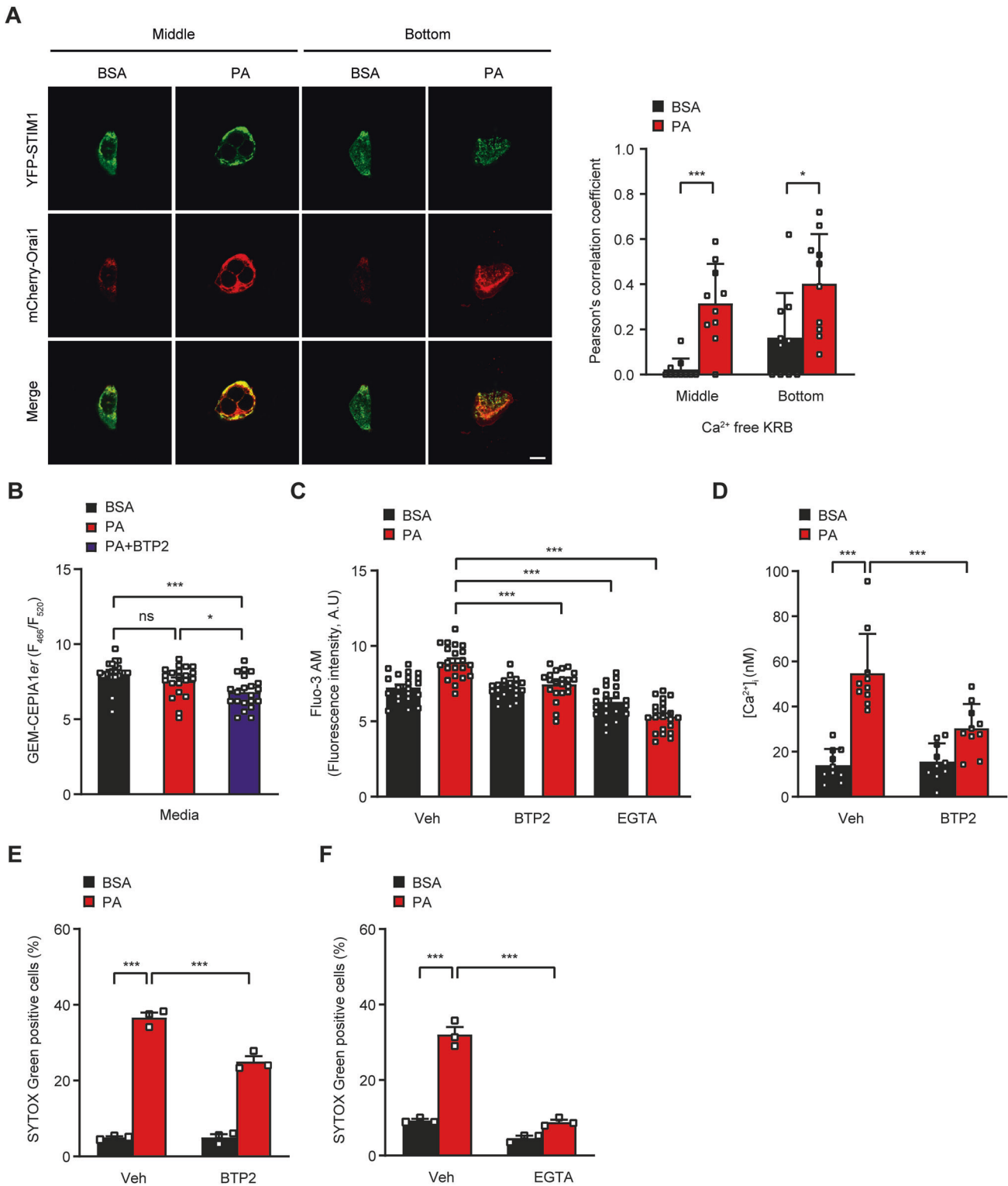


Fig. 4 SOCE in lipotoxicity. **A** After treatment of HepG2 cells transfected with *YFP-STIM1* and *mCherry-Orai1* with PA for 6 h, STIM1 aggregation and its co-localization with ORAI1 were examined at the middle and bottom of the cells by confocal microscopy (left). Scale bar, 10 μ m. Pearson's correlation coefficient was calculated as an index of co-localization of YFP-STIM1 and mCherry-Orai1 (right). **B** After treatment of *GEM-CEPIA1er*-transfected cells with PA in the presence or absence of 10 μ M BTP2 for 6 h, [Ca²⁺]_{ER} was determined by confocal microscopy. **C, D** After the same treatment as in (B), [Ca²⁺]_i was determined by confocal microscopy after Fluo-3 AM loading (C) or ratiometric analysis after Fura-2 loading (D). **E, F** After treatment of cells with PA in the presence or absence of 10 μ M BTP2 (E) or 2 mM EGTA (F) for 24 h, cell death was evaluated using SYTOX Green. Data are expressed as the means \pm SD of three independent experiments. Statistical comparisons were performed using two-tailed Student's *t*-test (A) or one-way ANOVA with Tukey's multiple comparison test (B–F). (**P* < 0.05; ***P* < 0.01; ****P* < 0.001; ns not significant).

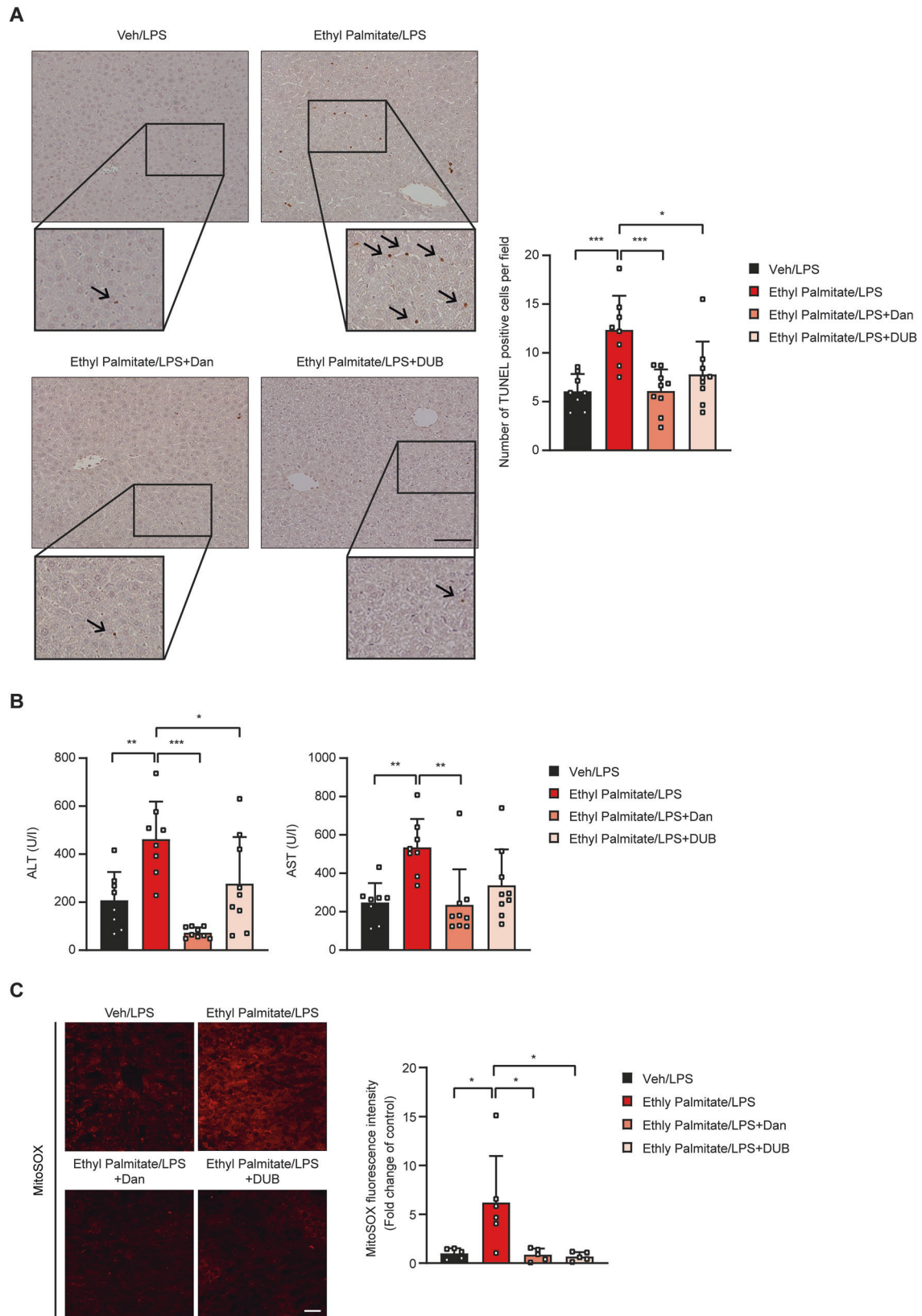


Fig. 5 In vivo lipotoxicity. **A** After injection of ethyl palmitate and LPS to C57BL/6 mice with or without pretreatment with dantrolene or DUB, cell death was evaluated by TUNEL staining of hepatic sections (right). Representative TUNEL staining is presented (left) ($n = 8-9$ /group). Insets were magnified. Scale bar, 200 μm . **B** In serum from mice of (A), ALT and AST levels were determined using a blood chemistry analyzer ($n = 8-9$ /group). **C** In frozen hepatic sections from the mice of (A), mitochondrial ROS accumulation was determined by MitoSOX staining ($n = 5-6$ /group) (right). Representative fluorescence images are shown (left). Scale bar, 20 μm . Data are presented as the means \pm SD. All statistical comparisons were performed using one-way ANOVA with Tukey's multiple comparison test. (* $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; ns not significant).

clarified which lysosomal Ca^{2+} channel is involved in lipotoxicity due to lysosomal Ca^{2+} -mediated mPT.

It has been demonstrated that $\text{ER} \rightarrow$ lysosome Ca^{2+} flux occurs when lysosomal Ca^{2+} content is lowered due to lysosomal Ca^{2+} release by mitochondrial stressors [12, 31]. Such refilling of lysosomal Ca^{2+} pool from ER Ca^{2+} pool occurring when lysosomal Ca^{2+} content is lowered, is likely due to small lysosomal volume accommodating Ca^{2+} which is 10% of ER Ca^{2+} volume [30]. Thus, while $[\text{Ca}^{2+}]_{\text{Lys}}$ is comparable to $[\text{Ca}^{2+}]_{\text{ER}}$, lysosomal Ca^{2+} content might not be sufficient for progression of events requiring Ca^{2+} flux. Besides ER Ca^{2+} pool, another source of lysosomal Ca^{2+} could be endocytic pathway. However, most of endocytic Ca^{2+} has been reported to be dissipated before reaching lysosome [46], supporting the importance of ER Ca^{2+} as a dominant source of lysosomal Ca^{2+} pool. The role of $\text{ER} \rightarrow$ lysosome Ca^{2+} flux in diverse pathological and physiological conditions could be an intriguing topic to be explored in future studies. When we studied whether similar phenomenon occurs after PA treatment of HepG2 cells, $\text{ER} \rightarrow$ lysosome Ca^{2+} refilling could be clearly demonstrated by simultaneous monitoring of $[\text{Ca}^{2+}]_{\text{ER}}$ and $[\text{Ca}^{2+}]_{\text{Lys}}$ during incubation in a Ca^{2+} -free medium ensuring recovery of $[\text{Ca}^{2+}]_{\text{Lys}}$ after removal of PA. Furthermore, blockade of $\text{ER} \rightarrow$ lysosome Ca^{2+} refilling with dantrolene, an inhibitor of RyR Ca^{2+} exit channel on ER or chelation of ER Ca^{2+} by TPEN could inhibit lipotoxicity, which indicates that intracellular Ca^{2+} flux from ER occurs to sustain lysosomal Ca^{2+} release, resulting in cell death. Ca^{2+} flux from ER to lysosome, in turn, induced ER Ca^{2+} emptying which activated SOCE. Occurrence of SOCE after PA treatment of HepG2 cells was evidenced by no decrease of $[\text{Ca}^{2+}]_{\text{ER}}$ in cells treated with PA without removal of extracellular Ca^{2+} , reappearance of the decrease of $[\text{Ca}^{2+}]_{\text{ER}}$ by BTP2 even in the presence of extracellular Ca^{2+} and STIM1 aggregation co-localized with ORAI1. Functional role of SOCE after treatment with PA was demonstrated by inhibition of lipotoxicity by BTP2 or extracellular Ca^{2+} chelation with EGTA. Furthermore, occurrence of SOCE and no decrease of $[\text{Ca}^{2+}]_{\text{ER}}$ in cells treated with PA without removal of extracellular Ca^{2+} supports that increase of $[\text{Ca}^{2+}]_{\text{i}}$ after PA treatment is not due to direct release of Ca^{2+} from ER through SERCA inhibition, since increase $[\text{Ca}^{2+}]_{\text{i}}$ due to SERCA inhibition would not be affected by extracellular Ca^{2+} [32]. While SERCA inhibition is an important component of ER stress associated with lipid overload or obesity in vivo inducing altered membrane phospholipid composition [47], SERCA might not be important in acute lipotoxicity of HepG2 cells in vitro. However, we do not eliminate the possibility that ER stress might contribute to lipotoxicity in vitro. In fact, we have observed that CHOP, an important player in ER stress-induced cell death, can be activated by mitochondrial ROS produced after PA treatment [48].

In our experiments to investigate the role of increased Ca^{2+} and mPT in lipotoxicity in vivo, we observed that DUB inhibiting mPT and dantrolene abrogating $[\text{Ca}^{2+}]_{\text{i}}$ increase and cell death in vitro by PA could inhibit lipotoxic death of hepatocytes in vivo. Reduction of mitochondrial ROS by dantrolene or DUB on mitochondrial ROS could be due to a feed-forward response leading to further ROS release following Ca^{2+} -mediated mPT [49, 50] which was inhibited by dantrolene or DUB. Dantrolene is a candidate for therapeutic drug against malignant hyperthermia or Alzheimer's disease [51, 52]. DUB has also been studied as a potential therapeutic agent against Friedreich's ataxia [53] or tumor-induced angiogenesis [54]. Our results suggest the possibility that dantrolene or DUB could be considered candidates for drug agents against diseases characterized by lipotoxicity such as nonalcoholic steatohepatitis or cardiomyopathy. While we have shown the contribution of mPT induced by lysosomal Ca^{2+} release as a mechanism of lipotoxicity in vitro and in vivo, other previously reported mechanisms of lipotoxicity such as ER or mitochondrial stress and generation of ceramide or lysophosphatidylcholine [1–4] could also play an important role depending on the cellular context or environmental condition, and optimal strategy against lipotoxicity could be different accordingly.

MATERIALS AND METHODS

Reagents

Reagents in this study were purchased from following sources: Oregon Green-488 BAPTA-1 dextran (OGBD, O6789), Fluo-3 AM (F23915), MitoSOX™ Red (M36008), SYTOX Green (S7020), BAPTA-AM (O6807), calcein-AM (C1430), Fura-2 (F1221) from Thermo Fisher Scientific; palmitate (PA, P0500), decylubiquinone (DUB, D7911), bovine serum albumin (BSA, A9418), cobalt chloride (CoCl_2 , 232696), ethylene glycol-bis (2-aminoethyl-ether)-N,N,N',N'-tetraacetic acid (EGTA, 03777), MitoTEMPO (SML0737), CA-074Me (205531) from Sigma-Aldrich; Gly-Phe β -naphthylamide (GPN, ab145914) from Abcam; BTP2 (S8380) from Selleckchem; CytoTox 96® Non-Radioactive Cytotoxicity Assay (G1780) from Promega™ Corporation.

Cell culture and treatment

HepG2 cells obtained from Korean Cell Line Bank (KCLB) were grown in DMEM medium (Welgene, LM-001-05)-1% penicillin–streptomycin–amphotericin B mixture (Lonza, 17-745E) supplemented with 10% fetal bovine serum (Corning, 35-010-CV). Cells were tested for mycoplasma contamination using a Mycoplasma PCR Detection Kit (e-Myco™, 25236, iNTRON Biotechnology). PA stock solution was prepared by dissolving palmitate in 70% ethanol and heating at 56 °C. PA stock solution was diluted in 2% fatty acid-free BSA-DMEM before treatment. For in vitro treatment, following concentrations were employed: OGBD, 100 $\mu\text{g}/\text{ml}$; Fluo-3 AM, 5 μM ; MitoSOX™ Red, 5 μM ; SYTOX Green Nucleic Acid Stain, 1 μM ; BAPTA-AM, 30 μM ; calcein-AM, 1 μM ; palmitic acid (PA), 500 μM ; decylubiquinone (DUB), 200 μM ; cobalt chloride (CoCl_2), 2 mM; Gly-Phe β -naphthylamide (GPN), 200 μM ; BTP2, 10 μM ; MitoTEMPO, 10 μM ; EGTA, 2 mM; CA-074Me, 10 μM .

SYTOX Green nucleic acid staining

Cell death was evaluated using SYTOX™ Green Nucleic Acid Stain kit (Thermo Fisher Scientific, S7020). Briefly, HepG2 cells were seeded in a 24-well plate. After 24 h, cells were treated with PA with or without indicated compound for 24 h. Cells were then incubated with SYTOX Green (1 μM) for 30 min at 37 °C. SYTOX Green fluorescence was measured by flow cytometry using BD FACSVerser and FACSCanto II (BD Biosciences, San Jose, CA, USA). Data analysis was performed using FlowJo software (10.8, FlowJo, LLC, BD Biosciences).

LDH release assay

Cell death was assessed using an LDH release assay kit (Promega™ Corporation, G1780). Briefly, HepG2 cells seeded 4×10^4 well in a 96-well plate were treated with PA with or without indicated compound for 24 h. Culture supernatant was collected and analyzed according to the manufacturer's protocol.

Transfection and plasmids

Cells were transfected with plasmids such as *GCaMP3-ML1*, *GEM-CEPIA1er*, *YFP-STIM1*, *3xFLAG-mCherry Red-Orai1/P3XFLAG7.1* using PolyJet™ In Vitro DNA Transfection Reagent (SigmaGen® Laboratories, SL100688), according to the manufacturer's protocol.

Measurement of cytosolic, ER and lysosome Ca^{2+} contents

To measure $[\text{Ca}^{2+}]_{\text{Lys}}$, HepG2 cells grown on a chambered coverglass (Thermo Fisher Scientific, 155383) were loaded with 100 $\mu\text{g}/\text{ml}$ OGBD, an indicator of lysosomal luminal Ca^{2+} for 16 h. After incubation in a fresh media for 4 h, cells were treated with BSA or 500 μM PA for 6 h. After washing with Ca^{2+} -free HEPES-buffered saline (HBS), fluorescence was measured using an LSM780 confocal microscope (Zeiss, Oberkochen, Germany) and quantified using ImageJ software.

To determine $[\text{Ca}^{2+}]_{\text{i}}$ by confocal microscopy, HepG2 cells grown on chambered coverglass were loaded with 5 μM Fluo-3 AM for 30 min, an indicator of cytosolic Ca^{2+} . After treatment with 500 μM PA for 6 h, cells were washed with Ca^{2+} -free PBS twice, and fluorescence was recorded using an LSM780 confocal microscope. Fluorescence intensity was quantified using ImageJ software. For ratiometric $[\text{Ca}^{2+}]_{\text{i}}$ measurement, cells were loaded with 2 μM of the acetoxymethyl ester form of Fura-2 in DMEM at 37 °C for 30 min and then washed with a basal Ca^{2+} solution (145 mM NaCl, 5 mM KCl, 3 mM MgCl_2 , 10 mM glucose, 1 mM EGTA, 20 mM HEPES, pH 7.4). Measurements were conducted using MetaFluor on an Axio Observer A1 (Zeiss) equipped with a 150 W xenon lamp Polychrome V (Till Photonics, Bloaom Drive Victor, NY, USA), a CoolSNAP-Hq2 digital camera (Photometrics, Tucson, AZ, USA), and a Fura-2 filter set.

Fluorescence at 340/380 nm was measured in phenol red-free medium, and converted to $[Ca^{2+}]_i$ using the following equation [23].

$$[Ca^{2+}]_i = K_d \times [(R - R_{min}) / (R_{max} - R)] \times [F_{min(380)} / F_{max(380)}]$$

where K_d = Fura-2 dissociation constant (224 nM at 37 °C), $F_{min(380)}$ = the 380 nm fluorescence in the absence of Ca^{2+} , $F_{max(380)}$ = 380 nm fluorescence with saturating Ca^{2+} , R = 340/380 nm fluorescence ratio, R_{max} = 340/380 nm ratio with saturating Ca^{2+} , and R_{min} = 340/380 nm ratio in the absence of Ca^{2+} .

To determine ER Ca^{2+} contents ($[Ca^{2+}]_{ER}$), HepG2 cells were grown on chambered coverglass and transfected with a *GEM-CEPIA1er* plasmid, a ratiometric fluorescent indicator of ER Ca^{2+} . After 24 h, cells were treated with BSA or 500 μ M PA for 6 h, and then washed with Ca^{2+} -free KRBB (Sigma-Aldrich, K4002). *GEM-CEPIA1er* fluorescence was measured using an LSM780 confocal microscope at an excitation wavelength of 405 nm and an emission wavelength of 466 or 520 nm. Fluorescence ratio F466/F520 was calculated as an index of $[Ca^{2+}]_{ER}$ [32].

GCaMP3-ML1 Ca^{2+} imaging

To detect perilyosomal Ca^{2+} release, HepG2 cells were transfected with a *GCaMP3-ML1* Ca^{2+} probe, a lysosome-targeted genetically-encoded Ca^{2+} sensor. Twenty-four h after transfection, cells were treated with BSA or 500 μ M PA for 6 h. Perilyosomal Ca^{2+} release was recorded by monitoring *GCaMP3* fluorescence intensity at 470 nm in a basal Ca^{2+} solution (145 mM NaCl, 5 mM KCl, 3 mM $MgCl_2$, 10 mM glucose, 1 mM EGTA, 20 mM HEPES, pH 7.4) using a Zeiss LSM780 confocal microscope. GPN, a lysosomotropic agent, of 200 μ M was added to evoke lysosomal Ca^{2+} release. *GCaMP3-ML1* fluorescence was calculated as a change *GCaMP3* fluorescence (ΔF) over baseline fluorescence (F_0).

STIM1 and ORAI1 co-localization

After transfection with *YFP-STIM1* and *3xFLAG-mCherry Red-Orai1/P3XFLAG7.1* (provided by Yuan J through Cha S-G) plasmids for 48 h, HepG2 cells were treated with BSA or 500 μ M PA for 6 h. Cells were then fixed with 4% paraformaldehyde (PFA) at room temperature for 10 min. Fluorescence images were acquired with an LSM780 confocal microscope, and the co-localization between STIM1 and ORAI1 was examined by calculating Pearson's correlation coefficient using ZEN software (Carl Zeiss Microscopy GmbH, Jena, Germany).

Measurement of mitochondrial ROS

Mitochondrial ROS was measured using MitoSOXTM Red. Briefly, HepG2 cells were pretreated with 10 μ M MitoTEMPO for 1 h, and then treated with 500 μ M PA for 24 h. After incubation with 5 μ M MitoSOXTM Red at 37 °C for 15 min, fluorescence was measured by flow cytometry using BD FACSVerser or FACSCanto II. Data were analyzed using FlowJo software.

To determine mitochondrial ROS accumulation *in vivo*, frozen sections of the liver tissue were incubated with MitoSOXTM Red at 37 °C for 15 min, and then washed with PBS. Fluorescence images were observed with an LSM710 confocal microscope and fluorescence intensity was quantified using ImageJ software (NIH, Bethesda, MD, USA).

mPTP opening

mPTP opening was assessed by Co^{2+} quenching of calcein-AM fluorescence. Cells were treated with PA (500 μ M) with or without DUB (200 μ M) or oleosoxime (100 μ M) for 24 h. Cells were then loaded with calcein-AM (1 μ M) and $CoCl_2$ (2 mM) at 37 °C for 15 min. Calcein fluorescence was measured by flow cytometry employing BD FACSVerser and FACSCanto II and analyzed using FlowJo software.

Mitochondrial membrane potential

Mitochondrial membrane potential was determined using tetramethylrhodamine ethyl ester perchlorate (TMRE). After treatment of HepG2 cells with PA for 24 h, cells were incubated with 100 nM TMRE at 37 °C for 30 min, and fluorescence was measured using LSM710 confocal microscope. Fluorescence intensity was quantified using ImageJ software.

Animals

Eight-week-old C57BL/6N male mice were purchased from Orient Bio (Seongnam, Korea). All experiments using mice were performed in accordance with the guidelines of the Public Health Service Policy in

Humane Care and Use of Laboratory Animals. The protocol was approved by the Institutional Animal Care and Use Committee (IACUC) of the Department of Laboratory Animal Resources of Soonchunhyang Institute of Medi-bio Science. Ethyl palmitate (Tokyo Chemical Industry, P0003) was dissolved in water with 4.8% lecithin (FUJIFILM Wako Pure Chemical Corporation, 120-00832) and 10% glycerol (Sigma-Aldrich, G2025) to make a mixture containing ethyl palmitate at a concentration of 300 mM. LPS (Sigma-Aldrich, E. coli O55:B5) was dissolved in PBS, and 0.025 mg/kg LPS was injected intraperitoneally into mice. Dantrolene and DUB were dissolved in DMSO and diluted with PBS. For *in vivo* administration, dantrolene (10 mg/kg), DUB (5 mg/kg) or DMSO solution diluted in PBS was injected into mice that were fasted for 24 h. After 1 h, 300 mM ethyl palmitate or vehicle was administered, followed by LPS injection 1 h later. Blood and liver tissue samples were obtained 24 h later.

Blood chemistry

Serum ALT and AST levels were measured using a Fuji Dri-Chem NX500i chemistry Analyzer (Fujifilm, Tokyo, Japan).

TUNEL staining

Paraffin-embedded liver tissue blocks were prepared by fixing in 10% neutral buffered formalin (Sigma, HT501128). TUNEL staining was conducted using In Situ Cell Death Detection kit (Roche, 11684817910) according to the manufacturer's instructions. Cell death index was expressed as the number of TUNEL-positive cells per field counted from more than 20 fields randomly selected ($\times 200$).

Statistical analysis

Data are presented as means \pm SD of three independent experiments. Two-tailed Student's *t*-test was used to compare values between two groups. One-way ANOVA with Tukey's test was used to compare values between multiple groups. Data were considered significant when $P < 0.05$. Data analysis was performed using GraphPad Prism 8 software (GraphPad Software, La Jolla, CA, USA).

DATA AVAILABILITY

The data generated or analyzed during the current study and materials are available from the corresponding author without imposed restriction on reasonable request.

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AUTHOR CONTRIBUTIONS

M-SL and KYH conceived the experiments. S-JO and YH conducted the experiments. M-SL, KYH, S-JO and YH wrote the manuscript.

FUNDING

This study was supported by a National Research Foundation of Korea (NRF) grant funded by the Korea government (MSIT) (NRF-2019R1A2C3002924) and by the Bio&Medical Technology Development Program (2017M3A9G7073521). M-S Lee is the recipient of Korea Drug Development Fund from the Ministry of Science & ICT, Ministry of Trade, Industry & Energy and Ministry of Health & Welfare (HN22C0278), and a grant from the Faculty Research Assistance Program of Soonchunhyang University (20220346).

COMPETING INTERESTS

M-SL in the CEO of LysoTech, Inc. The authors declare no competing interests.

ADDITIONAL INFORMATION

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