

SIRT3 Acts As a Negative Regulator of Autophagy Dictating Hepatocyte Susceptibility to

Lipotoxicity

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Abbreviation:

NAFLD, non-alcoholic fatty liver disease; SA, stearic acid; PA, palmitic acid; SFAs, saturated fatty acids; ER, endoplasmic reticulum; SIRT6, Sirtuins; NAD, nicotinamide adenine dinucleotide; AMPK, AMP-activated protein kinase; OA, oleic acid; BSA, bovine serum albumin; ALT, alanine aminotransferase; DMEM, Dulbecco's Modified Eagle Medium; LDH, lactate dehydrogenase; PI, propidium iodide; TEM, Transmission electron microscopy; CQ, chloroquine; IP immunoprecipitation; ROS, Reactive oxygen species; $O_2^{\cdot-}$, intracellular superoxide; H_2O_2 , hydrogen peroxide; MAP1, microtubule-associated protein 1; LC3, microtubule-associated protein 1 light chain 3; EBSS, Earle's Balanced Salt Solution; SCD-1, stearoyl-CoA desaturase-1; NAM, nicotinamide; Baf, bafilomycin A1; mTORC1, mammalian target of rapamycin C1; AICAR, 5-Aminoimidazole-4-carboxamide1- β -D-ribofuranoside; NAC, N-acetylcysteine, MnSOD, manganese superoxide dismutase; 2-ME, 2-methoxyestradiol.

Abstract

Lipotoxicity induced by saturated fatty acids (SFAs) plays a central role in the pathogenesis of non-alcoholic fatty liver disease (NAFLD); however, the exact mechanism(s) remain to be fully elucidated. SIRT3 is an NAD^+ -dependent deacetylase primarily located inside mitochondria. In this study, we demonstrated that a SFAs-rich high-fat diet (HFD) was more detrimental to the liver than an isocaloric unsaturated FAs-rich HFD. Unexpectedly, SIRT3 expression/activity were significantly elevated in the livers of mice exposed to the SFAs-rich HFD. Using cultured HepG2 and AML-12 hepatocytes, we demonstrated that unlike monounsaturated FAs, SFAs upregulates SIRT3 expression/activity. SIRT3 overexpression renders both the liver and hepatocytes susceptible to palmitate-induced cell death, which can be alleviated by SIRT3 siRNA transfection. In contrast, SIRT3 suppression protects hepatocytes from palmitate cytotoxicity. Further studies revealed that SIRT3 acts as a negative regulator of autophagy, whereby enhancing the susceptibility of hepatocytes to SFAs-induced cytotoxicity. Mechanistic investigations elucidate that SIRT3 overexpression causes manganese superoxide dismutase (MnSOD) deacetylation/activation, which depleted intracellular superoxide contents, leading to AMP-activated protein kinase (AMPK) inhibition and mTORC1 activation, resulting in autophagy suppression. In contrast, SIRT3 siRNA gene silencing enhances autophagy flux. The similar result was observed in the liver tissue from SIRT3 knockout mice.

Conclusion: our data identified SIRT3 to be a novel negative regulator of autophagy, whose activation by SFAs contributes to lipotoxicity in hepatocytes and suggest that restraining SIRT3 overactivation can be a potential therapeutic choice for the treatment of NAFLD as well as other metabolic disorders, with lipotoxicity being the principal pathomechanism.

Introduction

Lipotoxicity refers to cellular dysfunction and/or cell death derived from excess exposure to elevated levels of fatty acids and lipid accumulation in non-adipose tissues, including liver, heart, skeletal muscle, and pancreas.⁽¹⁾ Lipotoxicity accounts for many manifestations of the metabolic syndrome, including non-alcoholic fatty liver disease (NAFLD).⁽²⁾ Hepatocyte cell death resulting from lipotoxicity, also called lipoapoptosis, is a hallmark of NAFLD.⁽³⁾ Overall, lipotoxicity is most commonly induced by saturated fatty acids (SFAs), in specific palmitic acid (16:0, PA), the most abundant SFA in the diet and in the circulation.⁽⁴⁾ The cellular mechanisms underlying SFAs-induced lipotoxicity in hepatocytes remain to be fully understood; however, it is well documented that endoplasmic reticulum (ER) stress and oxidative stress are the major intracellular pathways mechanistically involved in lipoapoptosis.^(5, 6) Protecting hepatocytes from SFAs-induced hepatotoxicity provides a conceivable strategy for preventing the development of NAFLD.

Autophagy is an evolutionarily conserved physiological process that represents a system of bulk protein degradation aimed at removal and breakdown of cellular components (organelles and proteins) during starvation, thereby redistributing nutrients for maintenance of cellular energetic balance.⁽⁷⁾ It also plays a critical role in eliminating damaged proteins and organelles.⁽⁸⁾ We previously reported that autophagy activation protected hepatocytes against SFAs-induced lipotoxicity.⁽⁹⁾ Similarly, autophagy activation confers protection against palmitate-induced cytotoxicity in pancreatic beta-cells.⁽¹⁰⁾

Sirtuins (SIRT) are highly conserved nicotinamide adenine dinucleotide (NAD⁺)-dependent deacetylases that regulate a wide variety of biological functions.⁽¹¹⁻¹⁴⁾ Of the seven mammalian sirtuins, SIRT3, SIRT4, and SIRT5 are predominantly localized inside mitochondria,

with SIRT3 being considered as the primary modulator of mitochondrial protein acetylation.⁽¹⁵⁾ SIRT3 is highly expressed in metabolically active tissues, such as the liver, and is a key regulator of mitochondrial functions, including metabolism, energy production, and modulation of the response to oxidative stress.⁽¹⁶⁻²¹⁾ The function of SIRT3 in cellular response is stimulus- and cell type-specific,⁽²²⁾ with both anti-apoptotic and pro-apoptotic activities being reported.⁽²³⁻²⁵⁾

The present study aims to investigate the role of SIRT3 in SFAs-induced hepatotoxicity. Unexpectedly, we demonstrate that the sensitivity of hepatocytes to SFAs-induced lipotoxicity is positively associated with SIRT3 activity. Our detailed mechanistic investigations provide evidence for a novel role of SIRT3 in the regulation of autophagy induction via an AMP-activated protein kinase (AMPK)-dependent mechanism.

Materials and Methods

Animal Model and Experimental Protocol

All animal studies were performed in accordance with the animal care committee at the University of Illinois at Chicago. All mice were fed *ad libitum*, and housed on a 12:12 h light:dark cycle at 25°C. Male SIRT3^{-/-} mice and WT 129Sv mice (8-week of age) were purchased from Jackson Laboratory (Bar Harbor, ME), and maintained on a standard chow diet. Male C57BL/6 mice (8-week old, Jackson Laboratory, Bar Harbor, ME) were fed with normal (n = 8) or high-fat diet using either palm oil (n = 8) or corn oil (n = 8) as fat source (Supplementary data, Table. 1, Bio-Serv, Flemington, NJ). For liver-specific SIRT3 overexpression, the animals were injected in the lateral tail vein with recombinant adeno-associated viral (AAV) serotype 8 gene transfer vectors bearing a liver-specific promoter combination (albumin promoter) with mouse SIRT3 sequence. The detailed protocols were shown in the online supplementary data. Mice were sacrificed 8 weeks later.

Establishment of stable human SIRT3 over-expression cell lines

The eukaryotic expression vector plasmid pcDNA3.1+/hSIRT3 containing the human SIRT3 gene sequence was kindly provided by Dr. Yu Wang (University of Hong Kong).⁽²⁶⁾ HepG2 cells were transfected with 0.8 µg/well of the expression vector pcDNA3.1+/hSIRT3 or empty vector control pcDNA3.1+ (Invitrogen, Grand Island, NY) in 24-well plate using Lipofectamine 2000 reagent (Invitrogen, Grand Island, NY) following the manufacturer's guidelines. After 48 hours transfection, cells were trypsinized into 24-well plates at a ratio of 1:100 for monoclonal cell selection using 800 µg/ml G418. The stable cell line was established and confirmed by both western blot analysis and Real Time PCR.

MnSOD Overexpression

Recombinant lentivirus vector (pLV[Exp]-EGFP:T2A:Neo) containing human MnSOD (NM_001024465.1) under the control of EF-1alpha promoter was generated by Cyagen Biosciences Inc. (Guangzhou, China). A noncoding vector carrying the EF1alpha promoter was used to produce vector control particles. The optimal virus titer used for cell transfection was screened according to the manufacturer's instructions. HepG2 cells were transfected with either pLV[Exp]-EGFP:T2A:Neo-EF1alpha-hMnSOD to overexpress MnSOD or pLV[Exp]-EGFP:T2A:Neo-EF1alpha-null as a vector control.

Analysis of autophagic flux

The autophagic flux was measured as described previously.⁽²⁷⁾ Briefly, the cells were pretreated with chloroquine (CQ), an inhibitor of lysosome acidification, after SIRT3 overexpression or knockdown. The autophagic flux was determined by detecting GFP-LC3 puncta and LC3-II expression using laser scanning confocal microscope (Nikon A1R, Japan) and Western blot, respectively. Additionally, cells were transiently transfected with recombinant adenovirus mRFP-GFP-LC3 (Hanbio Biotechnology Co. Ltd., Shanghai, China). Yellow or red puncta was detected by confocal microscope from at least 50 cells for each individual experiment after different treatment.

Statistical Analysis

All data are expressed as mean \pm SD of at least three independent experiments. Statistical analysis was performed using a one-way ANOVA and analyzed by post-hoc test with Fisher's least significant difference (LSD). Differences between treatments were considered to be statistically significant at $p < 0.05$.

Results

SIRT3 Negatively Regulates Autophagy in Hepatocytes.

To directly test the effect of SIRT3 on autophagy in hepatocytes, a stable SIRT3-overexpressing HepG2 cell line was established (Supplementary data, Fig. 1). SIRT3 overexpression strongly decreased intracellular autophagosome formation and microtubule-associated protein 1 (MAP1) light chain 3 (LC3)-II conversion, the well-established markers of autophagy induction, while increased accumulation of p62, a protein specifically degraded by autophagy, was observed (Fig. 1A). Moreover, SIRT3 overexpression blunted rapamycin-induced autophagy activation (Supplementary data, Fig. 2) and attenuated autophagic flux, which was evidenced by reduced autophagosome formation and LC3-II conversion in the presence of CQ, and decreased autophagosomes (yellow puncta) and autolysosomes (red puncta) number (Fig. 1A). Conversely, SIRT3 knockdown via siRNA transfection led to increased LC3-II conversion, decreased p62 accumulation (Fig. 1B; Supplementary data, Fig. 3), and increased autophagic flux (Fig. 1B). Furthermore, SIRT3^{-/-} and the liver-specific SIRT3 overexpression mice were collected. As shown in Fig. 1C, SIRT3^{-/-} mice manifested enhanced hepatic LC3-II conversion and decreased accumulation of p62 in both fasting and fed states (Fig. 1C), while the mice with liver-specific SIRT3 overexpression manifested decreased LC3-II conversion and increased accumulation of p62 in the liver (Fig. 1D). These data altogether suggest that SIRT3 is a negative regulator of autophagy.

FFAs Differentially Regulate SIRT3 Expression in the Liver/Hepatocytes.

SFAs-induced hepatocyte cell death contributes to the pathogenesis of NAFLD ⁽⁹⁾. We showed that both PA and SA induced significant cell death (Supplementary data, Fig. 4A-F), while oleic acid (OA) showed no effect on cell viability (Supplementary data, Fig. 4G & H). In our *in vivo*

investigations, male C57BL/6 mice were exposed to either a SFAs-rich (palm oil) high-fat diet (HFD) or an isocaloric unsaturated fatty acids (unSFAs)-rich (corn oil) HFD for 8 weeks. We demonstrated here that the SFAs-rich HFD more prominent hepatic pathological alterations than the isocaloric unSFAs-rich HFD (Fig. 2A; Supplementary data, Fig. 5). Intriguingly, hepatic SIRT3 expression was significantly elevated in the setting of SFAs-rich HFD feeding (Fig. 2B; Supplementary data, Fig. 6A). Correspondingly, mitochondrial protein acetylation status in the liver was significantly reduced in mice fed the SFAs-rich HFD (Fig. 2C).

To directly determine the effect of different types of fatty acids on SIRT3 expression and activity, cultured hepatocytes were exposed to either PA or OA. PA exposure increased SIRT3 expression in a dose- and time-dependent manner in both human (HepG2) and mouse (AML-12, a non-transformed mouse hepatocyte cell line) hepatocytes (Fig. 2D; Supplementary data, Fig. 7A & B) without affecting its stability (Supplementary data, Fig. 6B & C). Accordingly, acetylated mitochondrial protein abundance was reduced by PA exposure (Fig. 2E; Supplementary data, Fig. 7C). OA exposure had no effect on SIRT3 expression (Fig. 2F).

SIRT3 Controls Sensitivity of Hepatocyte/Liver to SFAs-induced Lipotoxicity.

SIRT3 upregulation by PA, but not OA, spurred us to investigate the role of SIRT3 activity in SFA-induced lipotoxicity. Somewhat unexpectedly, SIRT3 overexpression increased cellular susceptibility to PA-induced cell death (Fig. 3A). In contrast, SIRT3 inhibition, with either genetic (siRNA transfection) or pharmacologic (nicotinamide, NAM) approach, alleviated PA-induced cell death in both control and SIRT3-overexpressing cells (Fig. 3B-D; Supplementary data, Fig. 8 & 9). Similar results were observed when SA was applied (Supplementary data, Fig. 10). These results suggest that SIRT3 activity modulates the sensitivity of hepatocyte to SFAs-induced lipotoxicity. This notion was corroborated by our *in vivo* investigation using the liver-

specific SIRT3 overexpression mice. As shown in Fig. 3E and Supplementary data, Fig. 11, liver-specific SIRT3 overexpression aggravated liver damage in response to an 8-week palm oil-rich HFD feeding.

SIRT3 Sensitizes Hepatocytes to Lipotoxicity via Negatively Mediating Autophagy.

In keeping with previous studies, including ours,⁽⁹⁾ we showed here that autophagy induction, via several well-established approaches, including rapamycin supplementation, amino acid and serum-depleted medium (EBSS), as well as serum-free medium, provided strong protection against PA-induced cell death (Supplementary data, Fig. 12A & B). Conversely, the autophagy inhibitors, bafilomycin A1 (Baf) and CQ, aggravated PA hepatotoxicity (Supplementary data, Fig. 12C & D). Similar results were observed when SA was used (Supplementary data, Fig. 12E & F). However, OA supplementation with or without autophagy inhibitors had no effect on cell death (Supplementary data, Fig. 12G & H).

To determine whether the negative regulation of SIRT3 on autophagy is involved in PA-induced lipotoxicity, rapamycin was exploited to activate autophagy in SIRT3-overexpressing cells before PA exposure. As shown in Fig. 4A, rapamycin pretreatment blocked PA-induced lipotoxicity in SIRT3-overexpressing cells. Conversely, autophagy suppression via either pharmacological (CQ) or genetic (siRNA for Atg5, Supplementary data, Fig. 13) approach blunted the protective effect of SIRT3 siRNA transfection on PA-induced cell death (Fig. 4B & C; Supplementary data, Fig. 14). These results suggest that SIRT3 controls the sensitivity of hepatocytes to PA-induced lipotoxicity via negatively regulating autophagy.

SIRT3 Negatively Regulates Autophagy via Suppressing AMPK.

AMPK activation induces autophagy by suppressing mammalian target of rapamycin C1 (mTORC1).⁽²⁸⁾ To determine the role of AMPK in SIRT3-regulated autophagy suppression, we

first examined the effect of SIRT3 on AMPK activation. As shown in Fig. 5A, SIRT3 overexpression was associated with suppressed AMPK activation, evidenced by decreased protein phosphorylation of AMPK and ACC, an AMPK target protein, as well as increased phospho-p70S6k1, a well-known downstream target of mTORC1 activation. Moreover, SIRT3 overexpression blunted AICAR-instigated AMPK activation (Supplementary data, Fig. 15). Conversely, SIRT3 siRNA silencing led to a robust enhancement of AMPK activation and p70S6k1 inhibition (Fig. 5A; Supplementary data, Fig. 16). Accordingly, the livers from SIRT3^{-/-} mice demonstrated upregulated p-AMPK and p-ACC levels, as well as decreased phospho-p70S6k1 abundance in comparison to wild type mice (Fig. 5B; Supplementary data, Fig. 17A), while the opposite changes of these proteins were observed in the livers from hepatocyte-specific SIRT3 overexpressing mice (Fig. 5C).

Inasmuch as the existence of a reverse association between SIRT3 and AMPK activity, we next sought to determine if AMPK is mechanistically implicated in the effect of SIRT3 on autophagy. AMPK siRNA silencing prevented LC3-II conversion and LC3 puncta accumulation resulting from SIRT3 suppression (Fig. 5D), indicating that AMPK is the main downstream target of SIRT3 in regulating autophagy. Although both ERK1/2 and Akt were reported to regulate autophagy and be potentially impacted by SIRT3 activation,⁽²⁹⁾ their involvement was not supported by our investigations (Supplementary data, Fig. 17).

PA inhibits AMPK activation in hepatocytes.⁽³⁰⁾ In accordance, we observed that PA strongly inhibited AMPK activation both *in vitro* and *in vivo*, whereas OA activated AMPK (Supplementary data, Fig. 18). Importantly, we showed that silencing AMPK blunted the protection against PA-induced cell death in SIRT3 siRNA- transfected HepG2 cells (Fig. 5E; Supplementary data, Fig. 19). In contrast, either EBSS (amino acid and serum-depleted) or GP

starvation (glucose, L-glutamine, pyruvate, and serum-depleted) medium protected against PA-induced cell death in both control and SIRT3-overexpressing cells, both being associated with the strong activation of AMPK and autophagy (Fig. 5F; Supplementary data, Fig. 20 & 21). These results altogether suggest that AMPK activation contributes to SIRT3-regulated autophagy and sensitivity to PA-induced lipotoxicity.

Altered Intracellular Redox Status Contributes to the Regulatory Role of SIRT3 on AMPK.

Other than altered AMP/ATP ratio, intracellular redox status modulates AMPK activation.⁽³¹⁾ To gain insight on mechanism(s) involved in SIRT3-regulated AMPK activation, we first examined the effect of SIRT3 on intracellular redox status. As shown in Fig. 6A, SIRT3 overexpression markedly decreased intracellular $O_2^{\bullet-}$ levels, while intracellular H_2O_2 content was significantly increased. In contrast, siRNA silencing of SIRT3 significantly enhanced intracellular $O_2^{\bullet-}$ levels but reduced H_2O_2 content (Fig. 6B; Supplementary data, Fig. 22). These observations directed us to subsequently analyze the effect of $O_2^{\bullet-}$ on AMPK activity and autophagy by treating HepG2 with rotenone, a well-established $O_2^{\bullet-}$ inducer that acts via inhibition of mitochondrial complex I. As expected, rotenone exposure led to a significant increase of intracellular $O_2^{\bullet-}$ production (Supplementary data, Fig. 23A). Interestingly, cellular p-AMPK level was also markedly increased in response to rotenone exposure (Supplementary data, Fig. 23B). Importantly, both rotenone-induced $O_2^{\bullet-}$ overproduction and AMPK activation were associated with autophagy induction, as shown by increased LC3-II conversion and LC3 puncta accumulation (Supplementary data, Fig. 23B & C). Reducing intracellular $O_2^{\bullet-}$ by N-acetylcysteine (NAC), a $O_2^{\bullet-}$ cleaner (Supplementary data, Fig. 24A), markedly reduced p-AMPK abundance, and inhibited autophagy (Supplementary data, Fig. 24B & C). Furthermore, we showed that AMPK siRNA silencing abolished rotenone-triggered LC3-II conversion, while AICAR-induced LC3-II

conversion was not affected by NAC supplementation (Fig. 6C). These data altogether suggest that decreased $O_2^{\cdot-}$ generation contributes to SIRT3 overexpression-induced AMPK inhibition and resultant autophagy suppression in hepatocytes.

Our data support that SIRT3 overexpression-induced autophagy suppression involves AMPK inhibition via depleting cellular $O_2^{\cdot-}$ levels. We next asked whether the alteration of intracellular redox status is able to affect PA-induced lipotoxicity. As shown in Supplementary data, Fig. 25A, increasing intracellular $O_2^{\cdot-}$ by rotenone markedly alleviated PA-induced cell death, which was abolished by both siRNA silencing of AMPK and autophagy inhibition (Fig. 6D; Supplementary data, Fig. 25B). In addition, rotenone also alleviated PA-induced cell death in SIRT3 overexpressing cells (Fig. 6E). Conversely, depleting cellular $O_2^{\cdot-}$ content by NAC aggravated PA-induced lipotoxicity (Supplementary data, Fig. 26). Accordingly, NAC blunted the protection conferred by SIRT3 siRNA silencing against PA-induced cell death (Fig. 6F).

SIRT3 Mediated Intracellular $O_2^{\cdot-}$ Homeostasis via Regulating Manganese Superoxide Dismutase (MnSOD) Activity.

SIRT3-mediated deacetylation reaction enhances MnSOD activity,⁽²⁰⁾ a mitochondrial antioxidant enzyme catalyzing the conversion of $O_2^{\cdot-}$ to H_2O_2 . In line with previous reports, our result showed that whereas SIRT3 overexpression robustly decreased acetylated MnSOD protein abundance, SIRT3 silencing by siRNA led to increased MnSOD protein acetylation (Fig. 7A & B). Of note, treatment of HepG2 cells with 2-methoxyestradiol (2-ME), a commonly used MnSOD inhibitor, mimicked the effect of SIRT3 siRNA knockdown on intracellular ROS production (Fig. 7C; Supplementary data, Fig. 27A & B), whereas MnSOD overexpression in HepG2 cells mimicked the effect of SIRT3 overexpression (Fig. 7D). Correspondingly, 2-ME prevented PA-induced cell death in both normal and SIRT3-overexpressing cells (Fig. 7E;

Supplementary data, Fig. 27C), which, as expected, was blunted by autophagy suppression (Supplementary data, Fig. 27D), while MnSOD overexpression suppressed autophagy and aggravated PA-induced cell death (Fig. 7F; Supplementary data, Fig. 28 & 29).

Palmitate Exposure is Associated with MnSOD Deacetylation and Reduced $O_2^{\cdot-}$ Generation.

Finally, the effects of PA exposure on MnSOD protein acetylation status and intracellular $O_2^{\cdot-}$ homeostasis were determined. In harmony with its positive effect on SIRT3, a 12-hour PA exposure (0.5 mM) evidently decreased MnSOD protein acetylation (Fig. 8A). In accordance, PA exposure reduced intracellular $O_2^{\cdot-}$ levels and attenuated rotenone-triggered intracellular $O_2^{\cdot-}$ increase (Fig. 8 B & C).

Discussion

Lipotoxicity plays a pathological role in the development of NAFLD.⁽³²⁾ However, the exact mechanism(s) underlying SFAs-triggered hepatocyte cell death remains to be fully elucidated. In the present study, we provide evidence for a novel SIRT3-dependent mechanism exacerbating SFAs-induced lipotoxicity in hepatocytes. We demonstrate that hepatic SIRT3 overexpression renders hepatocytes more susceptible to SFAs-induced cell death and aggravates NAFLD progression in mice in the setting of SFAs-rich HFD feeding through negatively regulating autophagy induction. Our results reveal that AMPK inhibition, resulting from MnSOD activation-derived $O_2^{\cdot-}$ depletion, is mechanistically involved in SIRT3-induced autophagy suppression and aggravated lipotoxicity in hepatocytes (see proposed model in Fig. 8D).

SIRT3 plays a pivotal role in maintaining mitochondrial protein acylation homeostasis, and thereby metabolism, by deacetylating critical enzymes in a variety of metabolic pathways.^(16-21, 25) SIRT3 activation is one of the major mechanisms accounting for many calorie restriction-derived benefits.^(20, 33) Similarly, the beneficial effects of SIRT3 activation have been reported by numerous studies in a variety of experimental and clinical settings.^(34, 35) In this work, we demonstrate that SFAs and unsaturated FAs differentially regulate SIRT3 expression in hepatocytes. Unexpectedly, our data suggest that SIRT3 activation is associated with increased susceptibility of hepatocytes to PA-induced cell death. These observations are in direct contrast with a previous report that HFD feeding downregulated hepatic SIRT3 expression and that SIRT3 activation conferred beneficial effect on PA-induced hepatotoxicity by preventing oxidative stress derived from mitochondrial dysfunction.⁽³⁶⁾ These discrepancies are difficult to be fully clarified at present. However, one possible explanation can be the difference in the SIRT3 overexpression protocol employed. In comparison to the transient SIRT3 overexpression

used by Bao et al.,⁽³⁶⁾ our stable overexpression system mimics *in vivo* experimental conditions and is therefore more clinically relevant. Also noteworthy is that, although oxidative stress has been mechanistically implicated in PA-induced lipotoxicity in several cell types, including hepatocytes, our previous study documented that PA exposure was associated with increased intracellular glutathione concentration in HepG2 cells,⁽³⁷⁾ suggestive of the existence of oxidative stress-independent mechanism(s) in PA-induced hepatotoxicity. Furthermore, exacerbated PA-induced hepatocyte cell death by the antioxidant NAC implied that a certain amount of ROS could in fact be required to protect against lipotoxicity.⁽³⁷⁾

Our studies point to a critical role of SIRT3 as a negative regulator of autophagy in hepatocytes, leading to exacerbated SFAs-induced lipotoxicity. Our data revealed that SIRT3 gene silencing/knockout is associated with AMPK activation in hepatocyte/the liver. These observations are consistent with a previous report which demonstrated that the reduced SIRT3 expression represents a fundamental mechanism in metformin-induced AMPK activation in hepatocytes.⁽³⁸⁾ Conversely, SIRT3 overexpression in HepG2 cells led to a significant inhibition of AMPK activation. This observation disagree with a previous report showing that SIRT3 overexpression in HepG2 cells was associated with AMPK activation.⁽³⁹⁾ The transient SIRT3 overexpression system employed in that study might potentially account for the discrepancy. AMPK activation induces autophagy by inhibiting mTORC1, a strong inhibitor of autophagy.⁽²⁸⁾ Given the well-documented positive role of AMPK activity in the regulation of autophagy, we therefore postulate that AMPK may represent a mechanistic link between SIRT3 and autophagy. This notion is supported by our data that genetic inhibition of AMPK alleviated autophagy induction in SIRT3 siRNA-transfected hepatocytes and abolished SIRT3 silencing-conferred protection against lipotoxicity, while activating AMPK significantly reduced PA-induced cell

death in SIRT3-overexpressed hepatocytes. Nevertheless, under regular diet, mice with liver AAV-SIRT3 OE intriguingly failed to recapitulate the effects of SIRT3 on AMPK and mTORC1 as observed in SIRT3 OE hepatocytes, in spite of apparent autophagy induction. This observation implied the existence of the AMPK/mTORC1 signaling pathway-independent mechanism in the regulation of SIRT3 OE-instigated autophagy suppression in the *in vivo* system. On the other hand, we observed that, under high palm oil-HFD feeding, AAV-SIRT3 OE mice manifested suppressed hepatic AMPK activation and increased mTORC1 activation (p-S6K1) in comparison to that in AAV-null mice (Fig. 5C), suggesting that the effect of SIRT3 on AMPK/mTOR signaling pathway is context-dependent, which may require environmental/dietary cues.

AMPK is a positive regulator of autophagy. It was previously reported that SIRT3^{-/-} mice manifested a significantly reduced hepatic ATP level in comparison to wild type animals.⁽¹⁶⁾ Moreover, AMPK activation by metformin was associated with SIRT3 downregulation and the subsequent reduction of cellular ATP levels in both hepatocytes and the liver, which was recovered by SIRT3 overexpression.⁽³⁹⁾ These observations provided the first line of evidence supporting the existence of a negative relationship between SIRT3 and AMPK. Other than altered AMP/ATP ratio, it was also reported that O₂^{•-} plays an important role in the activation of AMPK and autophagy.^(31, 40) To gain further insight on the mechanism(s) involved in SIRT3-triggered AMPK suppression, we investigate the effect of SIRT3 on ROS production and their role in AMPK activation. Our studies provide initial evidence suggesting that O₂^{•-} depletion plays a mechanistic role in SIRT3 overexpression-induced AMPK suppression. This notion is supported by several findings: 1. SIRT3-overexpressing HepG2 cells have significantly low levels of O₂^{•-}, whereas hydrogen peroxide was significantly increased. Conversely, SIRT3 gene

silencing was associated with markedly increased $O_2^{\bullet-}$ levels, while hydrogen peroxide levels were significantly reduced; 2. Rotenone, a potent $O_2^{\bullet-}$ inducer via suppressing mitochondrial complex I, increased cellular $O_2^{\bullet-}$ production, activated AMPK, induced autophagy, and protected HepG2 cells against lipotoxicity; 3. NAC, a potent ROS scavenger, inhibited AMPK and autophagy activation, and potentiated PA-induced hepatotoxicity; 4. MnSOD negatively regulates $O_2^{\bullet-}$ production and AMPK activation. Overall, our studies support the notion that SIRT3-mediated $O_2^{\bullet-}$ depletion mechanistically links SIRT3 activation to AMPK suppression.

MnSOD, the mitochondrial form of SOD converting $O_2^{\bullet-}$ into H_2O_2 , is crucial for the maintenance of mitochondria-generated $O_2^{\bullet-}$ homeostasis. Interestingly, MnSOD is a target of SIRT3, whose deacetylation by SIRT3 results in enhanced enzymatic activity.⁽²⁰⁾ In line with previous reports, we found that SIRT3 overexpression robustly decreased MnSOD protein acetylation, whereas its silencing by siRNA led to increased acetylated MnSOD. Moreover, we found that MnSOD inhibition mimics SIRT3 gene silencing in terms of the ROS generation pattern, whereas MnSOD overexpression mimics SIRT3 overexpression phenotype. Importantly, similar to SIRT3 siRNA transfection, MnSOD inhibition prevented PA-induced cell death in both normal and SIRT3-overexpressing HepG2 cells. These data suggest that SIRT3 mediates mitochondria-derived $O_2^{\bullet-}$ protection via affecting MnSOD acetylation status.

The negative effect of palmitate on AMPK was previously reported in several cell types, including hepatocytes, the exact mechanism(s) remain unclear. The ceramide-dependent PP2A activation seemed to play an important role in palmitate-induced AMPK dephosphorylation/suppression.^(41, 42) Moreover, palmitate-induced PPAR- α suppression was also reported to contribute to AMPK inhibition.⁽⁴³⁾ The cell culture experiments in the present study provided strong evidence that SIRT3 upregulation is mechanistically involved in

palmitate-elicited AMPK suppression. At the same time, we also recognized that our *in vitro* observations were not able to be completely recapitulated in our *in vivo* investigations. Specifically, our data showed that long-term high palm oil-HFD feeding exerted much profound effect on AMPK/mTORC1 signaling pathway than AAV-SIRT3 overexpression did. Therefore, the role of SIRT3-independent pathways in the regulation of palmitate-induced AMPK inhibition cannot be excluded and warrant further investigations.

In summary, this study provides initial experimental evidence that SFAs exposure of hepatocytes leads to SIRT3 upregulation and activation both *in vitro* and *in vivo*, which in turn exacerbates SFAs-triggered hepatotoxicity by negatively regulating autophagy induction. Intriguingly, our studies uncover that AMPK activation due to MnSOD activation-dependent $O_2^{\bullet -}$ depletion plays a mechanistic role in SIRT3-mediated autophagy suppression in hepatocytes. Given that the both muscle- and liver-specific SIRT3 knockout mice do not manifest any overt metabolic phenotype under either chow or high-fat diet conditions⁽⁴⁴⁾ and that SIRT3 knockout mice are protected from acetaminophen-induced hepatotoxicity,⁽⁴⁵⁾ our findings suggest that curbing SIRT3 overactivation can be a potential therapeutic choice for the treatment of NAFLD as well as other metabolic disorders, with lipotoxicity being the principal pathomechanism.

Conflicts of Interest

None.

Accepted Article

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FIGURE LEGENDS

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Mitochondrial protein acetylation. (D) SIRT3 protein expression. (E) Mitochondrial protein acetylation. (F) SIRT3 protein expression. Each *in vitro* test was performed at least three times ($n = 8$ for mice), and a representative blot was shown. All values are denoted as means \pm SD. Bars with different characters differ significantly, $p < 0.05$.

Figure 3. SIRT3-overexpressing increases the susceptibility of PA/palm oil-induced injury in HepG2 cells and liver. Vector control or SIRT3 OE HepG2 cells were suffered to PA (0.5 mM) exposure for 12 h. For SIRT3 inhibition, HepG2 cells were either transfected siSIRT3 or pretreated with NAM (5 mM), a pharmacologic inhibitor of SIRT3 for 1 h. Cell death was detected by the measurements of LDH release and PI staining. Liver specific SIRT3 OE mice model was established and fed with normal or high-fat diet using either palm oil or corn oil as described in the Methods. Plasma and liver were collected for analysis ($n = 8$). (A)-(D) Cell death was detected by the measurements of PI staining or LDH release after the indicated treatments. (E) Liver H&E stain (I); Plasma ALT (II). All values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.

Figure 4. Autophagy alteration contributes to SIRT3-regulated lipotoxicity in HepG2 cells. (A) SIRT3 OE and vector control HepG2 cells were treated with 0.5 mM PA for 12 h with or without 1 h pre-incubation with 50 nM rapamycin. Cell death was detected by the measurements of LDH release and PI staining, respectively. (B) HepG2 cells were transfected with either scramble siRNA or SIRT3 siRNA for overnight, followed with 0.5 mM PA exposure for 12 h. Autophagy inhibitor CQ (20 μ M) was added 1 h ahead of PA treatment. Cell death was detected. (C) HepG2

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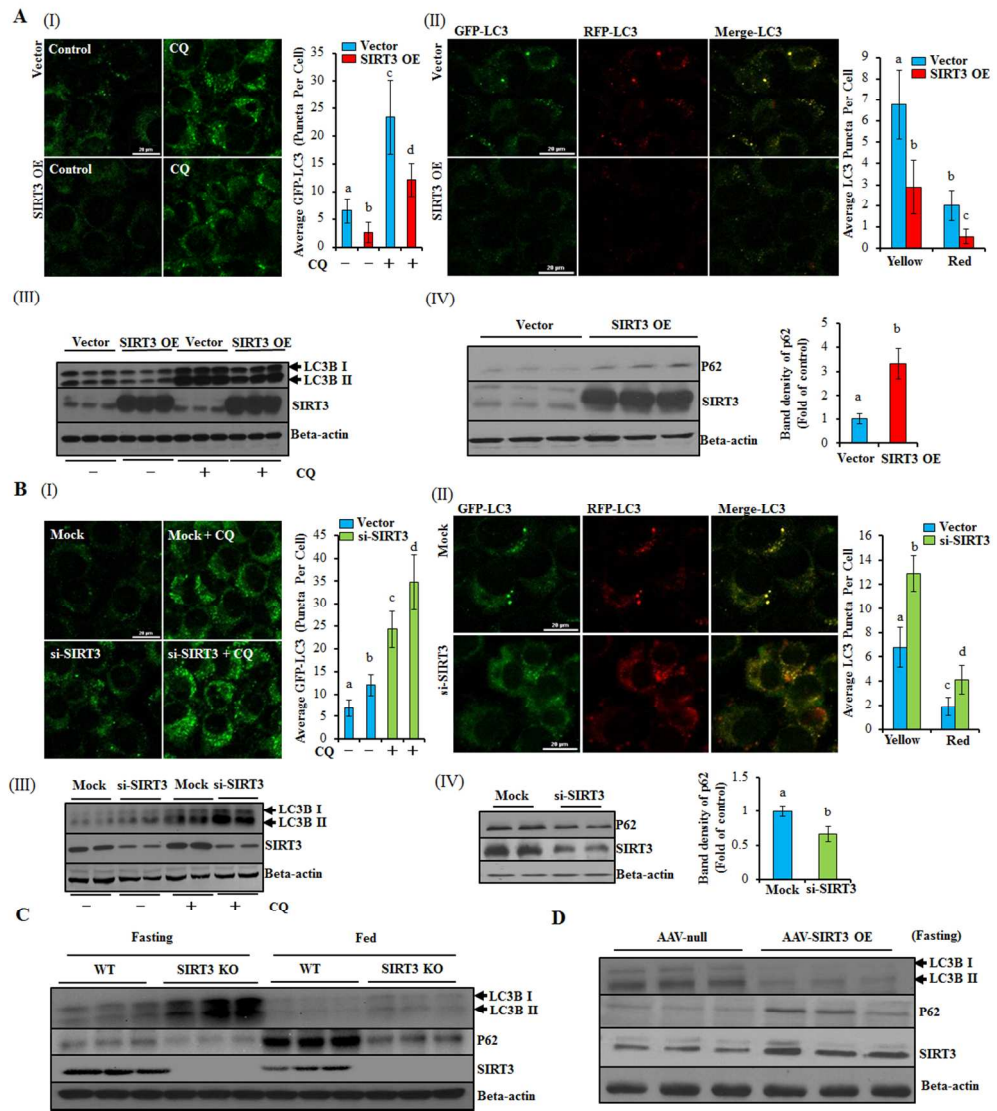


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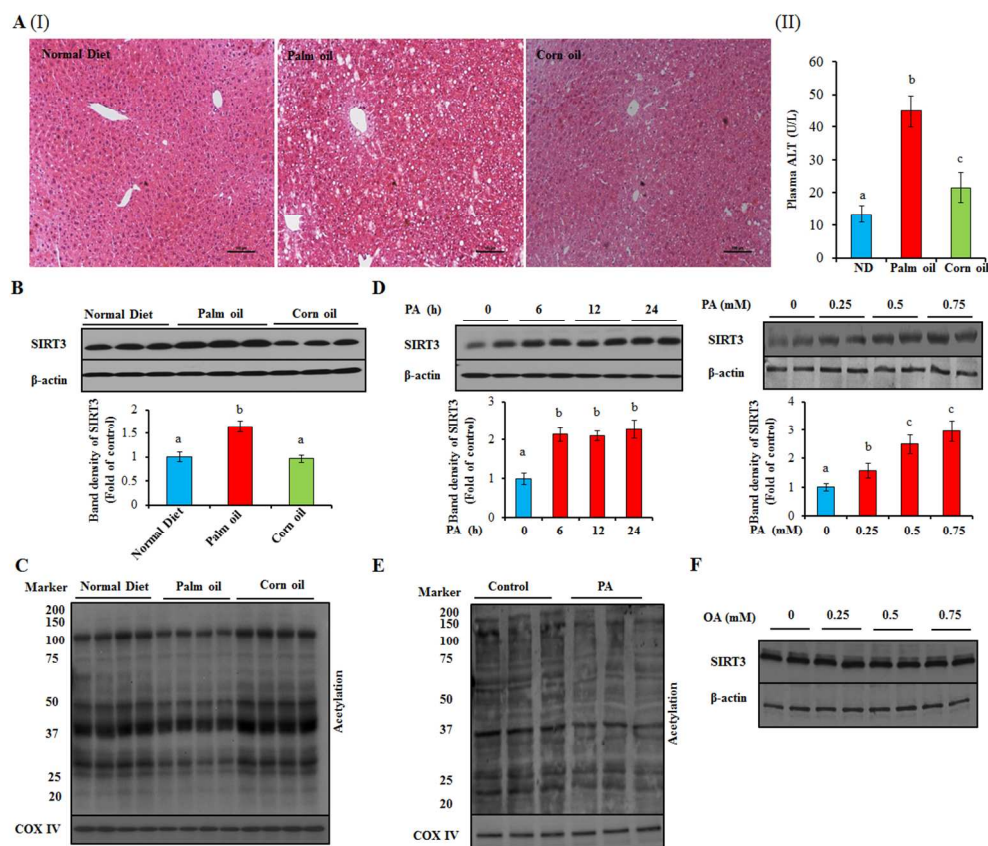


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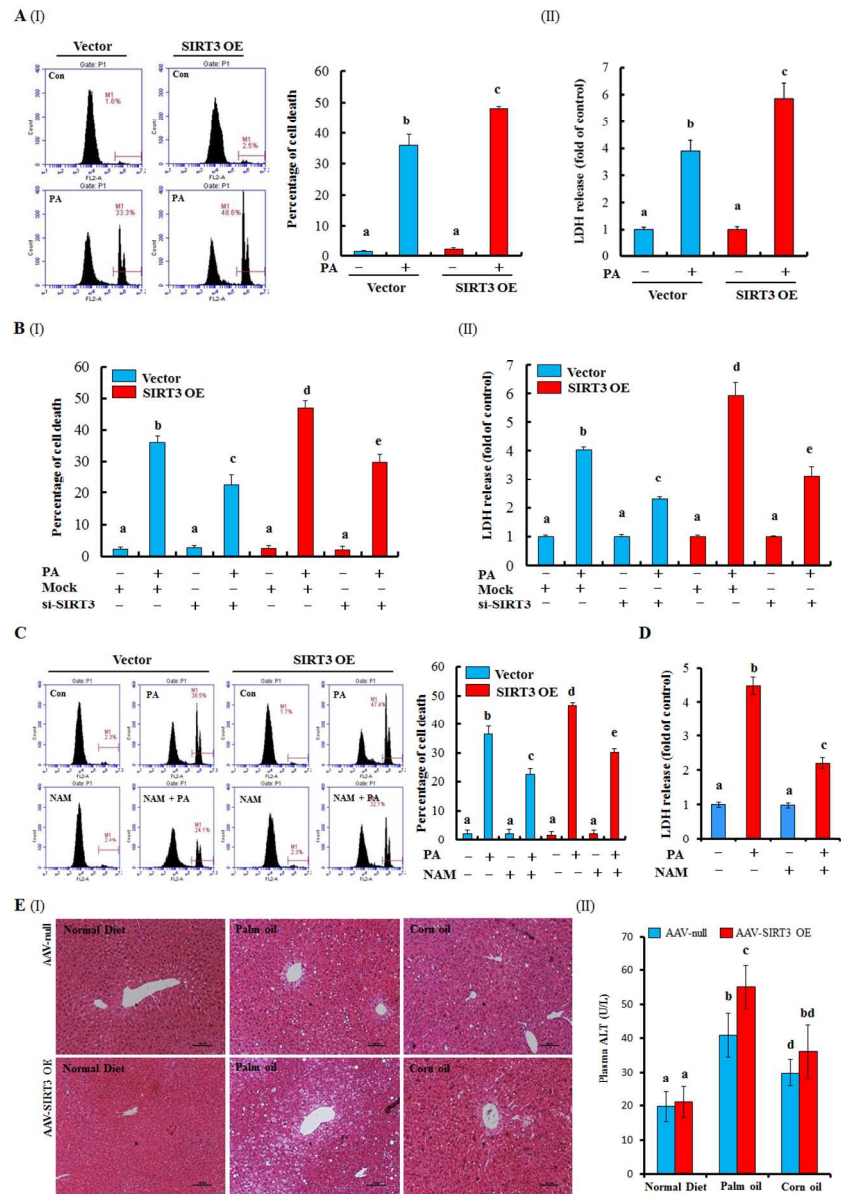


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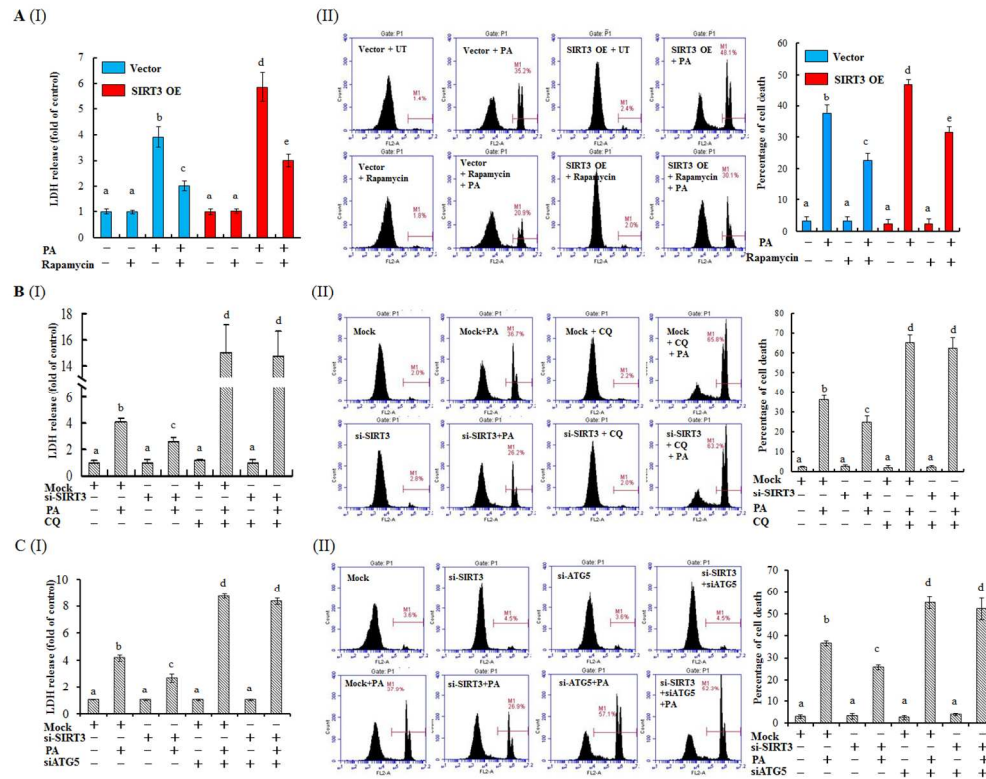


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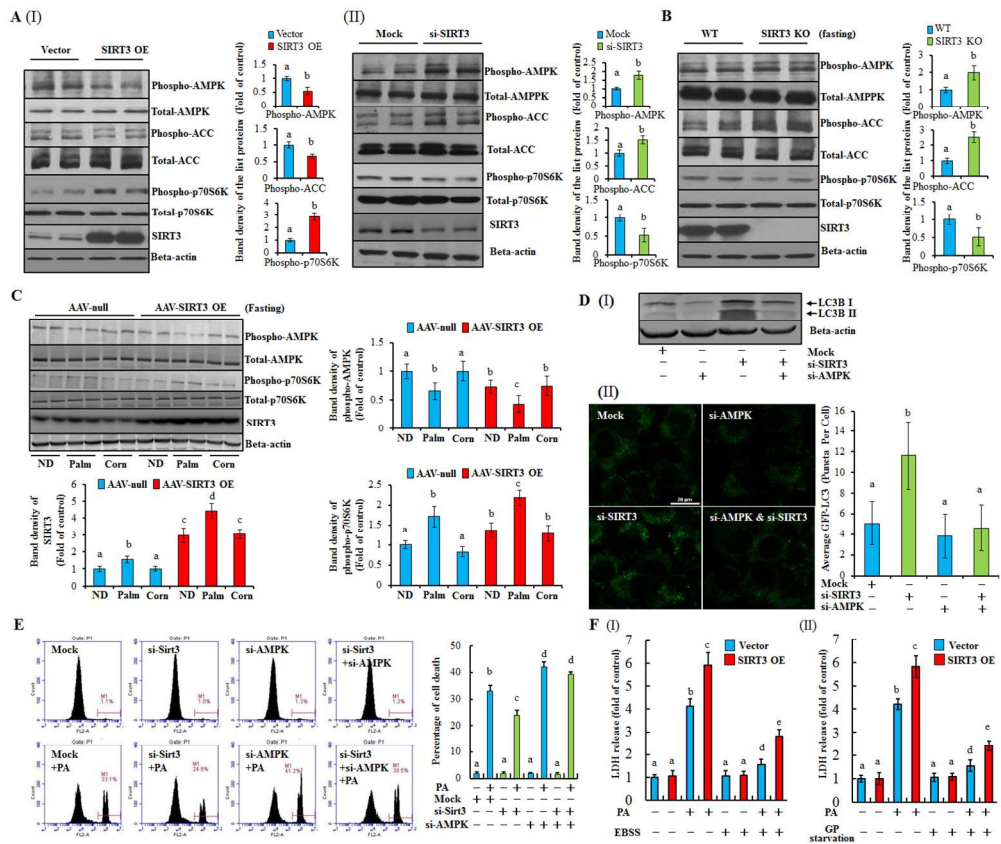


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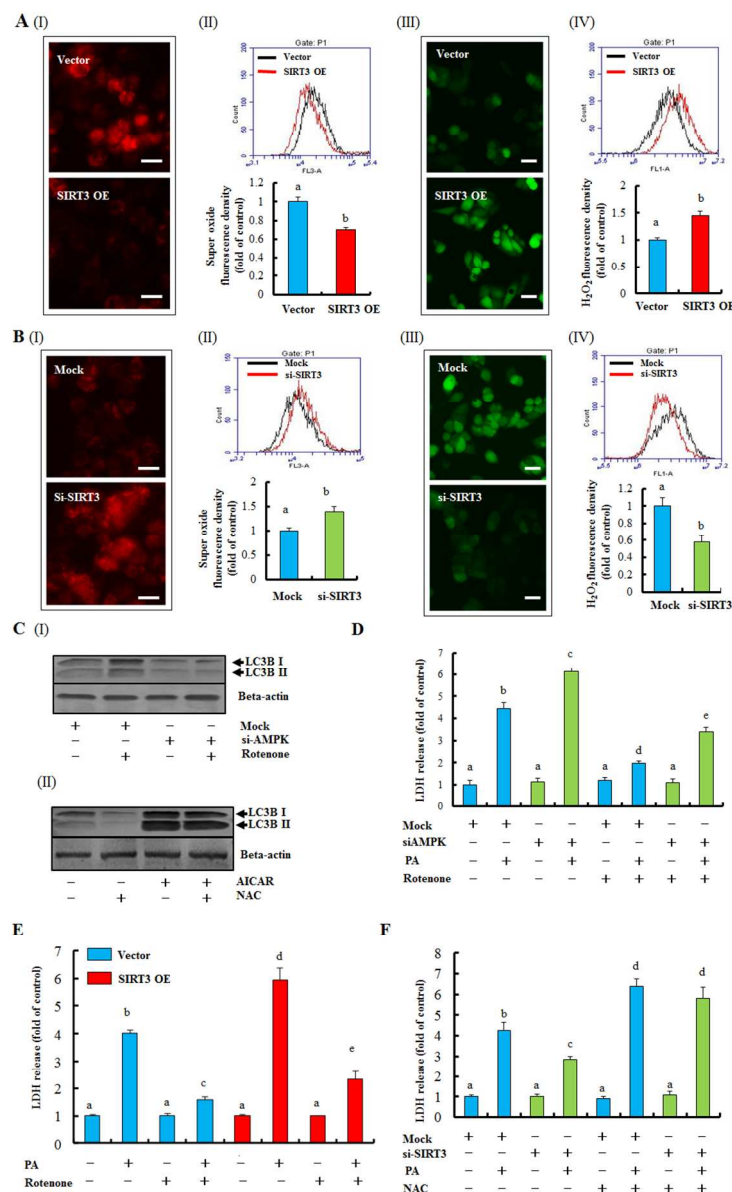


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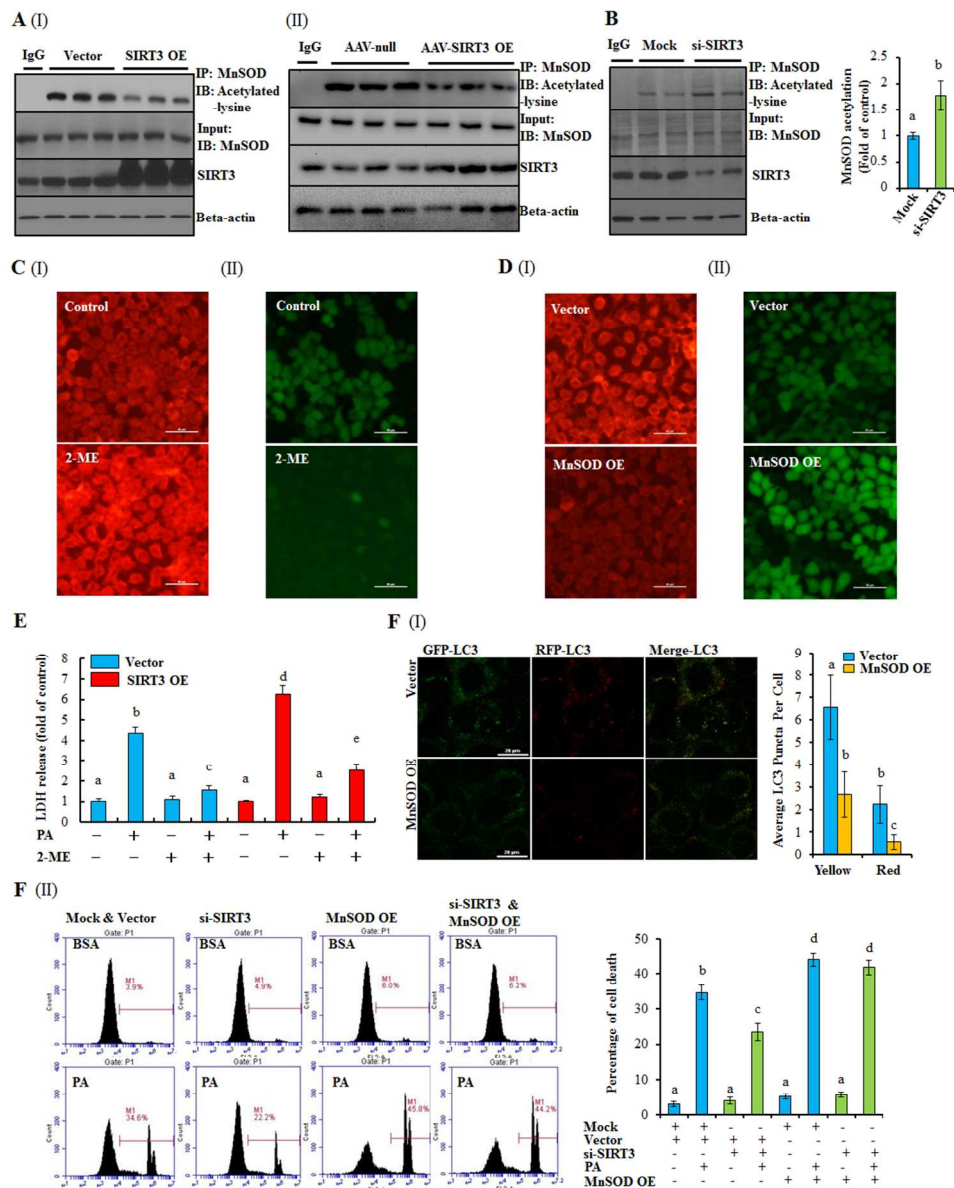


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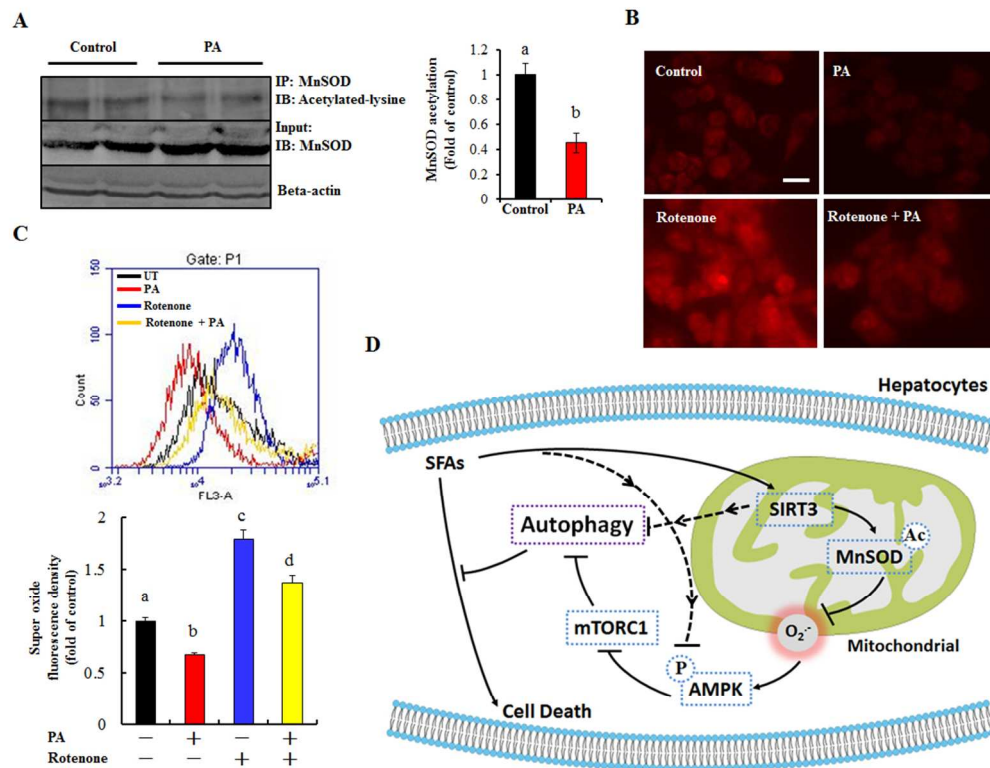


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Supplementary Material**SIRT3 Acts As a Negative Regulator of Autophagy Dictating Hepatocyte****Susceptibility to Lipotoxicity**

Songtao Li^{1,2,†}, Xiaobing Dou^{1,3,†}, Hua Ning^{2,5}, Qing Song², Wei Wei², Ximei Zhang¹,
Chen Shen¹, Jiabin Li¹, Changhao Sun^{2,5}, Zhenyuan Song^{1,3,4}

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Supplementary Methods

Animal Model and Experimental Protocol

All animal studies were performed in accordance with the animal care committee at the University of Illinois at Chicago. All mice were fed *ad libitum*, and housed on a 12:12 h light: dark cycle at 25°C. Male SIRT3^{-/-} mice and WT 129Sv mice (8-week of age) were purchased from Jackson Laboratory (Bar Harbor, ME), and maintained on a standard chow diet. Male C57BL/6 mice (8-week old, Jackson Laboratory, Bar Harbor, ME) were randomly assigned to normal diet, palm oil high-fat diet, and corn oil high-fat diet (Supplementary data Table 1). The liver-specific SIRT3 overexpression mice and corresponding control mice were generated via the injection of recombinant adeno-associated viral (AAV) serotype 8 gene transfer vectors containing a liver-specific promoter combination (albumin promoter) with either mouse SIRT3 sequence or empty as vector control (Cyagen Biosciences Inc., Guangzhou, China). Finally, there are 6 groups in total (n = 8 mice/group) and the animals were fed for 8 weeks. AAV8 vectors were administered by tail vein injection at a dose of 1×10^{12} viral titer/mL in a total volume of 100 μ L/mice at the beginning of the experiment. Recombinant AAV8 vectors encoding mouse SIRT3 (NM_022433.2) under the control of albumin promoter (a liver-specific promoter) were generated by Cyagen Biosciences Inc. (Guangzhou, China). A noncoding plasmid carrying only the albumin promoter was used to produce vector control particles. The overexpression efficiency was shown in Fig. 1D and Supplementary data, Fig. 30. Mice were sacrificed after 8-week of feeding. Plasma and liver were collected for analysis. Plasma alanine aminotransferase (ALT) content was determined by a commercial ALT kit (Thermo Fisher, Waltham, MA). Triglyceride was assayed using a

Triglycerides Assay Kit (Thermo Fisher, Waltham, MA). Liver redox state was detected using a commercial TBARS Kit (Nanjing Jiancheng Bioengineering Institute, Nanjing, China). Small pieces of liver were fixed immediately in 10% buffered formalin. After paraffin embedding, 5 μ m sections were deparaffinized in xylene and were rehydrated through a series of decreasing concentrations of ethanol. Sections were stained with hematoxylin and eosin (H&E).

Chemicals

All chemicals were purchased from Sigma-Aldrich (Sigma-Aldrich, St. Louis, MO), unless otherwise specified. Actinomycin D was obtained from Solarbio (Beijing, China). PA- and SA-BSA conjugates were prepared as described previously ⁽¹⁾. Briefly, PA or SA was dissolved in ethanol and saponified with sodium hydroxide. The sodium salt was dried, re-suspended in saline and heated at 80°C until completely dissolved. While the solution was still warm, isovolumetric 20% (w/v) BSA was added and the mixture was stirred at 50°C for 4 h to allow PA or SA to bind to BSA. The PA- or SA-BSA complex (3 mmol/L fatty acid: 1.5 mmol/L BSA; molar ratio, 2:1) was then sterilized by filtering, and aliquoted for future use. In all the experiments, the control group was exposed to an equal amount of solvent (e.g. BSA, ethanol, DMSO).

Cell Culture

The HepG2 cell line was obtained from the American Type Culture Collection (Manassas, VA). Cells were cultured in Dulbecco's Modified Eagle Medium (DMEM) containing 10% (v/v) fetal bovine serum, 100 U /ml penicillin, and 100 μ g /ml streptomycin at 37°C in a humidified O₂/CO₂ (19:1) atmosphere.

Alpha mouse liver (AML)-12 hepatocyte culture was established from a mouse transgenic for human transforming growth factor α , and was obtained from the American Type Culture Collection (ATCC, CRL-2254), and was cultured in Dulbecco's Modified Eagle Medium /Ham's Nutrient Mixture F-12, 1:1 (DMEM /F-12, Sigma-Aldrich, 051M8322) containing 10% (v/v) fetal bovine serum (PAA Laboratories, A15-701), 5 mg/ml insulin (Sigma-Aldrich, I9278), 5 μ g/ml transferrin (Sigma-Aldrich, T8158), 5 ng/ml selenium (Sigma-Aldrich, 229865), 40 ng/ml dexamethasone (Sigma-Aldrich, D4902), 100 U/ml penicillin, and 100 μ g/ml streptomycin (Life technologies, 15140-122) at 37°C in a humidified O₂/CO₂ (95:5) atmosphere.

Establishment of stable human SIRT3 over-expression cell lines

The eukaryotic expression vector plasmid pcDNA3.1⁺/hSIRT3 containing the human SIRT3 gene sequence was kindly provided by Dr. Yu Wang (University of Hong Kong) ⁽²⁾. HepG2 cells grown to 80–90% confluence were transfected with 0.8 μ g/well of the expression vector pcDNA3.1⁺/hSIRT3 or empty vector control pcDNA3.1⁺ (Invitrogen, Grand Island, NY) in 24-well plate using Lipofectamine 2000 reagent (Invitrogen, Grand Island, NY) following the manufacturer's guidelines. After 48 hours transfection, cells were trypsinized into 24-well plates at a ratio of 1:100 for monoclonal cell selection using 800 μ g/ml G418. The stable cell line was established and confirmed by both western blot analysis and Real Time PCR. SIRT3 over-expression cells and vector control cells were cultured in DMEM containing 10% (v/v) fetal bovine serum, 400 μ g/ml G418, 100 U/ml penicillin, and 100 μ g/ml streptomycin at 37°C in a humidified O₂/CO₂ (19:1) atmosphere.

MnSOD Overexpression

Recombinant lentivirus vector (pLV[Exp]-EGFP:T2A:Neo) containing human MnSOD (NM_001024465.1) under the control of EF-1alpha promoter was generated by Cyagen Biosciences Inc. (Guangzhou, China). A noncoding vector carrying the EF1alpha promoter was used to produce vector control particles. The optimal virus titer used for cell transfection was screened according to the manufacturer's instructions. HepG2 cells were transfected with either pLV[Exp]-EGFP:T2A:Neo-EF1alpha-hMnSOD to overexpress MnSOD or pLV[Exp]-EGFP:T2A:Neo-EF1alpha-null as a vector control.

Cell Death Assays

2×10^5 /ml cells were seeded in 24-well plates and, after the indicated treatments, cell death was determined by measurement of LDH release, propidium iodide (PI) staining, or Hoechst staining. For LDH assay, culture medium was collected and detected using an LDH assay kit (Thermo Scientific Inc, VA) according to the manufacturer's instructions. For PI staining, cells were trypsinized and stained with PI staining solution (BD Pharmingen, CA) according to the manufacturer's instructions. Fluorescence was measured by flow cytometry (Accuri c6, BD, CA). For Hoechst staining, cells were stained with Hoechst staining solution (Beyotime Biotechnology, Nantong, China) according to the manufacturer's instructions and imaged by Nikon eclipse Ti-S fluorescence microscope (Nikon, Tokyo, Japan).

Analysis of GFP-LC3 puncta

The recombinant adenovirus GFP-LC3 (1×10^{10} viral titer/mL) was obtained from Hanbio Biotechnology Co. Ltd. (Shanghai, China). Cells were transiently transfected with recombinant adenovirus GFP-LC3 according to the manufacturer's instruction. Puncta was detected by a laser scanning confocal microscope (Nikon A1R, Japan) from at least 50 cells

for each individual experiment after different treatment.

Analysis of autophagic flux

The autophagic flux was measured according to methods described previously ⁽³⁾. The cells were pretreated with chloroquine (CQ), inhibitor of lysosome acidification, after genetically overexpressing or knocking-down SIRT3. The autophagic flux was determined via detecting GFP-LC3 puncta and LC3-II expression using laser scanning confocal microscope and Western blot, respectively. Additionally, cells were transiently transfected with recombinant adenovirus mRFP-GFP-LC3 (Hanbio Biotechnology Co. Ltd., Shanghai, China). Yellow or red puncta was detected by a laser scanning confocal microscope (Nikon A1R, Japan) from at least 50 cells for each individual experiment after different treatment.

Real Time-PCR

Total RNA isolation, reverse transcription, and real time PCR were performed as described previously ⁽⁴⁾. Briefly, total RNA from cultured cells was isolated with a phenol-chloroform extraction. For each sample, 1 µg total RNA was reverse-transcribed using a high-capacity cDNA reverse transcription kit (Applied Biosystems, Foster City, CA). The cDNA was amplified in MicroAmp Optical 96-well reaction plates with a SYBR Green PCR Master Mix (Applied Biosystems, Foster City, CA) on an Applied Biosystems Prism 7000 sequence detection system. Relative gene expression was calculated after normalization by a house-keeping gene (18S rRNA).

Western-blot Analysis

Western-blot was performed as described previously ⁽⁴⁾ and the following antibodies were used: Anti-Sirt3, anti-phospho-AMPK, anti-AMPK, anti-phospho-Akt, anti-Akt,

anti-phospho-Erk1/2, anti-Erk1/2, anti-phospho-p70S6K, anti-p70S6K, anti-LC3B, anti-p62, anti-phospho-ACC, anti-ACC, anti-MnSOD, anti-acetylation, anti-COX IV, anti-caspase-3, anti-parp-1, anti-Atg5, and anti-beta-actin antibodies (Cell Signaling Technology, Danvers, MA).

Immunoprecipitation

Cells were lysed in an immunoprecipitation (IP) buffer (150 mM NaCl, 50 mM Tris-HCl, 1% Nonidet P-40, pH 7.8, and a mammalian cell-specific protease inhibitor cocktail. Total cellular extracts (200 µg of protein) were incubated with anti-MnSOD antibody (1 µg/ml) in IP buffer overnight at 4 °C on a rocker. The antibody-protein mixture was agitated at 150 rpm with Protein A/G agarose (Santa Cruz Biotechnology) for 1 hr at 4°C. The immunoprecipitates were washed four times with IP buffer. The washed immunoprecipitates were incubated in 50 µl of 1 × electrophoresis loading buffer and heated at 100 °C for 5 min. The beads were spun out and the supernatant was resolved by SDS-PAGE, and the modification of MnSOD by Sirt3 was analyzed by Western blot.

RNA interference

Cultured cells were transfected with human SIRT3 siRNA, human AMPK siRNA, human or Atg5 siRNA (Santa Cruz Biotechnology) using Lipofectamine 2000 according to the manufacturer's instructions. In the control group, cells were transfected with scrambled siRNA (Santa Cruz Biotechnology).

Reactive oxygen species (ROS) detection

The intracellular superoxide ($O_2^{\cdot-}$) and hydrogen peroxide (H_2O_2) were detected as described previously ⁽⁵⁾. Briefly, hepatocytes were stained with DCFH-DA for the

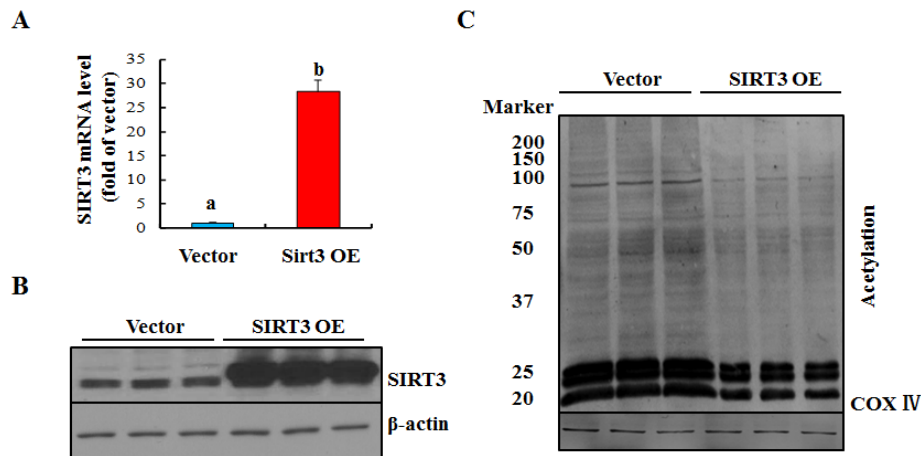
measurement of H_2O_2 after the indicated treatments. Intracellular $\text{O}_2^{\cdot-}$ was detected using a commercial Superoxide Detection kit (Enzo Life Science, NY) according to manufacturer's instructions. The fluorescent signal of listed ROS was collected by fluorescence microscope and flow cytometry.

Supplemental Table

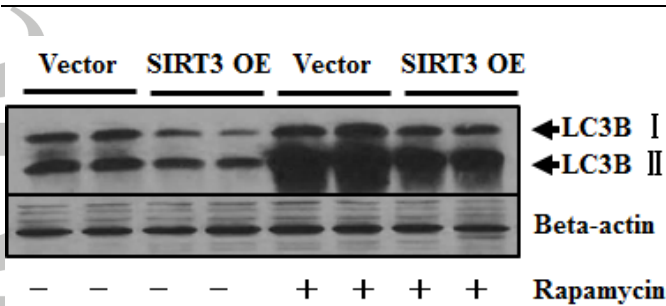
Supplementary Table 1

	Normal diet	High-fat diet	
		Palm oil	Corn oil
Corn Starch (g/kg)	417.5	227.5	227.5
Casein (g/kg)	200	200	200
Dextrinized cornstarch (g/kg)	132	132	132
Sucrose (g/kg)	100	100	100
Palm oil (g/kg)	-	190	-
Corn oil (g/kg)	-	-	190
Soybean oil (g/kg)	50	50	50
Fiber (g/kg)	50	50	50
Mineral mix(AIN-93G-MX, g/kg)	35	35	35
Vitamin mix(AIN-93G-MX, g/kg)	10	10	10
L-cystine (g/kg)	3	3	3
Choline bitartrate (g/kg)	2.5	2.5	2.5
Fat energy (kcal%)	11.7	45.0	45.0
Carbohydrates energy (kcal%)	67.5	38.3	38.3
Protein energy (kcal%)	20.8	16.7	16.7
Total energy (kcal/kg)	3848	4798	4798

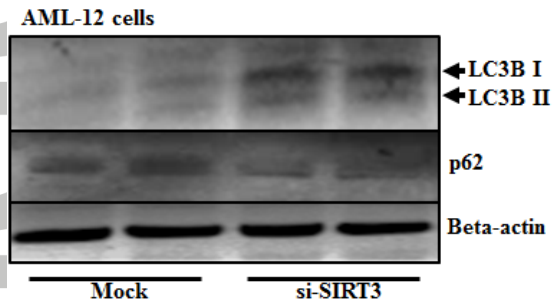
Supplemental Figures



Supplementary Fig. 1 Stable SIRT3-overexpressing (SIRT3 OE) hepatocyte cell line was established in HepG2 cells. (A) SIRT3 mRNA level. (B) SIRT3 protein level. (C) SIRT3 activity was detected by the assay of acetylated proteins abundance in the mitochondrial fractions. All values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.

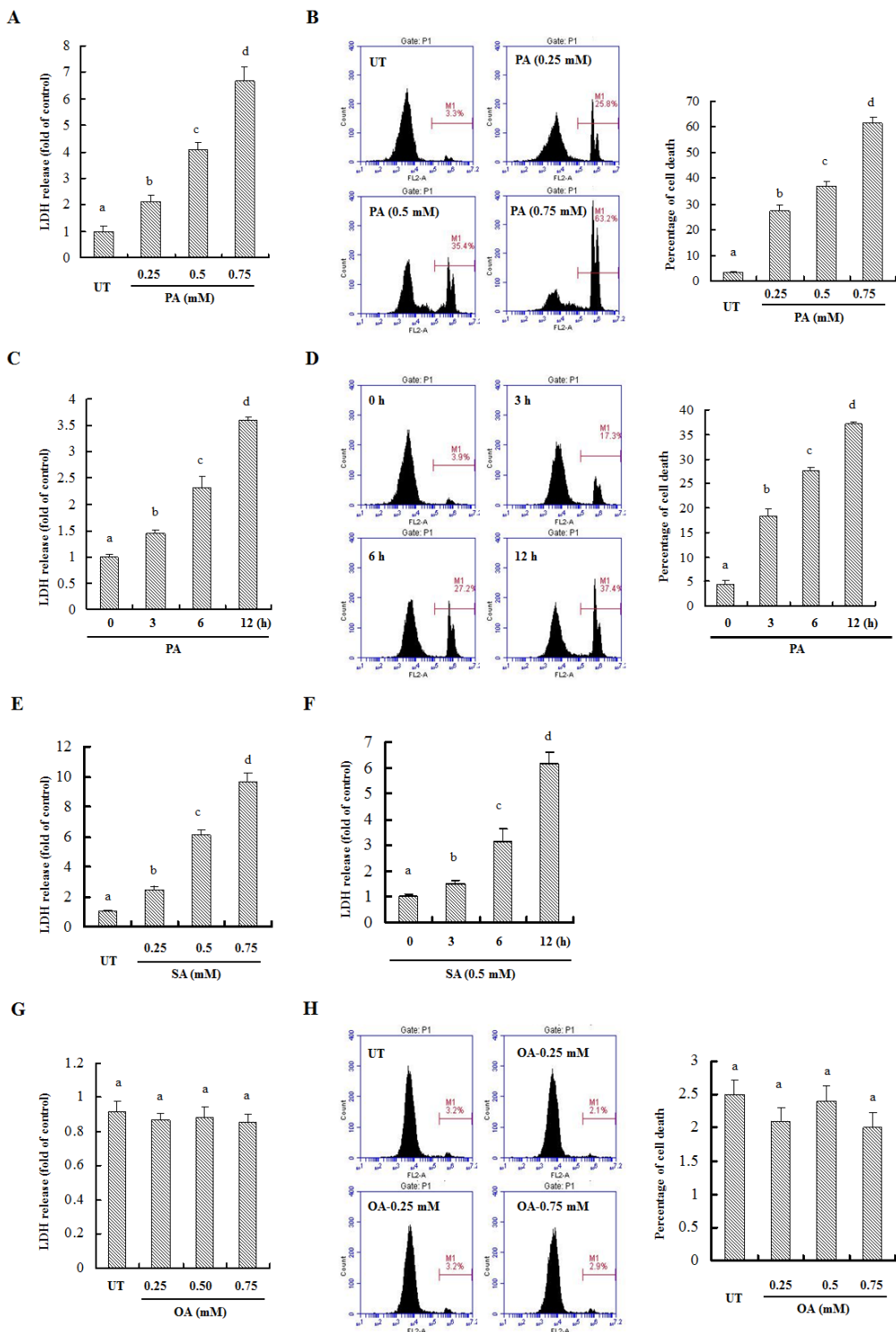


Supplementary Fig. 2 SIRT3-overexpressing inhibites LC3-II formation with or without autophagy activation. SIRT3 OE and vector control HepG2 cells were treated with rapamycin for 12 h. LC3-II level was detected by Western-blotting.



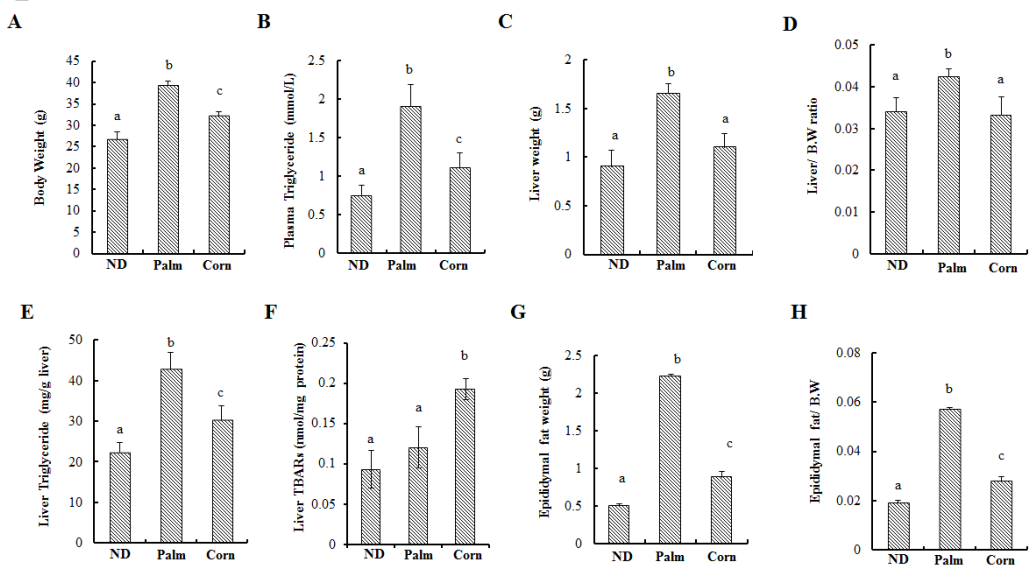
Supplementary Fig. 3 Knocking-down SIRT3 stimulates autophagy in mouse hepatocytes. AML-12 mouse hepatocytes were transfected with siSIRT3 or scramble siRNA. Total lysates were subjected to immunoblotting assay for LC3 and p62.

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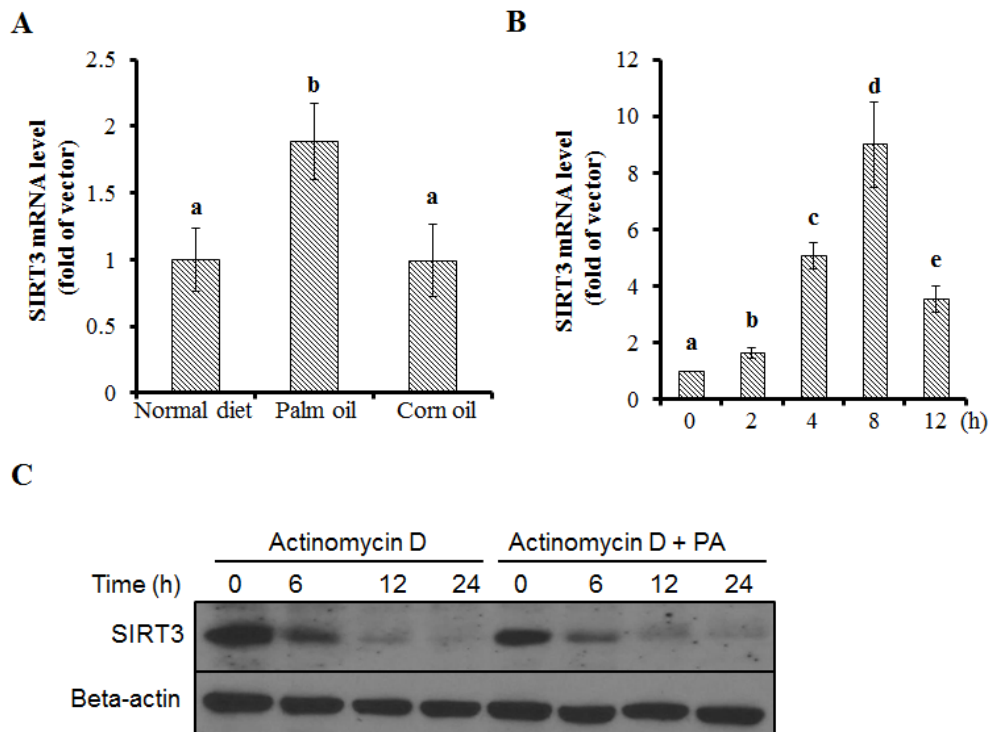


Supplementary Fig. 4 SFAs induces cell death in hepatocytes. (A) & (B) HepG2 cells were treated with 0.25, 0.5, 0.75 mM palmitic acid (PA) for 12 h. LDH release in the culture medium and propidium iodide (PI) staining were detected according to the

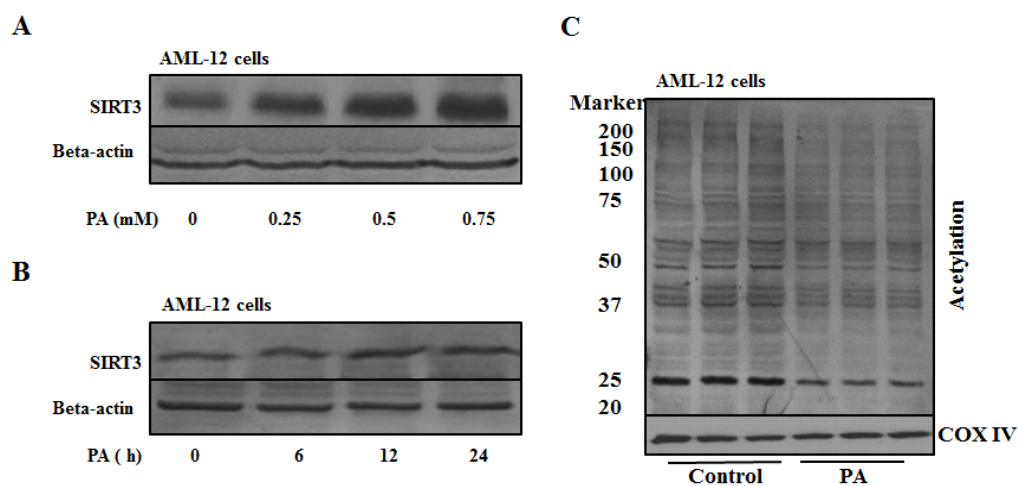
description in the Material and methods, respectively. (C) & (D) HepG2 cells were treated with 0.5 mM (PA) for the indicated duration. LDH release in the culture medium and propidium iodide (PI) staining were detected according to the description in the Material and methods, respectively. (E) & (F) HepG2 cells were exposed to 0.5 mM stearic acid (SA) for 12 h or as indicated in the figure. LDH release in the culture medium was detected according to the description in the Material and methods. (G) & (H) HepG2 cells were treated with 0.25, 0.5, 0.75 mM oleic acid (OA) for 12 h. LDH release in the culture medium and propidium iodide (PI) staining were detected according to the description in the Material and methods, respectively. All values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



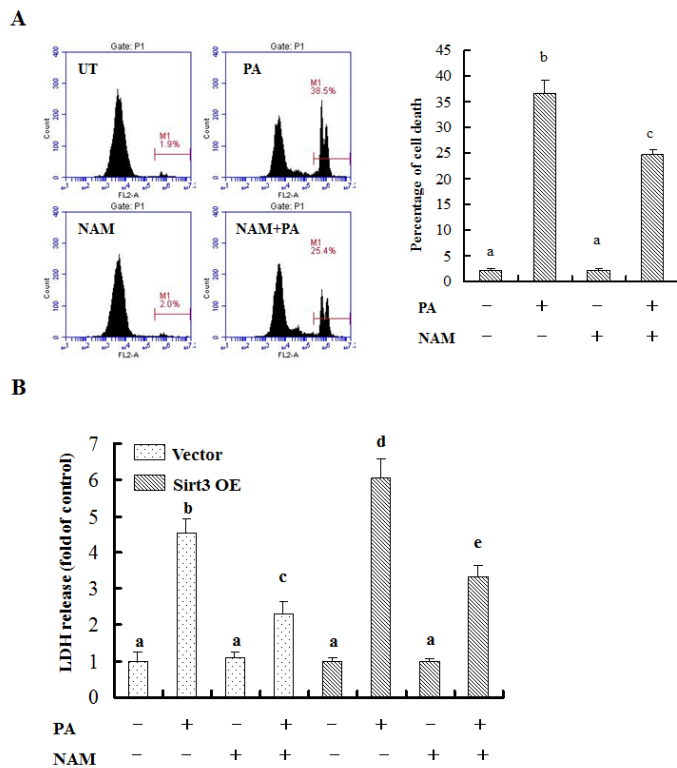
Supplementary Fig. 5 SFAs-rich (palm oil) high-fat diet induces more severe metabolic impairments than unsaturated fatty acids-rich (corn oil) high-fat diet in C57BL/6 mice. Male C57BL/6 mice (8-week of age) were fed with normal or high-fat diet using either palm oil or corn oil as fat source. Mice were sacrificed after 8-week of feeding. Plasma and liver were collected for analysis. (A) Body weight. (B) Plasma triglyceride was determined by a Triglycerides Assay Kit (Thermo Fisher, TR22421). (C) Liver weight. (D) Liver weight/body weight ratio. (E) Liver triglyceride content. (F) Liver TBARS. (G) Epididymal fat weight. (H) Epididymal fat/ body weight ratio. All values are denoted as means \pm SD (n = 8). Bars with different characters differ significantly, $p < 0.05$.



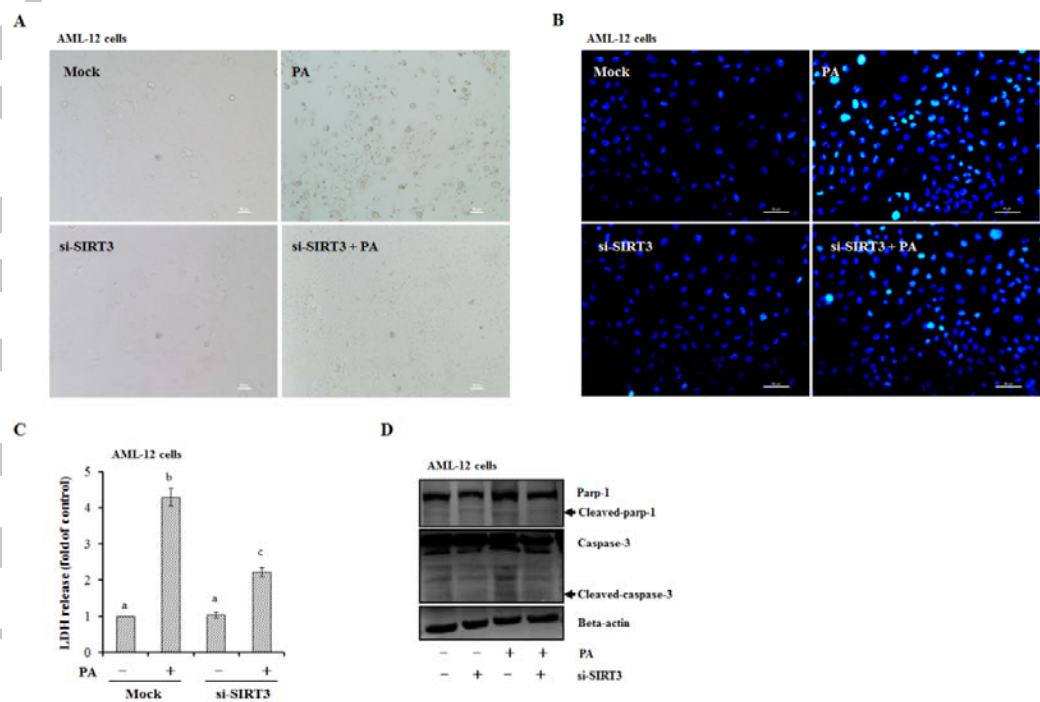
Supplementary Fig. 6 SFAs up-regulates SIRT3 mRNA expression in the liver and HepG2 cells. (A) Male C57BL/6 mice (8-week of age) were fed with normal or high-fat diet using either palm oil or corn oil as fat source. Mice were sacrificed after 8-week of feeding. Liver mRNA was extracted for the analysis of SIRT3 mRNA level. (B) HepG2 cells were treated with 0.5 mM PA for the indicated duration. The mRNA expression of SIRT3 was detected by RT-PCR. (C) HepG2 cells treated with or without 0.5 mM PA for the indicated duration in the presence of actinomycin D (5 μ g/ml). Immunoblotting assay was performed for SIRT3 expressions. Each *in vitro* test was performed at least three times ($n = 8$ for mice), and a representative blot was shown. All values are denoted as means \pm SD. Bars with different characters differ significantly, $p < 0.05$.



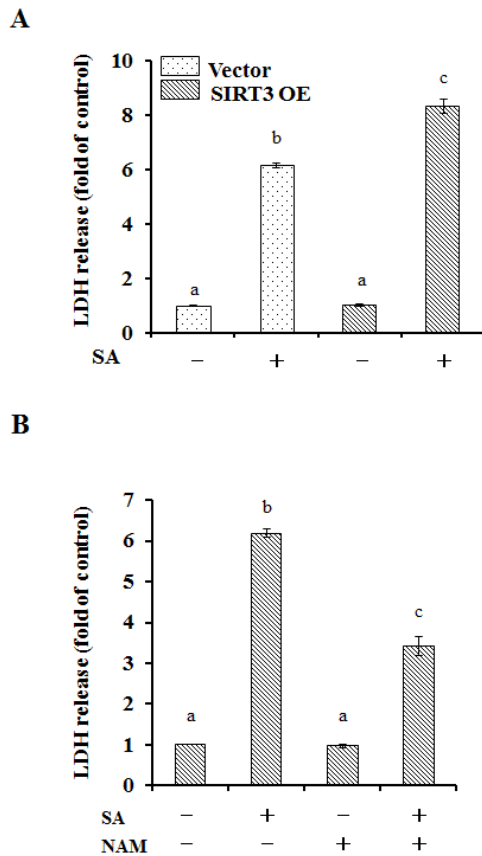
Supplementary Fig. 7 Palmitate increases SIRT3 expression and activity in mouse hepatocytes. (A) AML-12 cells were exposed to 0.25, 0.5, 0.75 mM palmitic acid (PA) for 12 h. Total cellular lysates were subjected to immunoblotting assay for SIRT3. (B) AML-12 cells were treated with 0.5 mM PA for the indicated duration. SIRT3 expression was detected. (C) AML-12 cells were incubated with 0.5 mM PA for 12 h. Mitochondrial proteins were extracted using a commercial Mitochondria Isolation kit (Beyotime, China) for the measurement of acetylated proteins degree.



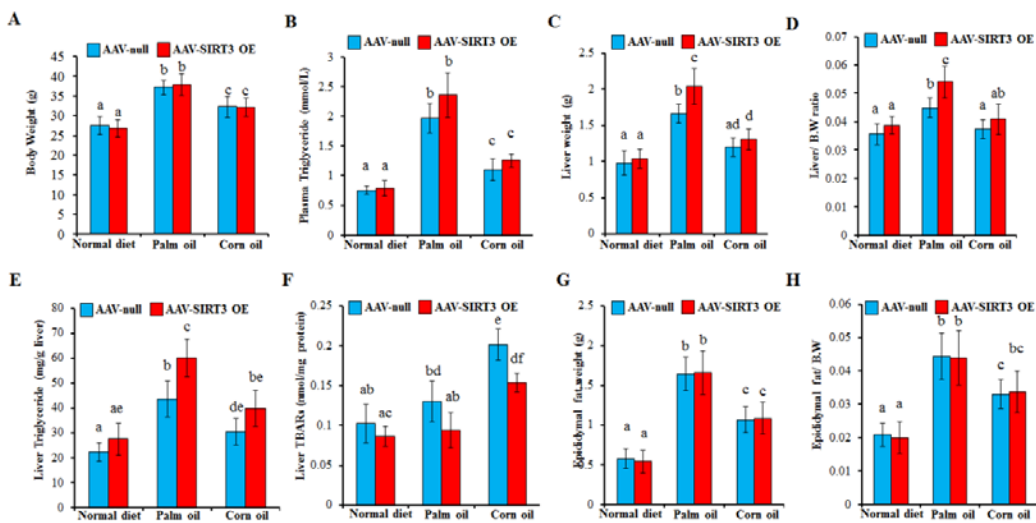
Supplementary Fig. 8 Nicotinamide (NAM) protects hepatocytes against palmitate-induced cell death. (A) HepG2 cells were treated with 0.5 mM palmitic acid (PA) for 12 h with or without 1 h pre-incubation of 5 mM NAM. Cell death was detected via propidium iodide (PI) staining using flow cytometry. (B) SIRT3 OE and vector control HepG2 cells were treated with 0.5 mM PA for 12 h with or without 1 h pre-incubation of NAM. Cell death was detected by the measurements of LDH release. All values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



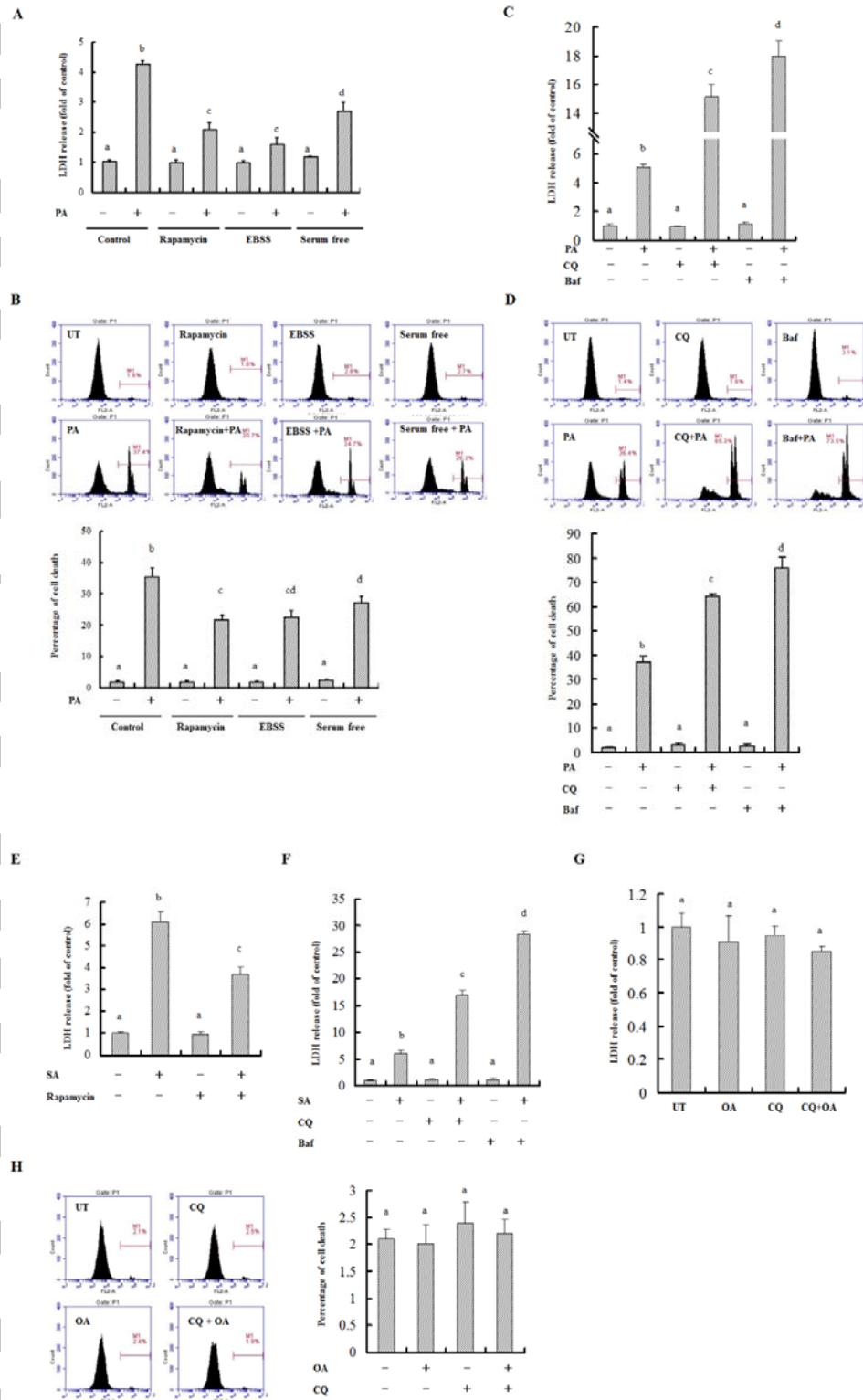
Supplementary Fig. 9 Knocking-down SIRT3 protects palmitate-induced cell death in mouse hepatocytes. AML-12 mouse hepatocytes were transfected with siSIRT3, and exposed to palmitic acid (0.5 mM) treatment for 12 h. Cell death was determined by the following assays. (A) Cellular morphological changes were examined by inverted phase contrast microscope at a scope of 100 ×. (B) Nuclear morphology was detected by Hoechst staining using fluorescence microscopy at a magnification of 200 ×. (C) LDH release was detected as described in the Methods. (D) Total lysates were subjected to immunoblotting assay for Caspase-3 and Parp-1. All values are denoted as means ± SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



Supplementary Fig. 10 SIRT3 overexpression aggravates stearic acid-induced cell death. (A) Vector control or SIRT3 OE HepG2 cells were suffered to stearic acid (SA, 0.5 mM) exposure for 12 h. Cell death was detected by the measurements of LDH release. (B) HepG2 cells were treated with 0.5 mM SA for 12 h with or without 1 h pre-incubation of 5 mM NAM. Cell death was detected by the measurements of LDH release. All values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



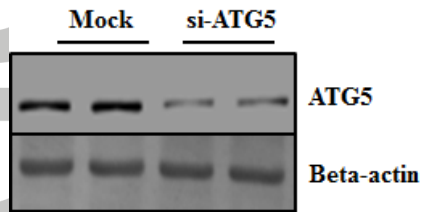
Supplementary Fig. 11 SIRT3 overexpression aggravates SFAs-rich (palm oil) high-fat diet induces metabolic impairments in C57BL/6 mice. Liver specific SIRT3 OE mice model was established and fed with normal or high-fat diet using either palm oil or corn oil as described in the Methods. Plasma and liver were collected for analysis (n = 8). (A) Body weight. (B) Plasma triglyceride was determined by a Triglycerides Assay Kit (Thermo Fisher, TR22421). (C) Liver weight. (D) Liver weight/body weight ratio. (E) Liver triglyceride content. (F) Liver TBARS. (G) Epididymal fat weight. (H) Epididymal fat/ body weight ratio. All values are denoted as means \pm SD (n = 8). Bars with different characters differ significantly, $p < 0.05$.



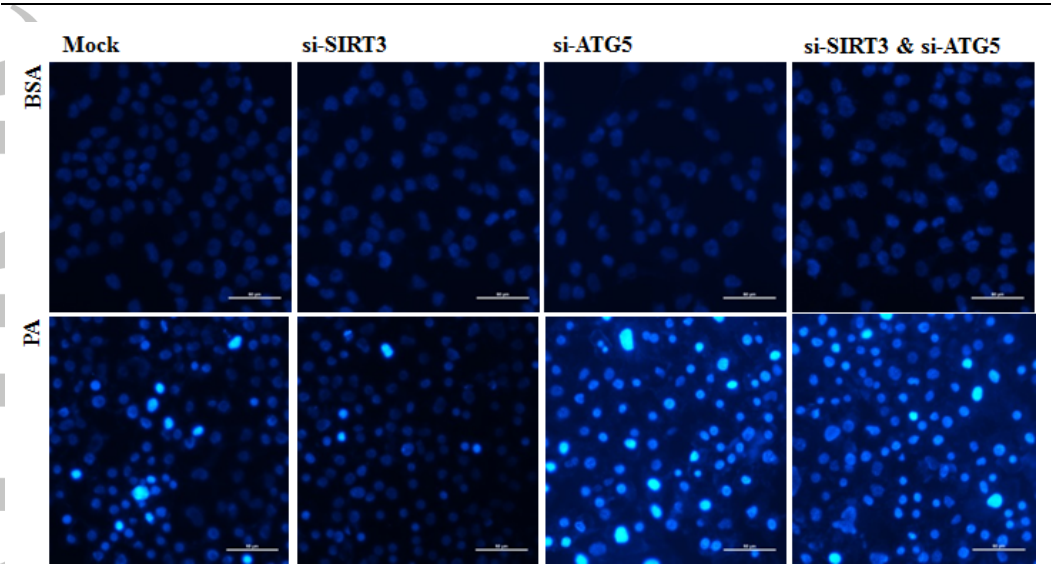
Supplementary Fig. 12 Autophagy regulates SFAs-induced lipotoxicity in HepG2

cells. (A) & (B) HepG2 cells were treated with 0.5 mM PA for 12 h in normal DMEM,

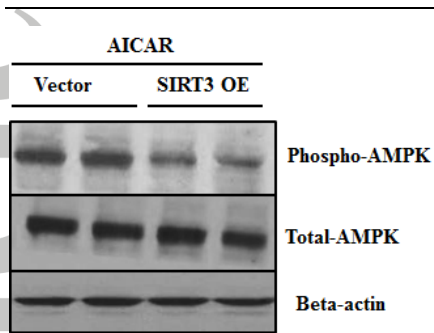
EBSS, or serum free DMEM culture medium, respectively. Rapamycin (50 nM) was added 1 h before PA treatment. Cell death was detected by the measurements of LDH release and propidium iodide (PI) staining using flow cytometry. (C) & (D) HepG2 cells were treated with 0.5 mM PA for 12 h. Autophagy inhibitor, chloroquine (CQ, 20 μ M) or bafilomycin A1 (Baf, 100 nM), was added 1 h before PA treatment, respectively. Cell death was detected by the measurements of LDH release and propidium iodide (PI) staining using flow cytometry. (E) & (F) HepG2 cells were treated with 0.5 mM SA for 12 h. Rapamycin, CQ, and Baf were added 1 h before SA treatment, respectively. Cell death was detected by the measurements of LDH release in the culture medium. (G) & (H) HepG2 cells were treated with 0.5 mM OA for 12 h. CQ was added 1 h before OA treatment. Cell death was detected by the measurements of LDH release and propidium iodide (PI) staining using flow cytometry. All values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



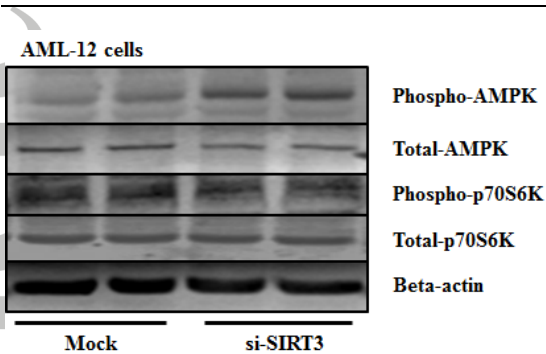
Supplementary Fig. 13 Knocking-down efficiency of siRNA for Atg5. Special siRNA for Atg5 was transfected into HepG2 cells as described in the Methods. Protein expressions of Atg5 were detected for testing the silencing efficiency.



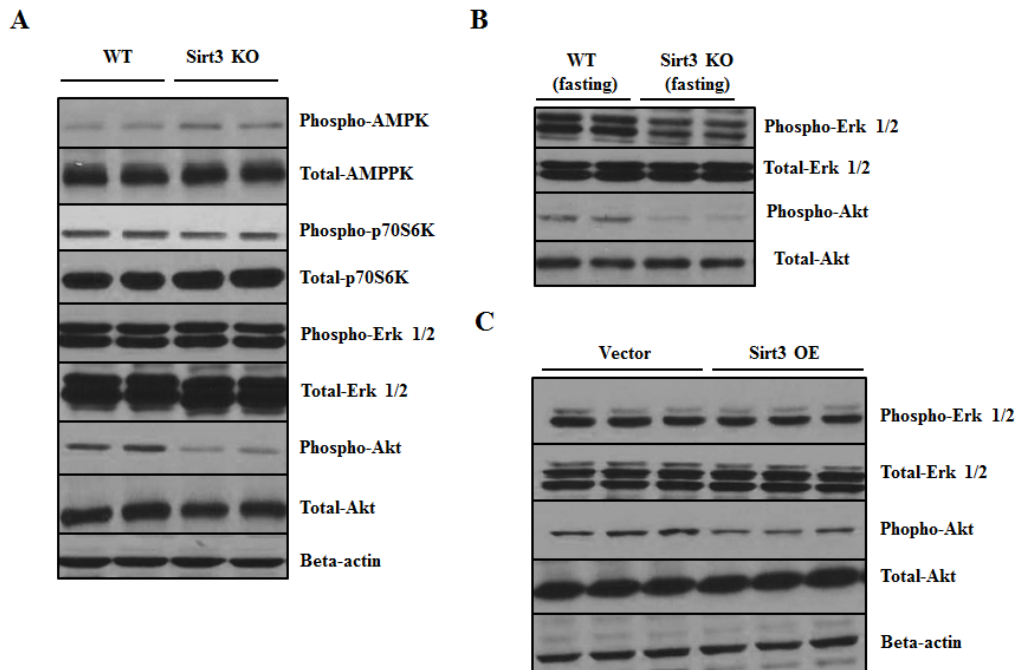
Supplementary Fig. 14 Inhibiting autophagy blocks knocking-down SIRT3-protected lipotoxicity in HepG2 cells. HepG2 cells were co-transfected with siRNA for SIRT3 and Atg5 siRNA, followed with 0.5 mM PA exposure for 12 h. Cell death was detected by Hoechst staining.



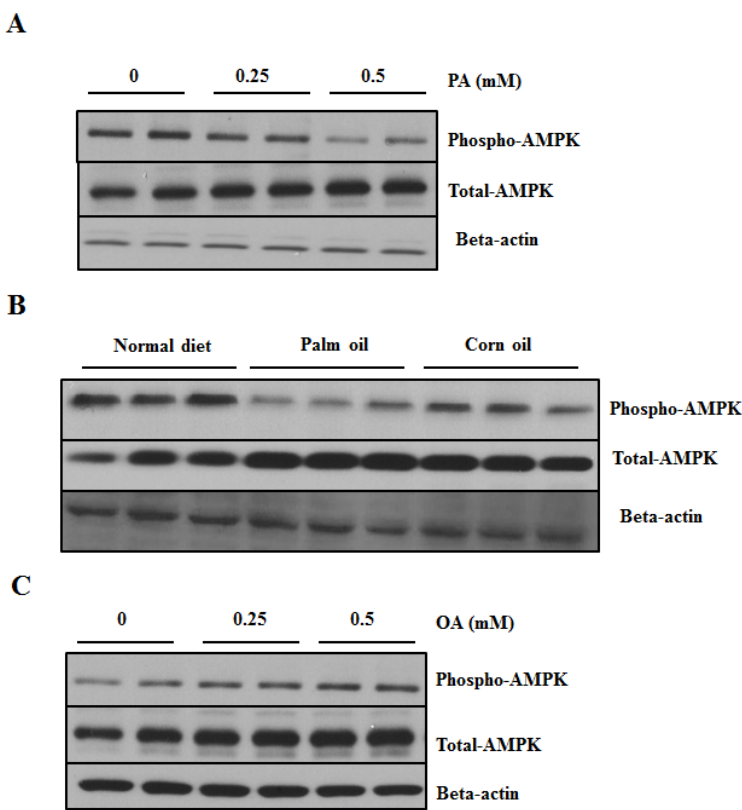
Supplementary Fig. 15 SIRT3-overexpressing decreases AICAR-activated AMPK phosphorylation. SIRT3 OE and vector control HepG2 cells were treated with AMPK agonist AICAR (2.5 mM) for 4 h. Immunoblotting assay was performed for phospho-AMPK.



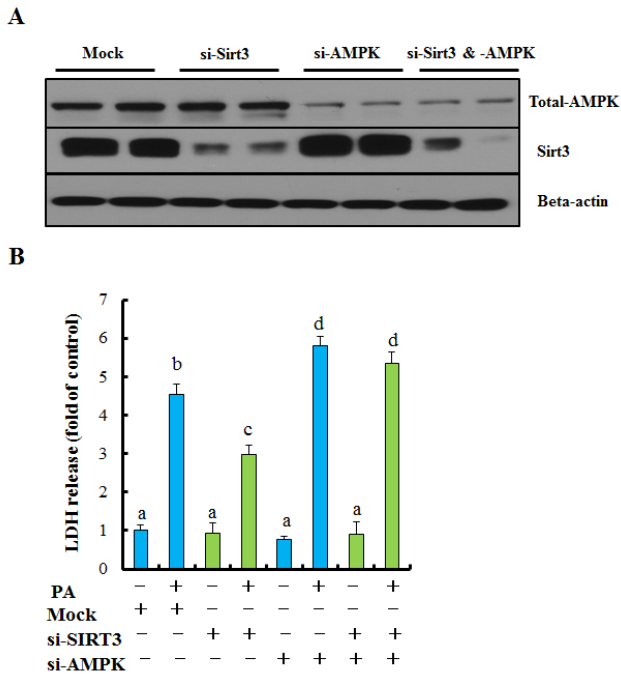
Supplementary Fig. 16 Knocking-down SIRT3 activates AMPK/mTOR axis in mouse hepatocytes. AML-12 mouse hepatocytes were transfected with siSIRT3 or scramble siRNA. Total lysates were subjected to immunoblotting assay for phosphorylated-AMPK and -p70S6K.



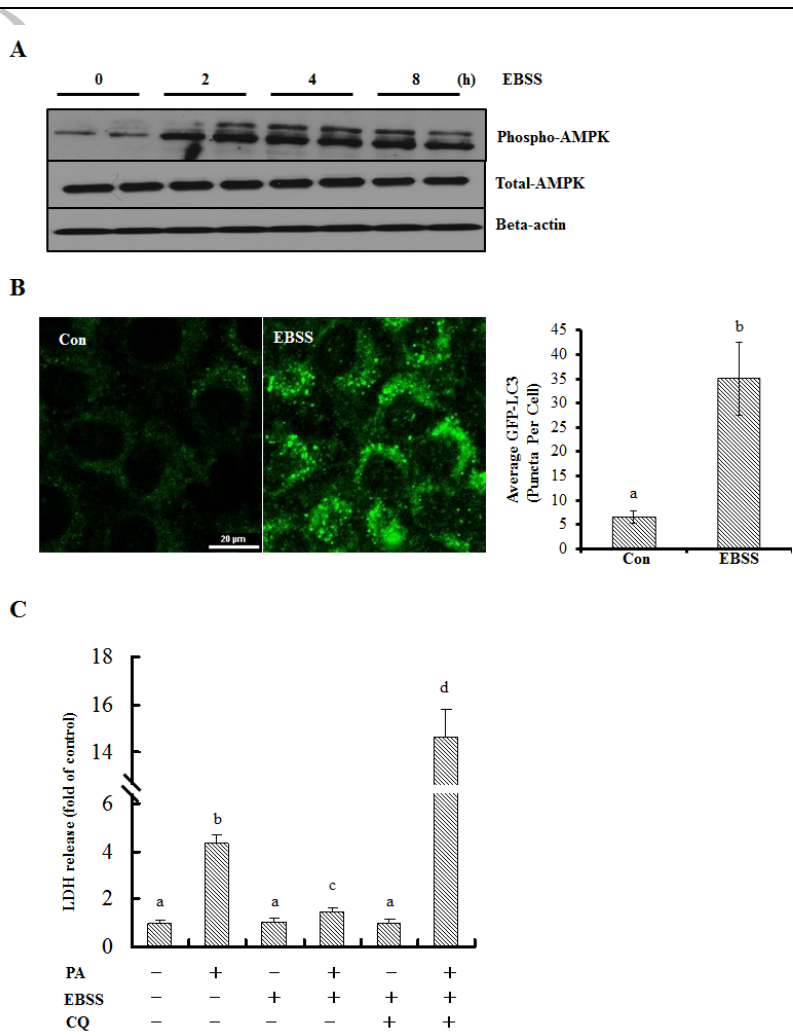
Supplementary Fig. 17 Neither ERK1/2 nor Akt is regulated by SIRT3. Liver tissue was collected from SIRT3 knock-out (KO) or wild type mice with or without 12 h fasting (n = 5). (A) Phosphorylated-AMPK, -p70S6K, -ERK1/2, and -Akt expressions were detected. (B) Phosphorylated-ERK1/2 and -Akt expressions were detected. (C) Immunoblotting assay for phosphorylated-ERK1/2 and -Akt in SIRT3 OE and vector control HepG2 cells.



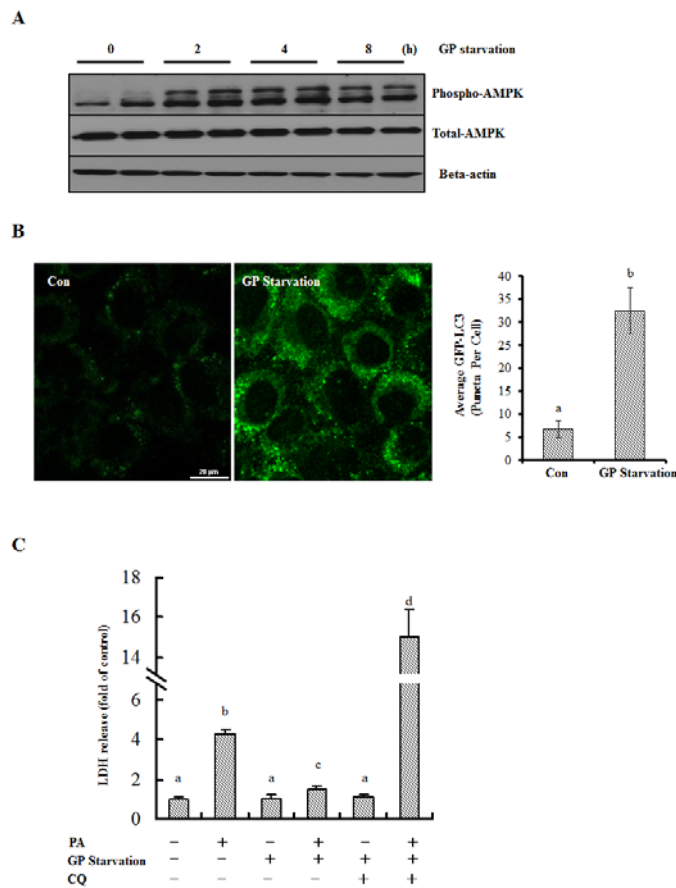
Supplementary Fig. 18 SFAs down-regulates AMPK phosphorylation in the liver and hepatocytes. Male C57BL/6 mice (8-week of age) were fed with normal or high-fat diet using either palm oil or corn oil as fat source. Mice were sacrificed after 8-week of feeding. Liver was collected for analysis. HepG2 cells were treated with PA or OA for 12 h. Total cellular lysates from hepatocytes and liver tissues (n = 8) were subjected to immunoblotting assay for phosphor-AMPK. (A) PA inhibits AMPK phosphorylation in HepG2 cells. (B) Palm oil feeding inhibits AMPK phosphorylation. (C) OA induces AMPK phosphorylation in HepG2 cells.



Supplementary Fig. 19 Silencing AMPK inhibits knocking-down SIRT3-protected lipotoxicity in HepG2 cells. HepG2 cells were co-transfected with siSIRT3 and siAMPK, and were exposed to 0.5 mM PA for 12 h. (A) The transfected efficiency was detected using Western-blotting. (B) Cell death was detected by LDH release. Values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.

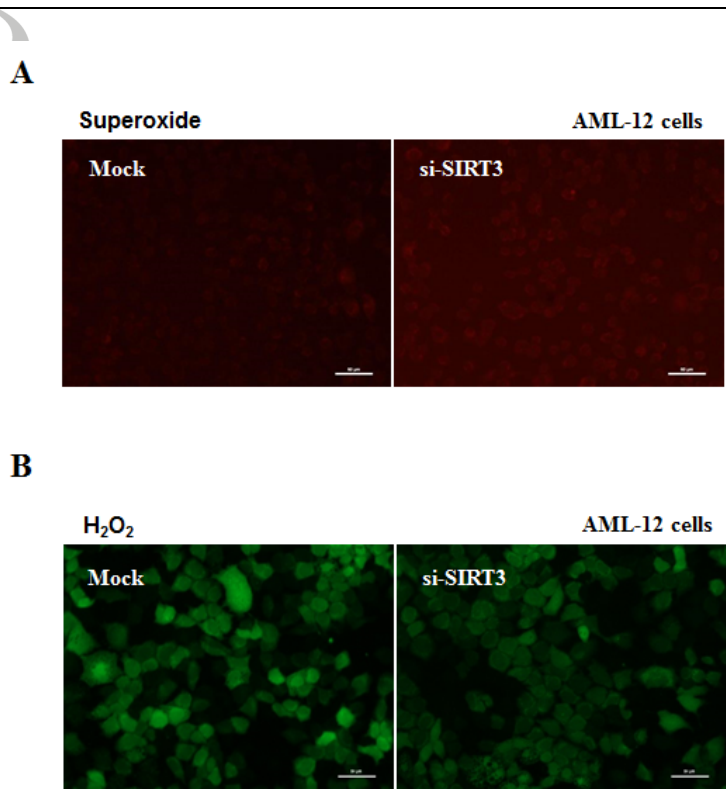


Supplementary Fig. 20 EBSS protects against PA-induced lipotoxicity via activating AMPK-regulated autophagy. (A) HepG2 cells were incubated in EBSS medium for the indicated duration. Phosphorylated-AMPK expression was detected. (B) HepG2 cells were transfected with recombinant adenovirus GFP-LC3 and treated with EBSS for 12 h. Puncta was detected. (C) HepG2 cells were treated with 0.5 mM PA for 12 h with or without EBSS incubation. CQ (20 μ M) was added 1 h before PA treatment. Cell death was detected by the measurements of LDH release. Values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.

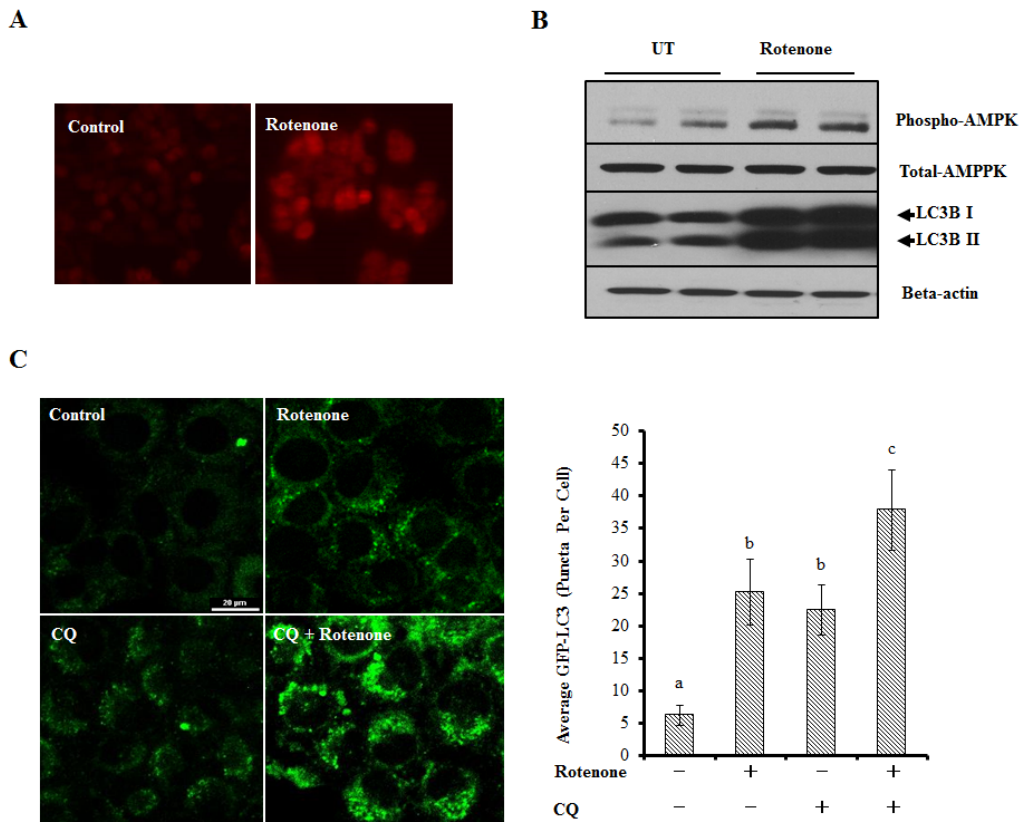


Supplementary Fig. 21 GP starvation protects against PA-induced lipotoxicity via activating AMPK-regulated autophagy. (A) HepG2 cells were incubated in GP starvation (glucose, L-glutamine, pyruvate, and serum-depleted) medium for the indicated duration. Phosphorylated-AMPK expression was detected. (B) HepG2 cells were transfected with recombinant adenovirus GFP-LC3 and treated with GP starvation medium for 12 h. Puncta was detected. (C) HepG2 cells were treated with 0.5 mM PA for 12 h with or without GP starvation incubation. CQ (20 μ M) was added 1 h before PA treatment. Cell death was detected by the measurements of LDH release. Values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.

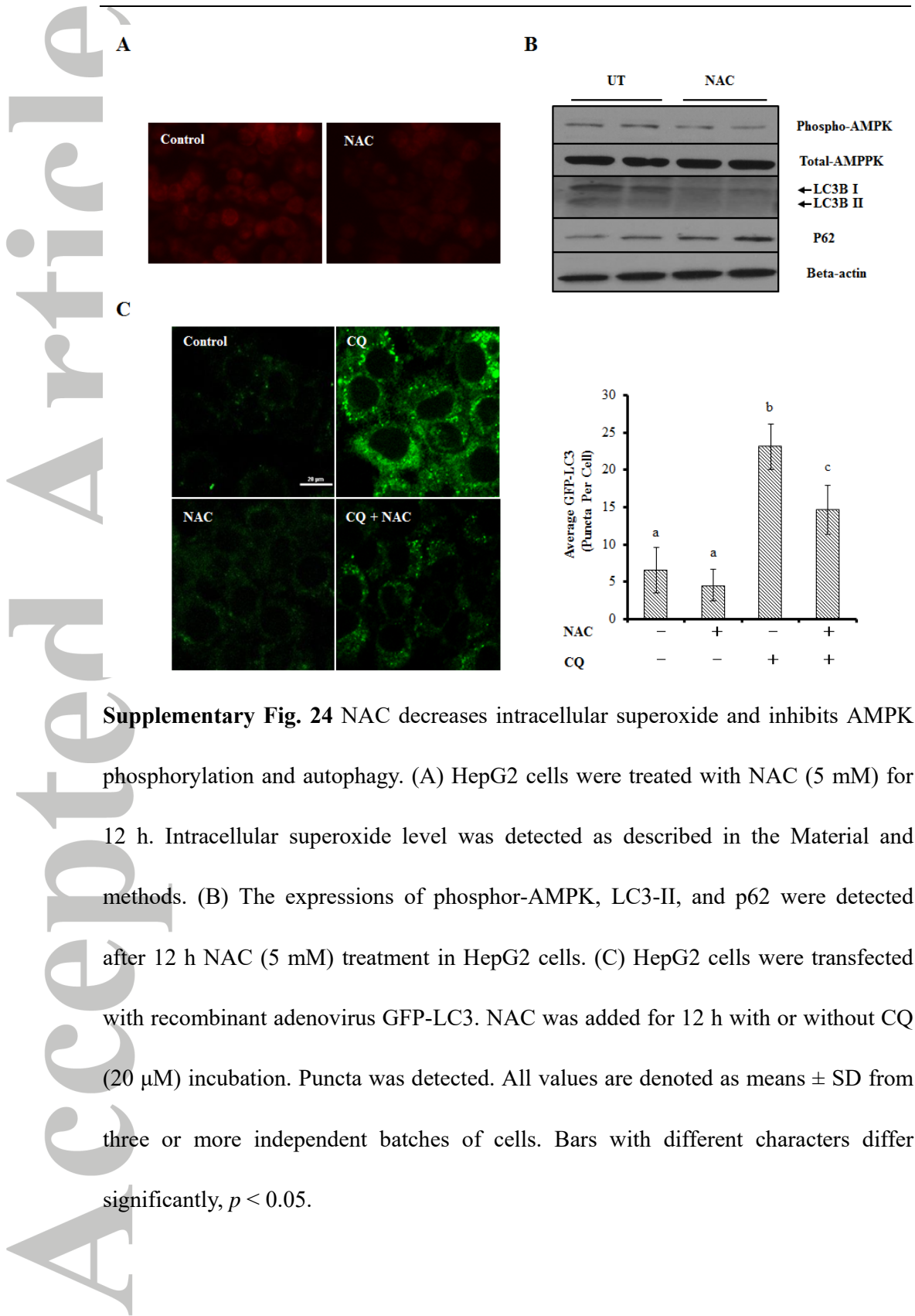
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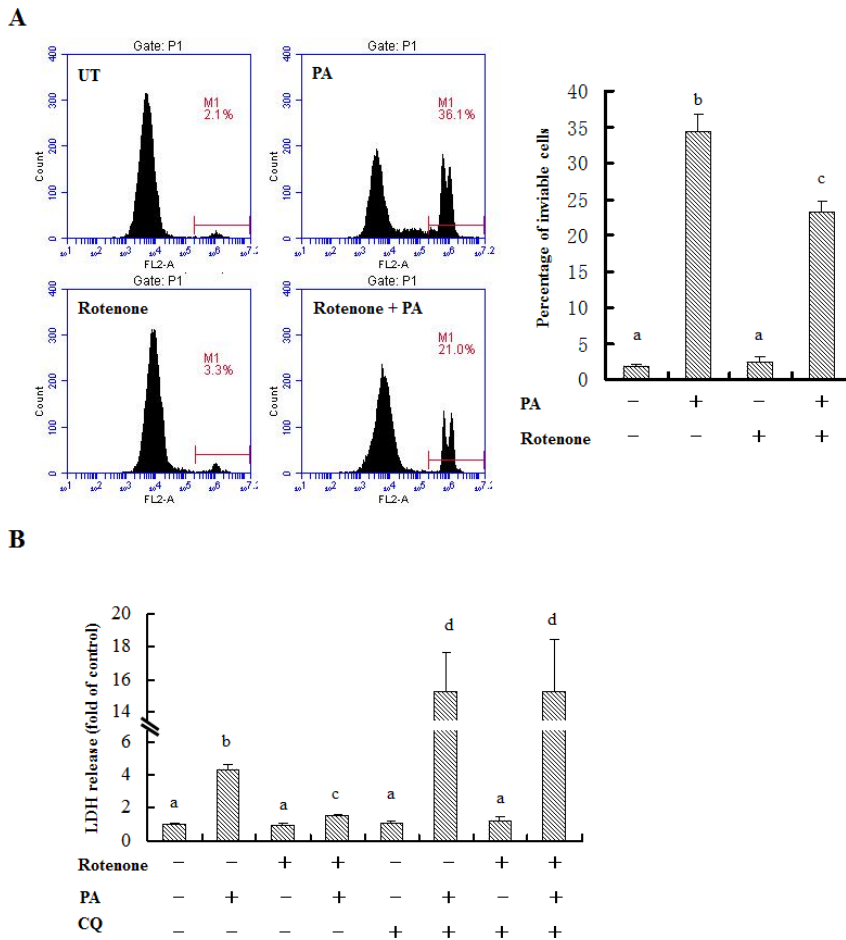
Supplementary Fig. 22 Knocking-down SIRT3 regulates intracellular redox in mouse hepatocytes. AML-12 mouse hepatocytes were transfected with siSIRT3 or scramble siRNA. (A) Intracellular superoxide level was detected as described in the Material and methods. (B) Intracellular H₂O₂ level was detected as described in the Material and methods.



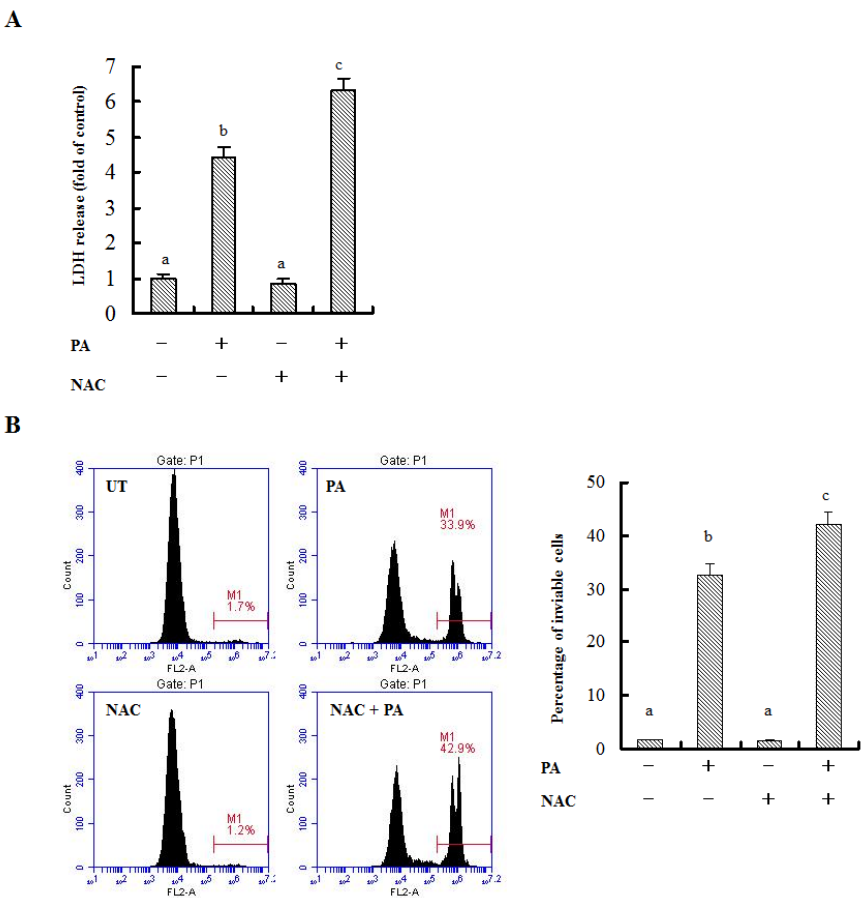
Supplementary Fig. 23 Rotenone increases intracellular superoxide and stimulates AMPK phosphorylation and autophagy. (A) HepG2 cells were treated with rotenone (10 nM) for 12 h. Intracellular superoxide level was detected as described in the Material and methods. (B) The expressions of phosphor-AMPK and LC3-II were detected after 12 h rotenone (10 nM) treatment in HepG2 cells. (C) HepG2 cells were transfected with recombinant adenovirus GFP-LC3 and treated with rotenone for 12 h. CQ (20 μ M) was added 1 h before rotenone treatment. Puncta was detected. All values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



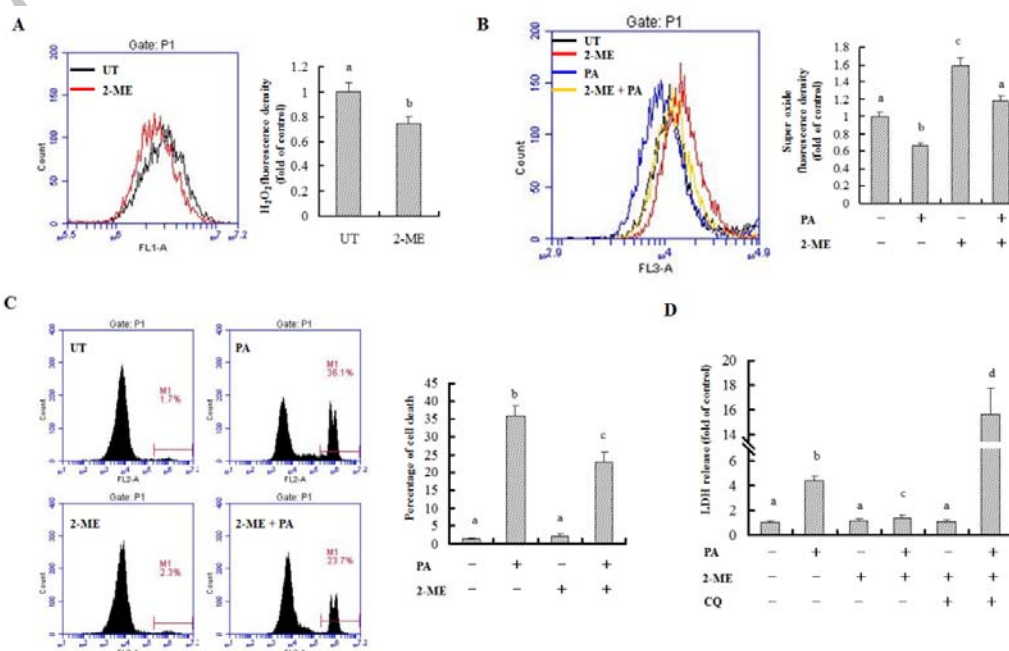
Supplementary Fig. 24 NAC decreases intracellular superoxide and inhibits AMPK phosphorylation and autophagy. (A) HepG2 cells were treated with NAC (5 mM) for 12 h. Intracellular superoxide level was detected as described in the Material and methods. (B) The expressions of phosphor-AMPK, LC3-II, and p62 were detected after 12 h NAC (5 mM) treatment in HepG2 cells. (C) HepG2 cells were transfected with recombinant adenovirus GFP-LC3. NAC was added for 12 h with or without CQ (20 μ M) incubation. Puncta was detected. All values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



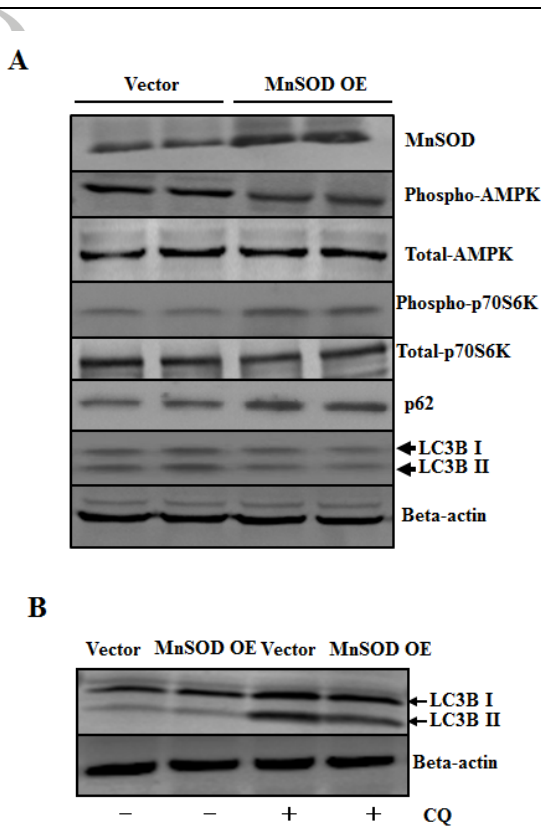
Supplementary Fig. 25 Rotenone protects against PA-induced lipotoxicity via activating autophagy. (A) HepG2 cells were treated with 0.5 mM PA for 12 h with or without rotenone (10 nM) pre-treatment. Cell death was detected by propidium iodide (PI) staining using flow cytometry. (B) HepG2 cells were treated with 0.5 mM PA for 12 h with or without rotenone (10 nM) pre-treatment. CQ (20 μ M) was added 1 h before rotenone treatment. Cell death was detected by the measurements of LDH release. Values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



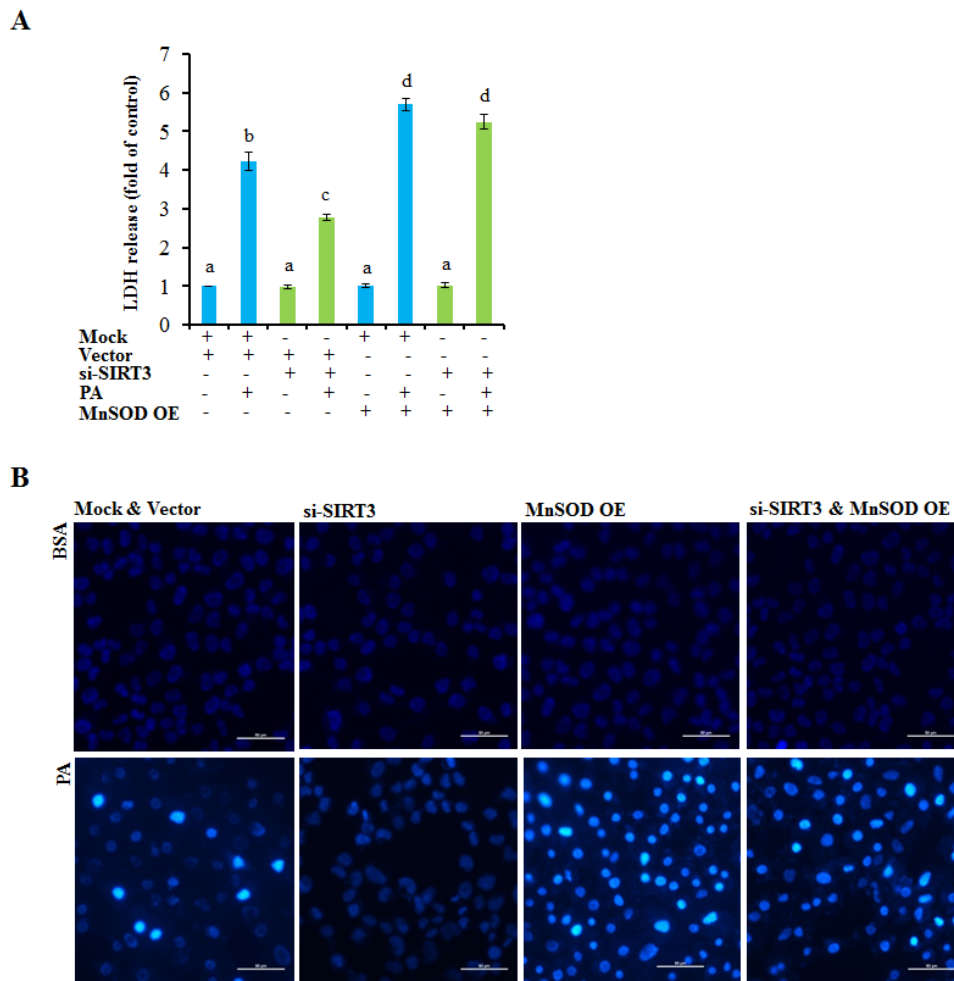
Supplementary Fig. 26 NAC aggravates PA-induced lipotoxicity. HepG2 cells were treated with 0.5 mM PA for 12 h with or without NAC (5 mM) pre-treatment. (A) Cell death was detected by propidium iodide (PI) staining using flow cytometry. (B) Cell death was detected by the measurements of LDH release. Values are denoted as means \pm SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



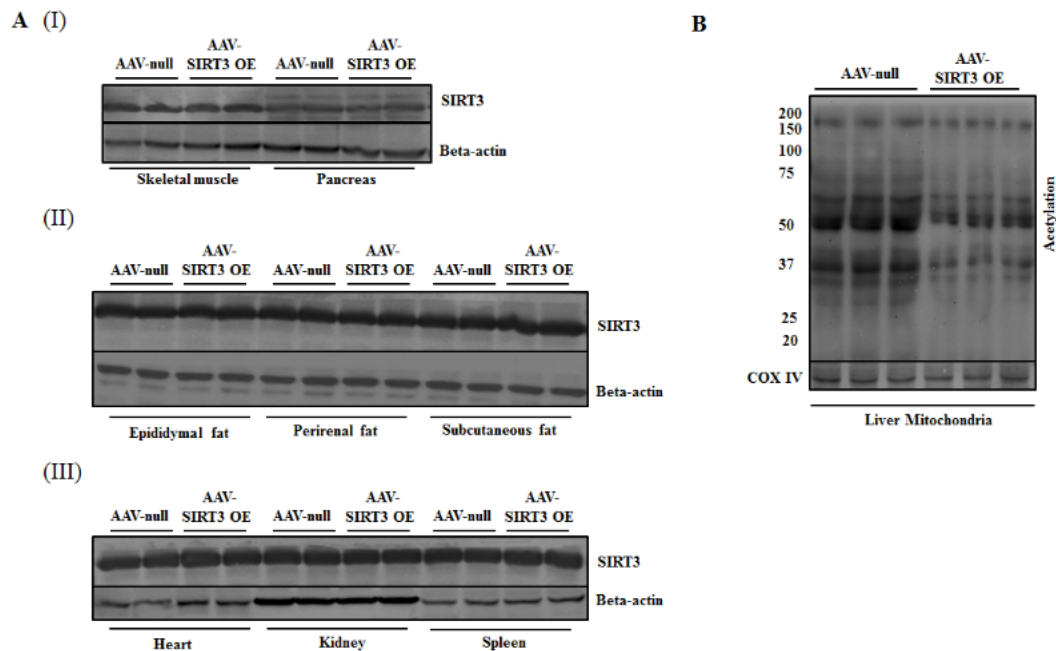
Supplementary Fig. 27 Inhibiting MnSOD protects against PA-induced lipotoxicity via activating autophagy. (A) HepG2 cells were treated with MnSOD inhibitor, 2-ME (100 μM), for 12 h. Intracellular hydrogen peroxide (H₂O₂) level was detected using the method as described in the Material and methods. (B) HepG2 cells were treated with 0.5 mM PA for 12 h with or without 2-ME pre-treatment. Intracellular superoxide level was detected using the method as described in the Material and methods. (C) HepG2 cells were treated with 0.5 mM PA for 12 h with or without 2-ME pre-treatment. Cell death was detected by propidium iodide (PI) staining using flow cytometry. (D) HepG2 cells were treated with 0.5 mM PA for 12 h with or without 2-ME pre-treatment. CQ (20 μM) was added 1 h before 2-ME treatment. Cell death was detected by the measurements of LDH release. All values are denoted as means ± SD from three or more independent batches of cells. Bars with different characters differ significantly, $p < 0.05$.



Supplementary Fig. 28 Over-expressing MnSOD inhibits autophagic flux in HepG2 cells. HepG2 cells were transfected with recombinant lentivirus containing human MnSOD or empty vector as control, and treated with or without CQ (20 μ M) for 12 h. (A) Immunoblotting assay was performed for MnSOD, phospho-AMPK, -p70s6k, p62, and LC3 expressions in MnSOD OE or vector control cells. (B) Immunoblotting assay was performed for LC3-II expression.



Supplementary Fig. 29 Over-expressing MnSOD blocks knocking-down SIRT3-protected lipotoxicity in HepG2 cells. HepG2 cells were co-transfected with siRNA for SIRT3 and recombinant lentivirus containing human MnSOD or empty vector as control, followed with 0.5 mM PA exposure for 12 h. Cell death was detected by LDH release (A) and Hoechst staining (B), respectively. All values are denoted as means \pm SD. Bars with different characters differ significantly, $p < 0.05$.



Supplementary Fig. 30 SIRT3 protein expressions in different tissues from liver specific SIRT3 over-expression mice. (A) Total lysates from different tissues (n = 8) were subjected to immunoblotting assay for SIRT3. (B) Mitochondrial proteins were extracted from liver tissues using a commercial Mitochondria Isolation kit (Beyotime, China) for the measurement of acetylated proteins abundance.

Reference

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