# **1** Supplementary Material

# 2 SIRT3 Acts As a Negative Regulator of Autophagy Dictating Hepatocyte

# 3 Susceptibility to Lipotoxicity

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#### 26 Supplementary Methods

#### 27 Animal Model and Experimental Protocol

All animal studies were performed in accordance with the animal care committee at the 28 University of Illinois at Chicago. All mice were fed ad libitum, and housed on a 12:12 h 29 light: dark cycle at 25°C. Male SIRT3<sup>-/-</sup> mice and WT 129Sv mice (8-week of age) were 30 31 purchased from Jackson Laboratory (Bar Harbor, ME), and maintained on a standard chow 32 diet. Male C57BL/6 mice (8-week old, Jackson Laboratory, Bar Harbor, ME) were randomly 33 assigned to normal diet, palm coil high-fat diet, and corn oil high-fat diet (Supplementary data Table 1). The liver-specific SIRT3 overexpression mice and corresponding control mice 34 were generated via the injection of recombinant adeno-associated viral (AAV) serotype 8 35 gene transfer vectors containing a liver-specific promoter combination (albumin promoter) 36 with either mouse SIRT3 sequence or empty as vector control (Cyagen Biosciences Inc., 37 Guangzhou, China). Finally, there are 6 groups in total (n = 8 mice/group) and the animals 38 39 were fed for 8 weeks. AAV8 vectors were administered by tail vein injection at a dose of  $1 \times 10^{12}$  viral titer/mL in a total volume of 100 µl/mice at the beginning of the experiment. 40 Recombinant AAV8 vectors encoding mouse SIRT3 (NM 022433.2) under the control of 41 albumin promoter (a liver-specific promoter) were generated by Cyagen Biosciences Inc. 42 43 (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 44 Supplementary data, Fig. 30. Mice were sacrificed after 8-week of feeding. Plasma and liver 45 were collected for analysis. Plasma alanine aminotransferase (ALT) content was determined 46 by a commercial ALT kit (Thermo Fisher, Waltham, MA). Triglyceride was assayed using a 47

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 µm 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).

#### 54 *Chemicals*

55 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 56 SA-BSA conjugates were prepared as described previously <sup>(1)</sup>. Briefly, PA or SA was 57 dissolved in ethanol and saponified with sodium hydroxide. The sodium salt was dried, 58 59 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 60 61 4 h to allow PA or SA to bind to BSA. The PA- or SA-BSA complex (3 mmol/L fatty acid: 62 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. 63 64 BSA, ethanol, DMSO).

65 *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  $\mu$ g /ml streptomycin at 37°C in a humidified O<sub>2</sub>/CO<sub>2</sub> (19:1) atmosphere.

70	Alpha mouse liver (AML)-12 hepatocyte culture was established from a mouse transgenic
71	for human transforming growth factor $\alpha$ , and was obtained from the American Type Culture
72	Collection (ATCC, CRL-2254), and was cultured in Dulbecco's Modified Eagle Medium
73	/Ham's Nutrient Mixture F-12, 1:1 (DMEM /F-12, Sigma-Aldrich, 051M8322) containing 10%
74	(v/v) fetal bovine serum (PAA Laboratories, A15-701), 5 mg/ml insulin (Sigma-Aldrich,
75	19278), 5 µg/ml transferrin (Sigma-Aldrich, T8158), 5 ng/ml selenium (Sigma-Aldrich,
76	229865), 40 ng/ml dexamethasone (Sigma-Aldrich, D4902), 100 U/ml penicillin, and 100
77	$\mu$ g/ml streptomycin (Life technologies, 15140-122) at 37°C in a humidified O <sub>2</sub> /CO <sub>2</sub> (95:5)
78	atmosphere.

## 79 Establishment of stable human SIRT3 over-expression cell lines

The eukaryotic expression vector plasmid pcDNA3.1<sup>+</sup>/hSIRT3 containing the human SIRT3 80 gene sequence was kindly provided by Dr. Yu Wang (University of Hong Kong)<sup>(2)</sup>. HepG2 81 82 cells grown to 80-90% confluence were transfected with 0.8 µg/well of the expression 83 vector pcDNA3.1<sup>+</sup>/hSIRT3 or empty vector control pcDNA3.1<sup>+</sup> (Invitrogen, Grand Island, 84 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 85 into 24-well plates at a ratio of 1:100 for monoclonal cell selection using 800  $\mu$ g/ml G418. 86 The stable cell line was established and confirmed by both western blot analysis and Real 87 88 Time PCR. SIRT3 over-expression cells and vector control cells were cultured in DMEM containing 10% (v/v) fetal bovine serum, 400 µg/ml G418, 100 U/ml penicillin, and 100 89 90  $\mu$ g/ml streptomycin at 37°C in a humidified O<sub>2</sub>/CO2 (19:1) atmosphere.

91 MnSOD Overexpression

92 Recombinant lentivirus vector (pLV[Exp]-EGFP:T2A:Neo) containing human MnSOD 93 (NM\_001024465.1) under the control of EF-1alpha promoter was generated by Cyagen 94 Biosciences Inc. (Guangzhou, China). A noncoding vector carrying the EF1alpha promoter 95 was used to produce vector control particles. The optimal virus titer used for cell transfection 96 was screened according to the manufacturer's instructions. HepG2 cells were transfected 97 with either pLV[Exp]-EGFP:T2A:Neo-EF1alpha-hMnSOD to overexpress MnSOD or 98 pLV[Exp]-EGFP:T2A:Neo-EF1alpha-null as a vector control.

### 99 Cell Death Assays

 $2 \times 10^5$  /ml cells were seeded in 24-well plates and, after the indicated treatments, cell death 100 101 was determined by measurement of LDH release, propidium iodide (PI) staining, or Hoechst 102 staining. For LDH assay, culture medium was collected and detected using an LDH assay kit 103 (Thermo Scientific Inc, VA) according to the manufacturer's instructions. For PI staining, 104 cells were trypsinized and stained with PI staining solution (BD Pharmingen, CA) according 105 to the manufacturer's instructions. Fluorescence was measured by flow cytometry (Accuri c6, 106 BD, CA). For Hoechst staining, cells were stained with Hoechst staining solution (Beyotime 107 Biotechnology, Nantong, China) according to the manufacturer's instructions and imaged by Nikon eclipse Ti-S fluorescence microscope (Nikon, Tokyo, Japan). 108

#### 109 Analysis of GFP-LC3 puncta

110 The recombinant adenovirus GFP-LC3  $(1 \times 10^{10} \text{ viral titer/mL})$  was obtained from Hanbio 111 Biotechnology Co. Ltd. (Shanghai, China). Cells were transiently transfected with 112 recombinant adenovirus GFP-LC3 according to the manufacturer's instruction. Puncta was 113 detected by a laser scanning confocal microscope (Nikon A1R, Japan) from at least 50 cells 114 for each individual experiment after different treatment.

### 115 Analysis of autophagic flux

The autophagic flux was measured according to methods described previously <sup>(3)</sup>. The cells 116 were pretreated with chloroquine (CQ), inhibitor of lysosome acidification, after genetically 117 overexpressing or knocking-down SIRT3. The autophagic flux was determined via detecting 118 GFP-LC3 puncta and LC3-II expression using laser scanning confocal microscope and 119 120 Western blot, respectively. Additionally, cells were transiently transfected with recombinant 121 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 122 123 at least 50 cells for each individual experiment after different treatment.

#### 124 *Real Time-PCR*

125 Total RNA isolation, reverse transcription, and real time PCR were performed as described 126 previously <sup>(4)</sup>. Briefly, total RNA from cultured cells was isolated with a phenol-chloroform 127 extraction. For each sample, 1 µg total RNA was reverse-transcribed using a high-capacity 128 cDNA reverse transcription kit (Applied Biosystems, Foster City, CA). The cDNA was 129 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 130 131 detection system. Relative gene expression was calculated after normalization by a 132 house-keeping gene (18S rRNA).

#### 133 Western-blot Analysis

134 Western-blot was performed as described previously <sup>(4)</sup> and the following antibodies

135 were used: Anti-Sirt3, anti-phospho-AMPK, anti-AMPK, anti-phospho-Akt, anti-Akt,

136 anti-phospho-Erk1/2, anti-Erk1/2, anti-phospho-p70S6K, anti-p70S6K, anti-LC3B, 137 anti-p62, anti-phospho-ACC, anti-ACC, anti-MnSOD, anti-acetylation, anti-COX IV, 138 anti-caspase-3, anti-parp-1, anti-Atg5, and anti-beta-actin antibodies (Cell Signaling 139 Technology, Danvers, MA). *Immunoprecipitation* 140 141 Cells were lysed in an immunoprecipitation (IP) buffer (150 mM NaCl, 50 mM Tris-HCl, 1% 142 Nonidet P-40, pH 7.8, and a mammalian cell-specific protease inhibitor cocktail. Total 143 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 144 145 with Protein A/G agarose (Santa Cruz Biotechnology) for 1 hr at 4°C. The 146 immunoprecipitates were washed four times with IP buffer. The washed immunoprecipitates 147 were incubated in 50  $\mu$ l of 1 × electrophoresis loading buffer and heated at 100 °C for 5 min. 148 The beads were spun out and the supernatant was resolved by SDS-PAGE, and the 149 modification of MnSOD by Sirt3 was analyzed by Western blot.

#### 150 **RNA interference**

151 Cultured cells were transfected with human SIRT3 siRNA, human AMPK siRNA, human or 152 Atg5 siRNA (Santa Cruz Biotechnology) using Lipofectamine 2000 according to the 153 manufacturer's instructions. In the control group, cells were transfected with scrambled 154 siRNA (Santa Cruz Biotechnology).

## 155 *Reactive oxygen species (ROS) detection*

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

measurement of  $H_2O_2$  after the indicated treatments. Intracellular  $O_2$ <sup>--</sup> 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.

# 162 Supplemental Table

# 163 Supplementary Table 1

		High-fat diet			
	Normal 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		

## **166** 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

177 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.



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

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



198 Supplementary Fig. 5 SFAs-rich (palm oil) high-fat diet induces more severe 199 metabolic impairments than unsaturated fatty acids-rich (corn oil) high-fat diet in 200 C57BL/6 mice. Male C57BL/6 mice (8-week of age) were fed with normal or high-fat 201 diet using either palm oil or corn oil as fat source. Mice were sacrificed after 8-week 202 of feeding. Plasma and liver were collected for analysis. (A) Body weight. (B) Plasma 203 triglyceride was determined by a Triglycerides Assay Kit (Thermo Fisher, TR22421). 204 (C) Liver weight. (D) Liver weight/body weight ratio. (E) Liver triglyceride content. 205 (F) Liver TBARS. (G) Epididymal fat weight. (H) Epididymal fat/ body weight ratio. 206 All values are denoted as means  $\pm$  SD (n = 8). Bars with different characters differ significantly, p < 0.05. 207 208



Supplementary Fig. 6 SFAs up-regulates SIRT3 mRNA expression in the liver and 210 211 HepG2 cells. (A) Male C57BL/6 mice (8-week of age) were fed with normal or 212 high-fat diet using either palm oil or corn oil as fat source. Mice were sacrificed after 213 8-week of feeding. Liver mRNA was extracted for the analysis of SIRT3 mRNA level. 214 (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 215 216 without 0.5 mM PA for the indicated duration in the presence of actinomycin D (5  $\mu$ g/ ml). Immunoblotting assay was performed for SIRT3 expressions. Each in vitro test 217 218 was performed at least three times (n = 8 for mice), and a representative blot was 219 shown. All values are denoted as means  $\pm$  SD. Bars with different characters differ 220 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 acetylizad proteins degree.



231 **Supplementary** Fig. 8 Nicotinamide (NAM) protects hepatocytes against 232 palmitate-induced cell death. (A) HepG2 cells were treated with 0.5 mM palmitic acid 233 (PA) for 12 h with or without 1 h pre-incubation of 5 mM NAM. Cell death was 234 detected via propidium iodide (PI) staining using flow cytometry. (B) SIRT3 OE and 235 vector control HepG2 cells were treated with 0.5 mM PA for 12 h with or without 1 h 236 pre-incubation of NAM. Cell death was detected by the measurements of LDH release. 237 All values are denoted as means  $\pm$  SD from three or more independent batches of cells. 238 Bars with different characters differ significantly, p < 0.05.



240 Supplementary Fig. 9 Knocking-down SIRT3 protects palmitate-induced cell death in mouse hepatocytes. AML-12 mouse hepatocytes were transfected with siSIRT3, and 241 242 exposed to palmitic acid (0.5 mM) treatment for 12 h. Cell death was determined by 243 the following assays. (A) Cellular morphological changes were examined by inverted phase contrast microscope at a scope of 100 ×. (B) Nuclear morphology was detected 244 245 by Hoechst staining using fluorescence microscopy at a magnification of  $200 \times (C)$ 246 LDH release was detected as described in the Methods. (D) Total lysates were 247 subjected to immunoblotting assay for Caspase-3 and Parp-1. All values are denoted as 248 means  $\pm$  SD from three or more independent batches of cells. Bars with different 249 characters differ significantly, p < 0.05.



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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.





260 Supplementary Fig. 11 SIRT3 overexpression aggravates SFAs-rich (palm oil) 261 high-fat diet induces metabolic impairments in C57BL/6 mice. Liver specific SIRT3 262 OE mice model was established and fed with normal or high-fat diet using either palm 263 oil or corn oil as described in the Methods. Plasma and liver were collected for 264 analysis (n = 8). (A) Body weight. (B) Plasma triglyceride was determined by a 265 Triglycerides Assay Kit (Thermo Fisher, TR22421). (C) Liver weight. (D) Liver weight/body weight ratio. (E) Liver triglyceride content. (F) Liver TBARS. (G) 266 Epididymal fat weight. (H) Epididymal fat/ body weight ratio. All values are denoted 267 268 as means  $\pm$  SD (n = 8). Bars with different characters differ significantly, p < 0.05.



271 Supplementary Fig. 12 Autophagy regulates SFAs-induced lipotoxicity in HepG2
272 cells. (A) & (B) HepG2 cells were treated with 0.5 mM PA for 12 h in normal DMEM,

273	EBSS, or serum free DMEM culture medium, respectively. Rapamycin (50 nM) was
274	added 1 h before PA treatment. Cell death was detected by the measurements of LDH
275	release and propidium iodide (PI) staining using flow cytometry. (C) & (D) HepG2
276	cells were treated with 0.5 mM PA for 12 h. Autophagy inhibitor, chloroquine (CQ, 20
277	$\mu$ M) or bafilomycin A1 (Baf, 100 nM), was added 1 h before PA treatment,
278	respectively. Cell death was detected by the measurements of LDH release and
279	propidium iodide (PI) staining using flow cytometry. (E) & (F) HepG2 cells were
280	treated with 0.5 mM SA for 12 h. Rapamycin, CQ, and Baf were added 1 h before SA
281	treatment, respectively. Cell death was detected by the measurements of LDH release
282	in the culture medium. (G) & (H) HepG2 cells were treated with 0.5 mM OA for 12 h.
283	CQ was added 1 h before OA treatment. Cell death was detected by the measurements
284	of LDH release and propidium iodide (PI) staining using flow cytometry. All values
285	are denoted as means $\pm$ SD from three or more independent batches of cells. Bars with
286	different characters differ significantly, $p < 0.05$ .
287	



289 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

291 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.



299 Supplementary Fig. 15 SIRT3-overexpressing decreases AICAR-activated AMPK

- phosphorylation. SIRT3 OE and vector control HepG2 cells were treated with AMPK
- 301 agonist AICAR (2.5 mM) for 4 h. Immunoblotting assay was performed for

302 phospho-AMPK.



304 **Supplementary Fig. 16** Knocking-down SIRT3 activates AMPK/mTOR axis in 305 mouse hepatocytes. AML-12 mouse hepatocytes were transfected with siSIRT3 or 306 scramble siRNA. Total lysates were subjected to immunoblotting assay for 307 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 316 317 hepatocytes. Male C57BL/6 mice (8-week of age) were fed with normal or high-fat 318 diet using either palm oil or corn oil as fat source. Mice were sacrificed after 8-week 319 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 320 subjected to immunoblotting assay for phosphor-AMPK. (A) PA inhibits AMPK 321 322 phosphorylation in HepG2 cells. (B) Palm oil feeding inhibits AMPK phosphorylation. 323 (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.



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332 Supplementary Fig. 20 EBSS protects against PA-induced lipotoxicity via activating 333 AMPK-regulated autophagy. (A) HepG2 cells were incubated in EBSS medium for the indicated duration. Phosphorylated-AMPK expression was detected. (B) HepG2 cells 334 335 were transfected with recombinant adenovirus GFP-LC3 and treated with EBSS for 12 336 h. Puncta was detected. (C) HepG2 cells were treated with 0.5 mM PA for 12 h with or 337 without EBSS incubation. CQ (20 µM) was added 1 h before PA treatment. Cell death 338 was detected by the measurements of LDH release. Values are denoted as means  $\pm$  SD 339 from three or more independent batches of cells. Bars with different characters differ 340 significantly, p < 0.05.



Supplementary Fig. 21 GP starvation protects against PA-induced lipotoxicity via 342 343 activating AMPK-regulated autophagy. (A) HepG2 cells were incubated in GP 344 starvation (glucose, L-glutamine, pyruvate, and serum-depleted) medium for the 345 indicated duration. Phosphorylated-AMPK expression was detected. (B) HepG2 cells 346 were transfected with recombinant adenovirus GFP-LC3 and treated with GP 347 starvation medium for 12 h. Puncta was detected. (C) HepG2 cells were treated with 348 0.5 mM PA for 12 h with or without GP starvation incubation. CQ (20  $\mu$ M) was added 349 1 h before PA treatment. Cell death was detected by the measurements of LDH release. 350 Values are denoted as means  $\pm$  SD from three or more independent batches of cells. Bars with different characters differ significantly, p < 0.05. 351



**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 H<sub>2</sub>O<sub>2</sub> level was detected as described in the Material and methods.



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359 Supplementary Fig. 23 Rotenone increases intracellular superoxide and stimulates 360 AMPK phosphorylation and autophagy. (A) HepG2 cells were treated with rotenone 361 (10 nM) for 12 h. Intracellular superoxide level was detected as described in the 362 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 363 364 transfected with recombinant adenovirus GFP-LC3 and treated with rotenone for 12 h. CQ (20 µM) was added 1 h before rotenone treatment. Puncta was detected. All values 365 366 are denoted as means  $\pm$  SD from three or more independent batches of cells. Bars with different characters differ significantly, p < 0.05. 367



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369 Supplementary Fig. 24 NAC decreases intracellular superoxide and inhibits AMPK 370 phosphorylation and autophagy. (A) HepG2 cells were treated with NAC (5 mM) for 371 12 h. Intracellular superoxide level was detected as described in the Material and 372 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 373 374 with recombinant adenovirus GFP-LC3. NAC was added for 12 h with or without CQ (20  $\mu$ M) incubation. Puncta was detected. All values are denoted as means  $\pm$  SD from 375 376 three or more independent batches of cells. Bars with different characters differ significantly, p < 0.05. 377



379 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 380 381 without rotenone (10 nM) pre-treatment. Cell death was detected by propidium iodide 382 (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  $\mu$ M) was added 1 h 383 384 before rotenone treatment. Cell death was detected by the measurements of LDH 385 release. Values are denoted as means  $\pm$  SD from three or more independent batches of 386 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.



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Supplementary Fig. 27 Inhibiting MnSOD protects against PA-induced lipotoxicity 395 396 via activating autophagy. (A) HepG2 cells were treated with MnSOD inhibitor, 2-ME 397 (100  $\mu$ M), for 12 h. Intracellular hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) level was detected using the method as described in the Material and methods. (B) HepG2 cells were treated 398 399 with 0.5 mM PA for 12 h with or without 2-ME pre-treatment. Intracellular 400 superoxide level was detected using the method as described in the Material and 401 methods. (C) HepG2 cells were treated with 0.5 mM PA for 12 h with or without 402 2-ME pre-treatment. Cell death was detected by propidium iodide (PI) staining using 403 flow cytometry. (D) HepG2 cells were treated with 0.5 mM PA for 12 h with or 404 without 2-ME pre-treatment. CQ (20  $\mu$ M) was added 1 h before 2-ME treatment. Cell 405 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 406 characters differ significantly, p < 0.05. 407



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



418 **Supplementary Fig. 29** Over-expressing MnSOD blocks knocking-down 419 SIRT3-protected lipotoxicity in HepG2 cells. HepG2 cells were co-transfected with 420 siRNA for SIRT3 and recombinant lentivirus containing human MnSOD or empty 421 vector as control, followed with 0.5 mM PA exposure for 12 h. Cell death was 422 detected by LDH release (A) and Hoechst staining (B), respectively. All values are 423 denoted as means  $\pm$  SD. Bars with different characters differ significantly, p < 0.05.

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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.

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