1	Lysophosphatidylcholine acyltransferase 3 is decreased in
2	non-alcoholic steatohepatitis, resulting in caspase-
3	independent hepatocyte death
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29	Running title: Lysophosphatidylcholine acyltransferase 3 in lipotoxicity
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31	Number of:
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37	Abbreviations: BEL, bromoenol lactone; DAG, diacylglycerol; ER,
38	endoplasmic reticulum; FFA, free fatty acids; GSK-3, glycogen synthase
39	kinase-3; HPH, human primary hepatocytes; JNK, c-Jun N-terminal kinase;
40	LPC, lysophosphatidylcholine; lysophosphatidylcholine acyltransferase,

- 41 LPCAT; NAFLD, nonalcoholic fatty liver disease; NASH, nonalcoholic
- 42 steatohepatitis; $PACOCF_3$, palmityl trifluoromethyl ketone; PC,
- 43 phosphatidylcholine; PLA2, phospholipase A2; RIP, receptor-interacting
- 44 protein kinases

1 Background & Aims: Lipotoxicity causes liver inflammation, which leads to 2 non-alcoholic steatohepatitis (NASH). Lysophosphatidylcholine (LPC) is a 3 causal agent of lipotoxicity. Recently, lysophosphatidylcholine acyltransferase (LPCAT) was identified as an enzyme that catalyzes the 4 5 esterification of LPC, which potentially decreases LPC levels. However, the 6 effect of LPCAT in lipotoxicity of the liver is not fully understood. Our aim 7 was to determine whether LPCAT attenuates lipotoxicity in the liver. Methods: Mice fed a high-fat diet with/without sucrose (HFDS/HFD) and Huh-7 cells 8 9 treated with palmitate were used. Results: Mice fed HFDS showed advanced 10 liver fibrosis as compared to mice fed HFD or normal chow. LPCAT3 mRNA expression in the liver was significantly decreased in the HFDS liver, and LPC 11 content in the HFDS liver was significantly increased as compared to the other 12groups. When Huh-7 cells with shRNA-mediated knockdown of LPCAT3 (shLPCAT3 13 cells) were treated with palmitate, the intracellular LPC concentration and 14 15 cell death were significantly higher than those in wild-type Huh-7 cells. Palmitate-induced cell death in shLPCAT3 was attenuated by a combination of 16 17receptor-interactive protein kinase 1 inhibitor with pan-caspase inhibitor. In LPC 18 contrast, intracellular and palmitate-induced cell death were 19 significantly lower in LPCAT3-overexpressing Huh-7 cells than in wild-type 20 cells. Conclusion: Depletion of LPCAT3 in a mouse model of NASH leads to

caspase-independent cell death, and LPCAT3 is a potential therapeutic target
 in NASH.

1 Introduction

2 Non-alcoholic fatty liver disease (NAFLD), which is a manifestation of the metabolic syndrome in the liver, is highly prevalent in western countries. 3 Non-alcoholic steatohepatitis (NASH) is a subset of NAFLD that is accompanied 4 by inflammation and may lead to cirrhosis ^{1, 2}. Serum free fatty acids (FFA) 5 6 are higher in patients with NASH than in healthy subjects ³. Furthermore, 7 saturated FFA, such as palmitate (PA) can induce apoptosis in hepatocytes ⁴⁻⁷. 8 Thus, toxic lipids, such as saturated FFA, are one of the suspected causes of 9 NASH.

10 Lysophosphatidylcholine (LPC), a metabolite of FFA, is a major phospholipid generated from phosphatidylcholine (PC) by phospholipase A2 11 12 (PLA2)⁸. Inhibition of PLA2 decreases the concentration of LPC in hepatocytes 13and attenuates FFA-induced lipoapoptosis ⁹. In a rat model of NAFLD, serum alanine aminotransferase (ALT) levels were increased in correlation with serum 14 LPC levels ¹⁰. LPC directly induces apoptosis via a pathway that is largely 15 indistinguishable from saturated FFA-induced lipoapoptosis ^{9, 11}. Therefore, LPC 16 is considered as a casual substrate for lipotoxicity in the liver. 17

18 Shindou et al. identified lysophosphatidylcholine acyltransferase 19 (LPCAT) as an enzyme that catalyzes the esterification of LPC ¹². LPCAT 20 regulates cell-membrane glycerolipid remodeling in the so-called "<u>Lands</u> 21 <u>cycle.</u>" LPCAT is classified into four subtypes, LPCAT1-4, on the basis of the

phospholipid substrate ¹². According to its enzymatic activity, LPCAT can
 decrease FFA-induced LPC generation. However, the influence of LPCAT in
 lipoapoptosis remains unclear.

The present study aimed to investigate whether and how LPCAT affects bipotoxicity in the liver. To this end, we employed a NASH mouse model for *invivo* study and a hepatoma cell line (Huh-7) for *in-vitro* study.

1 EXPERIMENTAL PROCEDURES

2 Animals. Male 4-week-old C57BL/6J mice were obtained from Charles River 3 Laboratories (Charles River, Yokohama, Japan) and were maintained on a 12-h light/12-dark cycle in humidity-controlled rooms at 22 ° C with ad libitum 4 access to drinking water. After 1 week of habituation, 10 mice each were 5 assigned to normal chow (Cont), high-fat diet (HFD-60; Oriental Yeast Co., 6 7 Japan) alone (HFD), and HFD with sucrose supplementation (42 g/L) Tokvo. 8 groups (HFDS) and were fed their respective diets for 16 weeks. Three mice of 9 each group were examined by intraperitoneal glucose tolerant test (IPGTT). The 10 remaining seven mice of each group were sacrificed using isoflurane anesthesia 11 after overnight fasting at 21 weeks of age. Three of the seven mice of each 12 group were analyzed for liver LPC content. All of the animal experiments were 13approved by the Animal Care and Use Committee of Iwate Medical University Japan; 28-001). Liver samples were subjected hematoxylin-eosin 14 (Morioka, and Masson-Goldner for fibrosis 15staining for steatosis staining for 16 histological evaluation. Serum aspartate aminotransferase (AST) and total 17cholesterol (TC) were measured with an autoanalyzer (JCA-BM2250; JEOL, Tokyo, 18 Japan).

19 *Cells.* Huh-7 cells, a human hepatocellular carcinoma cell line, were 20 maintained in Dulbecco's modified Eagle's medium containing glucose (25 mM) 21 supplemented with 10% fetal bovine serum, 100,000 IU/L penicillin, and 100

mg/L streptomycin. In addition, we employed Huh-7 cells that stably expressed 1 2 a short-hairpin RNA (shRNA) complementary to LPCAT3 (shLPCAT3 cells) and 3 LPCAT3 overexpression (LPCAT3 OE) cells generated from Huh-7 cells by using lentivirus vector (RC209485L2; OriGene, Rockville, MD). The cells were treated 4 with palmitate (PA) (#P5585; Sigma-Aldrich, St. Louis, MO). PA was dissolved 5 6 in isopropanol at a stock concentration of 160 mM. The final concentration of 7 PA was $\leq 0.2\%$ in the medium, and the corresponding isopropanol concentration 8 was used as a vehicle control (Veh).

9 *Measurement of the <u>phospholipid</u> concentration in cells or the liver.* 10 Cellular LPC in *in-vitro* experiments was measured by an enzymatic assay as 11 reported by Kishimoto et *al.* ¹³. We previously reported details on this assay 12 for *in-vitro* setting ⁹.

13 Total lipids of the liver were extracted by the method of Bligh and Dyer ¹⁴. Phospholipids were separated from total lipids using a diethylaminoethyl-14 15 cellulose column. Liquid chromatography-electrospray ionization-tandem mass 16 spectrometry was carried out using a TSQ-Vantage (Thermo Fisher Scientific, 17Waltham, MA, USA) with an UltiMate 3000 LC system (Thermo Fisher Scientific) 18 equipped with an HTC PAL autosampler (CTC Analytics, Zwingen, Switzerland). 19 LPC and PC were measured by selected reaction monitoring in the negative ion 20 mode.

1 Statistical analysis. All data represent at least three independent 2 experiments and are expressed as the mean \pm SD. Differences between groups 3 were compared using Student's *t*-test and one-way analysis of variance with a 4 *post-hoc* Dunnett test. Significance was accepted at p < 0.05.

5 Materials and methods for intraperitoneal glucose tolerant test, 6 quantitation of cell death, quantitative real-time PCR, immunoblot analysis, 7 XBP1 splicing analysis, and antibodies and reagents are provided in the 8 Supplementary Information.

1 **RESULTS**

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3

HFD induces fatty liver and insulin resistance in mice, and sucrose supplementation enhances HFD-induced inflammation and fibrosis in the liver

To confirm the effect of HFD and sucrose supplementation in mice, we 4 evaluated histological findings of the liver and laboratory data, and we 5 6 monitored body weight. Body weight was significantly higher in HFD and HFDS 7 than in Cont animals (Figure 1A). AST and TC were significantly higher in HFD 8 and HFDS than in Cont (Figure 1B and 1C). Furthermore, HFD and HFDS showed 9 prolonged glucose elevation after intraperitoneal administration of glucose 10 (Figure 1D). In histological evaluation, HFD and HFDS revealed overt lipid accumulation in the liver (Figure 1E). Furthermore, pericellular fibrosis and 11 12 lymphocyte accumulation were observed in the HFDS liver (Figure 1E). mRNA 13expression of alpha smooth muscle actin and collagen I and collagen III was significantly higher in the HFDS than in the HFD and Cont livers (Supplemental 14 Figure 1A-C). These data revealed that HFDS in the present study induced 15 steatohepatitis with insulin resistance in mice, 16 which were therefore 17considered a NASH model. Since LPC is considered as a causal substance for 18 liver injury thorough hepatocyte apoptosis, we evaluated the LPC content in the liver in each group. The liver LPC concentration was significantly higher 19 20 in HFDS than in HFD and Cont (Figure 1F). In addition, the level of PC, a

1 source of LPC, was also elevated in HFDS and HFD compared to Cont 2 animals (Figure 1F).

3

LPCAT3 expression is decreased in mice fed HFDS

As the LPC content in the liver of NASH mice was increased (Figure 1F), 4 we reasoned that LPCAT family activity would be altered in these mice. 5 Therefore, we evaluated the expression of LPCATs in the liver among Cont, HFD, 6 7 and HFDS animals (Figure 1G). Although the expression of LPCAT1 and LPCAT2 8 tended to be decreased in HFD and HFDS as compared to Cont, the difference was 9 not significant. In contrast, LPCAT3 expression was significantly lower in the 10 HFDS liver than in the Cont and the HFD liver. LPCAT4 expression was lower in the HFDS liver than in the Cont, but not the HFD liver. On the basis of these 11 12 data, we hypothesized that the decrease in LPCAT3 expression was associated 13with NASH progression. Therefore, we focused on the effect of LPCAT3 in lipotoxicity. 14

15 Knockdown of LPCAT3 increases PA-induced cell death, which is mediated 16 by LPC

To evaluate the influence of LPCAT3 in lipotoxicity, we used PA, which is known as a toxic lipid. By using shRNA-mediated RNA interference, we knocked down LPCAT3 expression in Huh7 cells (Supplemental Figure 2A). As anticipated, PA cytotoxicity was enhanced in shLPCAT3 as compared to wild-type (WT) Huh7 cells as indicated by a biochemical assay (Figure 2A) and

morphological assessment (Figure 2B). To investigate whether PA-induced cell 1 2 death in shLPCAT3 was mediated by the generation of LPC, we used 3 pharmacological phospholipase A2 inhibitors, bromoenol lactone and palmityl 4 trifluoromethyl ketone. The two inhibitors significantly reduced PA-induced 5 cell death in shLPCAT3 cells as well as the PA-induced increase LPC content 6 (Figure 2C-E). These data indicated that the enhanced cytotoxicity of PA in 7 shLPCAT3 was mediated by an increase in the LPC level due to knockdown of 8 LPCAT3.

9 Endoplasmic reticulum stress and JNK phosphorylation do not show 10 significant differences between shLPCAT3 and WT during lipotoxic insult

11 We next evaluated in detail how lipotoxicity in shLPCAT3 cells induced 12 cell death. To this end, we evaluated several signaling molecules that were signals during lipoapoptosis^{3, 15, 16}. 13previously reported as key TNK phosphorylation and the ER stress response, assessed as eIF2a phosphorylation 14 and XBP1 splicing, respectively, were increased by incubation with PA in both 1516 WT and shLPCAT3 cells; however, all these signals did not show a significant 17difference between the WT and shLPCAT3 cells (Figure 2F and 2G).

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PA-induced cell death in shLPCAT3 cells is mediated by both caspasedependent and -independent mechanisms

20 Because PA-induced cell death was considered as caspase-dependent 21 apoptosis, we checked whether the pan-caspase inhibitor QVD-OPh was able to

prevent PA-induced cell death in shLPCAT3s. While QVD-OPh prevented PA-induced 1 cell death in both WT and shLPCAT3 cells, its effect in shLPCAT3 cells was 2 3 lower than that in WT cells (Figure 3A and 3B). We next evaluated the effect of the RIPK1 inhibitor necrostatin because necroptosis was suggested as 4 another cause of hepatocyte death in NASH¹⁷⁻¹⁹, and RIPK1 is known as a key 5 6 molecule in necroptosis. The combination of necrostatin and QVD-OPh 7 significantly reduced PA-induced cell death in shLPCAT3 cells (Figure 3A and 8 3B).

9 LPCAT3 overexpression decreases PA-induced CHOP expression and reduces 10 PA-induced cell death

11 To confirm the protective effect of LPCAT3 during lipotoxicity, we generated LPCAT3-overexpressing Huh7 cells (LPCAT30E). Overexpression of 12LPCAT3 was confirmed at the mRNA level, and clones 1 and 2 significantly 13 overexpressed LPCAT3 when compared with WT cells (Supplemental Figure 2B). PA-14 induced expression of CHOP, which is a transcriptional factor that serves as 15 16 an ER stress marker, was significantly decreased in LPCAT30E cells (Figure 4A). 17Moreover, phosphorylation of eIF2 and PA-induced cell death were significantly decreased in LPCAT30E compared to WT cells (Figure 4B). Finally, the PA-18 19 induced increase in intracellular LPC was significantly attenuated in LPCAT30E 20 as compared to WT cells (Figure 4C).

21

1 DISCUSSION

2 NASH is expected to become primary indication for liver а 3 transplantation due to hepatocellular carcinoma, liver cirrhosis, and liver failure ^{20, 21}. Because inflammation due to hepatocyte death promotes fibrosis 4 5 in the NASH liver, lipotoxicity-induced hepatocyte death needs to be eliminated. To aid in the establishment of therapeutic targets on the basis of 6 7 the pathophysiology of lipotoxicity-induced hepatocyte death, we investigated 8 the effect of LPCAT3 during lipotoxicity in the liver. The present study 9 revealed that (1) LPCAT3 expression in the liver was decreased in a HDF-10 induced mouse model of NASH, (2) depletion of LPCAT3 enhanced PA-induced 11 hepatocyte death, (3) PA-induced cell death under LPCAT3 depletion was executed by caspase-independent machinery, and (4) LPCAT3 overexpression 1213 decreased PA-induced CHOP expression and attenuated PA-induced cell death. These results indicate that LPCAT3 might serve as a therapeutic target in NASH 14 through a decrease in hepatocyte death. 15

In the present study, LPC was elevated and LPCAT3 expression was downregulated in the livers of NASH model mice. In an *in-vitro* study, the PAinduced increase in intracellular LPC and, accordingly, in hepatocyte death, was enhanced in LPCAT3 knockdown- and attenuated in LPCAT3-overexpressing cells. These data indicated that LPCAT3 protected against lipotoxicity and that depletion of LPCAT3 might be associated with the pathophysiology of NASH.

1	However, how LPCAT3 is suppressed in mice on HFDS remains uncertain. We
2	speculated that a disturbance of the phospholipid metabolism might affect
3	LPCAT3 expression. In NASH patients, PC levels in both the liver and the
4	plasma are significantly higher than those in healthy subjects. Similarly,
5	this study showed that PC levels in both HFD and HFDS mice were higher than
6	those in control group (Figure 1F). In addition, phospholipase A2 inhibitor
7	decreased the PA-induced LPC elevation (Figure 2E). Thus, PC levels in NASH
8	are high, although PC is normally converted to LPC by phospholipase A2. LPCAT3,
9	as a reverse pathway, converts PC to LPC. Thus, we hypothesize that the excess
10	PC might suppress the LPCAT3 pathway. However, LPCAT3 has enzymatic activity
11	toward other phospholipids, such as phosphatidylserine and
12	phosphatidylethanolamine, and thus, these phospholipids might affect LPCAT3
13	expression, although the association between these phospholipids and LPCAT3
14	expression in this study remains unclear. Thus, our hypothesis requires further study.
15	RIPKs, which constitute a family of seven members, are crucial
16	regulators of cell survival and death. Interaction between RIPK1 and RIPK3 is
17	important for necroptosis 22 . Necroptosis occurs when caspase 8 is inactivated
18	on the extrinsic cell death pathway. During necroptosis, RIPK1 and RIPK3 form
19	a complex that is involved in organelle swelling and rupture ²² . As a result,
20	necroptosis promotes inflammation by regulating the release of intracellular
21	damaged-associated molecular patterns ^{23, 24} . A previous study showed that a

RIPK1 inhibitor attenuated inflammation of the NASH liver in vivo ¹⁹. Thus, 1 2 necroptosis was reasoned to be one of the causes of hepatocyte death in NASH 3 ¹⁹. However, how necroptosis was induced in the NASH liver remained unclear. Obviously, the caspase-dependent pathway of lipoapoptosis is the main form of 4 hepatocyte death in the NASH liver ²⁵. However, lipid-induced cell death in 5 6 LPCAT3-depleted hepatocytes was considered as necroptosis because it was 7 attenuated by a combination of pan-caspase inhibitor and RIPK1 inhibitor. 8 Because hepatocyte death in this study was associated with caspase-dependent 9 as well as -independent pathways, LPCAT3-depleted hepatocytes showed lipid-10 induced cell death heterogeneity. The present study newly suggests that 11 impaired phospholipid metabolism induces necroptosis upon abundant execution 12of apoptosis due to lipotoxic insult.

We recognize several limitations of the present study. First, 13 the functions of other LPCATs remain unclear. The expression of all LPCATs tended 14 to decrease in the HFD as well as the HFDS liver, although these changes were 15not significant. We did not exclude whether or not these changes would be 16 17compensative. In the present study, we only demonstrated that LPCAT3 might be a potential therapeutic target of NASH. Second, heterogeneous cell death was 18 19 not elucidated in the mouse model. Since the *in-vivo* experiment in the present 20 study was an observational study based on histological findings, two major 21types of cell death, apoptosis and necroptosis, were not distinguishable.

Third, LPCAT3 expression in the NASH liver was not confirmed. We assessed mRNA 1 2 LPCAT3 expression in the liver by qRT-PCR; however, an antibody to detect the 3 protein is currently not available. Finally, the mechanism of necroptosis in LPCAT3-depleted hepatocytes remains unclear. Although LPCAT3 depletion may 4 lead to two types of cell death during lipotoxic insults, there was no 5 significant difference in endoplasmic reticulum stress and JNK phosphorylation 6 7 between WT and shLPCAT3 cells. Thus, how the disturbance of lipid metabolism due to LPCAT3 depletion induces the two types of cell death is not fully 8 9 understood.

10 In conclusion, the present study led to the following significant findings: LPCAT3 is depleted in the NASH mouse model and leads to caspase-11 dependent/-independent cell death, and LPCAT3 attenuates lipotoxicity by 12decreasing the intracellular LPC content. Based on these findings, we conclude 13 that LPCAT3 is a potential therapeutic target for NASH. Further study on the 14 15 detailed characteristics of LPCAT3-overexpressing hepatocytes is needed because LPCAT3 will affect phospholipid metabolism and cellular membrane 16 17homeostasis.

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ACKNOWLEDGEMENTS

2 This work was supported by KAKENHI Grant Number JP18K07980 (to K.K.) and 3 TaNeDS grant program from Daiichi Sankyo Co., Ltd. (to K.K.). We thank Gregory 4 J. Gores for mentorship, Steve F. Bronk for generation of LPCAT3 knockdown 5 Huh-7 cells, Hiroki Nakanishi (Akita Lipid Technology) for measurement of 6 phospholipids in the liver.

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8 CONFLICTS OF INTEREST

9 The authors declare that they do not have conflict of interest with 10 respect to this manuscript.

1 FIGURE LEGENDS

2 Figure 1. Mice fed high-fat diet with/without sucrose show overweight, 3 impaired glucose tolerance, high liver enzyme content, and hypercholesteremia, and HFDS induces inflammation and fibrosis in the liver and decreases liver 4 5 LPCAT3 mRNA expression. A, Time course of changes in body weight for mice fed normal chow (Cont), high-fat diet (HFD) and high fat diet with sucrose 6 7 (HFDS). *B* and *C*, Serum aspartate aminotransferase (AST; *B*) and total 8 cholesterol (TC; C) levels at 16 weeks of feeding. The serum concentration of 9 the indicated parameter is shown on the vertical axis, treatment groups on the 10 horizontal axis. D, Intraperitoneal glucose tolerance test (IPGTT) results. 11 Each mouse received 2 g/kg of glucose intraperitoneally. The plasma glucose 12 concentration and time points after glucose administration are indicated on 13the vertical and the horizontal axis, respectively. Ε, Representative microscopic images of liver sections subjected to hematoxylin and eosin 14 staining (upper panels, magnification, $50 \times$) and Masson's trichrome staining 15 16 (lower panels, magnification, 50×) for histological evaluation. *F*, 17Lysophosphatidylcholine (LPC) and phosphatidylcholine (PC) levels of the liver 18 were measured by liquid chromatography-electrospray ionization-tandem mass 19 G, mRNA expression of the indicated genes as analyzed by qRTspectrometry. 20 PCR. Gene expression relative to that of GAPDH is indicated on the vertical

1 axis, treatment groups on the horizontal axis. Data are the mean \pm SD from 2 seven experiments; *P<0.05.

3 Figure 2. LPCAT3 knockdown enhances lipotoxicity by increasing LPC production in hepatocytes, but lipotoxicity is not enhanced by JNK, eIF2a, 4 5 and endoplasmic reticulum stress. A and B, Wild-type Huh-7 cells (WT) and 6 LPCAT3 knockdown Huh-7 cells (shLPCAT3) were incubated with palmitate (PA; 800 7 μ M) for 6 h. Vehicle-treated cells (Veh) were used as controls. Caspase 3/78 catalytic activity was assessed by a fluorogenic assay and expressed as fold 9 change (A). Cell death was assessed by fluorescence positivity after propionic 10 iodine staining (B). C and D, Huh-7 cells were incubated with PA (800 μ M) in 11 the presence of a phospholipase A2 inhibitors (20 μM bromoenol lactone [BEL] 12and 60 μ M palmityl trifluoromethyl ketone [PACOCF3]) for 6 h. Caspase 3/713 catalytic activity was assessed by a fluorogenic assay and expressed as fold 14 change (\mathcal{C}). Cell death was assessed by fluorescence positivity after propionic iodine staining (D). E, Intracellular lysophosphatidylcholine (LPC) levels 15 16 were measured by an enzyme-linked colorimetric assay. PA induced an increase 17in LPC above basal levels. WT and shLPCAT3 cells were treated with PA at 400 18 or 800 μ M for 6 h. These cells were also incubated with/without 20 μ M BEL and 19 60μ M PACOCF3 for 6 h. F, Immunoblot analysis of phosphorylated JNK, total 20 JNK, phosphorylated eIF2a and total eIF2a. Whole cell lysates were prepared from WT cells and clones 1 and 2 of shLPCAT3 cells. The cells were incubated 21

with PA (800 µM) for 6 h. <u>The data (phosphorylated-to-total JNK ratio) for</u> <u>each treatment as normalized to it's own Veh control was shown in the lower</u> <u>graph.</u> G: WT and shLPCAT3 cells were treated with either Veh or 800 µM PA for 6 h. XBP-1 cDNA was amplified by PCR followed by 1 h incubation with *Pst*I. 5 Unspliced and spliced forms are presented. <u>Data are the mean ± SD from four</u> 6 <u>experiments; *PK0.05.</u>

7 Figure 3. PA-induced cell death in LPCAT3 knockdown Huh-7 cells is 8 ameliorated by a combination of pan-caspase inhibitor and RIPK1 inhibitor. A 9 and B, Huh-7 cells (WT) and LPCAT3 knockdown Huh-7 cells (shLPCAT3) were 10 incubated with PA (800 μ M) in the presence of QVD-OPh (20 μ M) alone, necrostatine (100 μ M) alone or combination of these agents for 6 h. Cells 11 treated with vehicle (Veh) were used as controls. Caspase 3/7 catalytic 1213activity was assessed by a fluorogenic assay and expressed as fold change (A). Cell death was assessed by fluorescence positivity after propionic iodine 14 staining (B). Data are the mean \pm SD from four experiments; *P(0.05. 15

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Figure 4. LPCAT3 overexpression ameliorates PA-induced cell death by suppressing the increase in intracellular LPC and CHOP expression. *A*, *B* and *C*, Huh-7 cells (WT) and LPCAT3-overexpressing Huh-7 cells (LPCAT30E) were incubated with PA (800 μM) for the indicated time. Vehicle (Veh)-treated

1 cells were used as controls. Immunoblot analysis of CHOP. Whole cell lysates 2 were prepared from WT cells and clones 1 and 2 of LPCAT30E cells. The cells were incubated with PA (800 $\mu\text{M})$ for 16 h (A). Cell death was assessed by 3 4 fluorescence positivity after propionic iodine staining. WT and LPCAT30E cells 5 were treated with PA for 16 h (B). Intracellular LPC levels were measured by 6 an enzyme-linked colorimetric assay. PA induced an increase in LPC above basal 7 levels in WT cells treated with Veh. WT and LPCAT30E cells were treated with PA for 8 h (C). All data in (B) and (C) are expressed as the mean \pm SD from 8 9 three experiments; **P*<0.05.

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Figure 2



Figure 3



Figure 4



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Supplemental material and method

Intraperitoneal glucose tolerant test. The mice were administrated glucose (2 g/kg body weight) intraperitoneally after 6 h of fasting. Blood samples were obtained from the tail vein before glucose administration and at 15, 30, 60, 90, and 120 min after glucose administration.

Quantitation of cell death. Cell death was evaluated by morphological approaches and biochemical methods. For morphological approaches, nuclei were stained by propidium iodide for 30 min at 37 °C and analyzed by fluorescence microscopy (Nikon Eclipse TE200; Nikon, Tokyo, Japan). The number of dead cells was expressed as a percentage of total cells counted. Cell death was also examined by spectrophotometry using Cell Count Reagent SF (Catalog No. 07553; Nakalai Tesque Inc.). The results were presented as the ratio of absorption in the vehicle control group to that in the treatment group. For biochemical methods, we evaluated caspase activity using the Apo-One Caspase-3/7 assay (G7790; Promega, Madison, WI), according to the manufacturer's protocol.

Quantitative reverse transcription (qRT)PCR. Total cellular RNA was extracted using Trizol reagent (Invitrogen, Camarillo, CA, USA) and was reverse-transcribed into cDNA with Moloney murine leukemia virus reverse transcriptase (Invitrogen,

Camarillo, CA, USA) and random primers (Invitrogen, Camarillo, CA, USA) as previously described ¹⁵. The cDNA was used as a template for qRT-PCR, which was carried out on a LightCycler instrument (Roche Applied Science) using SYBR green (Molecular Probes) as a fluorophore. PCR primers were as follows: for mouse LPCAT1 (NM_145376.5): forward 5'-ccctgggacctcctgataa-3' and reverse 5'-gcaggaagtccacgacctt-3' (69 bp), for mouse LPCAT2 (NM_173014.1): forward 5'-tgtactaatcgctcctgtttgatt-3' and reverse 5'-cactggaactcctgggatg-3' (63 bp), for mouse LPCAT3 (NM_145130.2): forward 5'-ggcctctcaattgcttatttca-3' and reverse 5'-agcacgacacatagcaagga-3' (65 bp), for mouse LPCAT4 (NM_207206.2): forward 5'-ggcctccagagggttaagtt-3' and reverse 5'-aaaagctagaagtactcggattgg-3' (69 bp), for mouse GAPDH (NM_001289726.1): forward 5'-gggttcctataaatacggactgc-3' and reverse 5'-ccattttgtctacgggacga-3' (112 bp), for human LPCAT3 (NM 005768.5): forward 5'-accaggaaagataccaaacagc-3' and reverse 5'-ggtagaaaaggcccagactca-3' (65 bp), for human GAPDH (NM_002046.5): forward 5'agccacatcgctcagacac-3' and reverse 5'-gcccaatacgaccaaatcc-3' (66 bp). GAPDH was used as an internal control. Gene expression was quantified by the $2^{-\Delta\Delta CT}$ method, and the target mRNA expression levels were expressed relative to GAPDH per sample as previously described ¹⁵.

Immunoblot analysis. Whole cell lysates were prepared as previously

described ¹⁵. Equal amounts of protein (20 to 80 µg) were resolved by SDS-PAGE on a 12.5–15% acrylamide gel, transferred to a nitrocellulose membrane, and incubated with primary antisera. Membranes were incubated with appropriate horseradish peroxidase-conjugated secondary antibodies (Santa Cruz Biotechnology, Santa Cruz, CA). Bound antibody complexes were visualized using chemiluminescent substrate (ECL; Amersham, Arlington Heights, IL) and exposure to Kodak X-OMAT film (Eastman Kodak, Rochester, NY).

XBP1 splicing analysis. RNA was extracted from shLPCAT3 cells subjected to indicated treatments and transcribed into cDNA. XBP1 was amplified by PCR using forward primer, 5'-AAACAGAGTAGCAGCTCAGACTGC-3', and reverse primer, 5'-TCCTTCTGGGTAGACCTCTGGGAG-3'. The PCR product was digested using the *Pst*I restriction enzyme for 1 h at 37 °C, and then electrophoresed on a 1.7 % agarose gel. The gels were photographed under UV transillumination.

Antibodies and reagents. Antisera used were obtained from the following sources: rabbit anti-phospho-eIF2α (1:1000; #9721), rabbit anti-eIF2α (1:1000; #9722), rabbit anti-phospho-JNK (1:1000; #9251), rabbit anti-JNK (1:1000; #9252, Cell Signaling Technology, Danvers, MA, USA); mouse anti-C/EBP homologous protein (CHOP [1:500; sc-575]), and goat anti-β-actin (1:1000; sc-1616, Santa Cruz Biotechnology, Santa Cruz, CA, USA). The pan-caspase inhibitor Q-Val-Asp-OPh (QVD-OPh) was obtained from MP Biomedicals Japan (Tokyo, Japan). The receptor interacting protein 1 kinase (RIPK1) inhibitor was obtained from Selleckchem (Houston, TX, USA). To determine effect of these pharmacological inhibitors to Huh-7 with PA, the cells were incubated with PA (800 μ M) in the presence of QVD-OPh (20 μ M) alone, necrostatine (100 μ M) alone or combination of these agents for 6 h. Bromoenol lactone (BEL; B1552), palmityl trifluoromethyl ketone (PACOCF₃; P8727), BSA, Bradford reagent, and other chemicals were all obtained from Sigma-Aldrich (St. Louis, MO, USA). To determine effect of each phospholipase A2 inhibitors to Huh-7 with PA, BEL of 20 μ M or PACOCF3 of 60 μ M was incubated with PA for 6 h.

Supplemental Figure Legends

Supplemental Figure 1. Expression of genes associated with liver fibrosis in mice fed normal chow, high-fat diet, and high-fat diet with sucrose. *A*, *B*, and *C*. Total RNA was prepared from the livers of mice fed normal chow (Cont), high-fat diet (HFD), or high-fat diet with sucrose (HFDS). mRNA levels of α SMA (A), collagen I (B), and collagen III (C) were quantified by qRT-PCR, normalized to GAPDH, and expressed as fold change over vehicle. All data are expressed as the mean ± SD from four experiments; **P*<0.05.

Supplemental Figure 2.

A and B, Total RNA was prepared from Huh-7 cells (WT), LPCAT3 knockdown Huh-7 cells (shLPCAT3), and LPCAT3-overexpressing Huh-7 cells (LPCAT3OE). LPCAT3 expression in shLPCAT3 (A) and LPCAT3OE (B) cells was quantified by qPCR, normalized to GAPDH, and expressed as fold change over WT. All data are expressed as the mean \pm SD from four experiments; **P*<0.05.

Supplemental Figure 1



Supplemental Figure 2

