



Limonin, an AMPK Activator, Inhibits Hepatic Lipid Accumulation in High Fat Diet Fed Mice

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NAFLD is the most prevalent liver disease in human history. The treatment is still limited yet. In the current study, we reported that limonin inhibited hepatic lipid accumulation and fatty acid synthesis in HFD fed mice. Using AMPK inhibitor and AMPK deficient *C. elegans*, we revealed the effect was dependent on the activation of AMPK. We found that limonin activated AMPK through inhibition of cellular energy metabolism and increasing ADP:ATP ratio. Furthermore, the treatment of limonin induced AMPK mediated suppression of the transcriptional activity of SREBP1/2. Our study suggests that limonin may a promising therapeutic agent for the treatment of NAFLD.

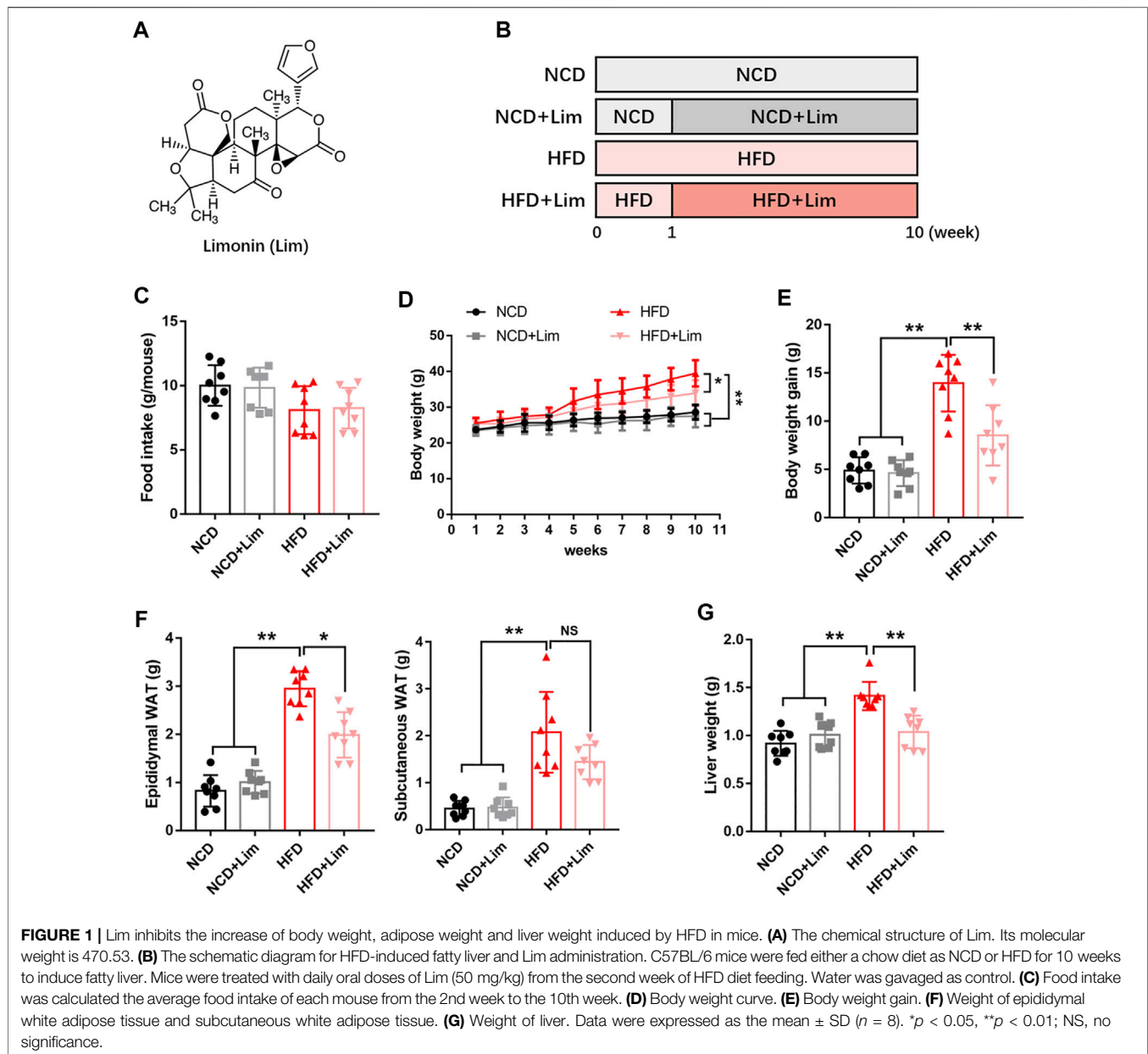
Keywords: limonin, AMPK, lipid accumulation, NAFLD, SREBP

INTRODUCTION

Non-alcoholic fatty liver disease (NAFLD), which is also called metabolic associated fatty liver disease (MAFLD) (Eslam et al., 2020), is characterized by pathological accumulation of triglycerides (TG) and other lipids in hepatocytes (Heeren and Scheja 2021; Ooi et al., 2021). It can progress to nonalcoholic steatohepatitis (NASH) and fibrosis, which eventually lead to liver cirrhosis, hepatocellular carcinoma. Although a variety of small molecule chemical drugs are undergoing clinical trials (Zhu et al., 2021), the treatment for NAFLD is still limited.

AMP-activated protein kinase (AMPK) is a key metabolic regulator that senses energy status and controls energy expenditure and storage, whose activation has been proposed to be therapeutically beneficial for the treatment of NAFLD (Garcia et al., 2019; Zhao et al., 2020). AMPK is activated in response to energy stress by sensing increases in AMP: ATP and ADP: ATP ratios and inhibited by adenosine triphosphate (ATP) (Garcia and Shaw 2017). Liver specific AMPK knockout can aggravate hepatic lipid accumulation, steatosis, inflammation, fibrosis and hepatocyte apoptosis (Garcia et al., 2019; Zhao et al., 2020). Hepatic activation of AMPK by the synthetic polyphenol protects against hepatic steatosis by suppressing sterol regulatory element binding protein 1 (SREBP1) activity (Li et al., 2011). Activating AMPK also inhibit hepatic cholesterol synthesis by suppression of SREBP2 activity (Tang et al., 2016). Our previous study showed that activation of AMPK by flavonoids could ameliorate hepatic steatosis in mice (Wang et al., 2020a).

Limonin (Figure 1A), a tetracyclic triterpenoid compound, is a secondary metabolite with high biological activity in plants (Fan et al., 2019). It is abundant in many traditional Chinese medicines and fruits (Fan et al., 2019). It has been recognized as one of the most beneficial and active components of medicinal foods (Gu et al., 2019). In recent years, pharmacological investigations have uncovered various bioactivities of limonin including anti-cancer (Chidambara Murthy et al., 2021), anti-inflammatory (Yang et al., 2021), anti-oxidation (Yu et al., 2005) and liver protection



activity (Yang et al., 2020). Limonin reduces LDL cholesterol in HepG2 cells (Bhathena and Velasquez 2002) and regulates the expression of genes related to lipid metabolism in mice (Fan et al., 2019). However, the effect of limonin in hepatic lipid metabolism and the mechanism is still unclear.

Here, we reported that limonin inhibited hepatic lipid accumulation and fatty acid synthesis in HFD-induced mice. Further mechanistic studies revealed that limonin suppressed the transcriptional activity of SREBP1/2 by activating AMPK. In general, our study reveals that limonin may be a promising therapeutic agent for the treatment of NAFLD.

MATERIALS AND METHODS

Materials

Limonin (Lim, CAS# 1,180-71-8, HPLC \geq 95%) was purchased from TCI (Shanghai) Development Co., Ltd., China. Compound C (#B3252) was purchased from ApexBio, United States. Antibody sources are as follows: phospho-AMPK α (#2535, Cell Signaling Technology), AMPK α (#5831, Cell Signaling Technology), phospho-ACC (#3661, Cell Signaling Technology), ACC (#3662, Cell Signaling Technology), phospho-LKB1 (#3055, Cell Signaling Technology), LKB1

(#3050, Cell Signaling Technology), phospho-CAMKK2 (#12818, Cell Signaling Technology), CAMKK2 (#16810, Cell Signaling Technology), PP2A (#2038, Cell Signaling Technology), PP2C (#3549, Cell Signaling Technology), phospho-TAK1 (#4508, Cell Signaling Technology), TAK1 (#5206, Cell Signaling Technology), SREBP1 (ab28481, Abcam), SERBP2 (ab30682, Abcam), β -actin (#3700, Cell Signaling Technology). Commercial kits used in measurement of plasma parameters are as follows: triglyceride (TG), total cholesterol (TC) assay kits were purchased from Dongou Diagnostics Co., Ltd., Zhejiang, China; alanine aminotransferase (ALT), aspartate aminotransferase (AST), alkaline phosphatase (ALP), non-esterified fatty acid (NEFA), high-density lipoprotein cholesterol (HDL-C), and low-density lipoprotein cholesterol (LDL-C) assay kits were purchased from Nanjing Jiancheng Bioengineering Institute, China.

Animal Experiments

All animal experiments were approved by the Animal Care and Use Committee of Zhejiang Chinese Medical University, where the experiments were conducted. 32 C57BL/6 male mice (eight-week-old), which were purchased from GemPharmatech Co., Ltd., Jiangsu, China (license number of animal production: SYXK 2015-0001), were housed two per cage in a temperature and humidity-controlled room with a 12:12 h light/dark cycle. After 1 week of acclimation, the mice were randomly separated into four groups: 1) NCD group: mice ($n = 8$) were orally administered with ultrapure water as control vehicle and fed with 10% Kcal high-fat, 7% sucrose control diet match D12492 (Research diet D12450J, Research Diet, NJ); 2) NCD + Lim group: mice ($n = 8$) were orally administered with limonin (Lim, 50 mg/kg/day) and fed with 10% Kcal high-fat, 7% sucrose control diet match D12492 (Research diet D12450J, Research Diet, NJ); 3) HFD group: mice ($n = 8$) were orally administered with ultrapure water as control vehicle and fed with 60% Kcal high-fat diet (Research diet D12492, Research Diet, NJ); 4) HFD + Lim group: mice ($n = 8$) were orally administered with limonin (Lim, 50 mg/kg/day) and fed with 60% Kcal high-fat diet (Research diet D12492, Research Diet, NJ). As described, the Lim administration started from the second week and the experiment lasted for 10 weeks in total (**Figure 1B**). Body weight and food intake were recorded weekly. At the end of the study, the blood samples were collected and the plasma parameters were detected using the indicated kits according to the manufacturers' instructions. The tissues were dissected, weighed, immediately frozen in liquid nitrogen and stored at -80°C .

Oral glucose tolerance tests and insulin tolerance test

OGTT and ITT were conducted at 9 weeks of the experiment. After fasting for 6 h, mice were oral D-glucose (2 g/kg) or i. p. injected with insulin (0.75 U/kg). Blood glucose levels were measured at 0, 30, 60, 90, and 120 min, which was measured by tail vein using a standard glucometer (Johnson & Johnson, United States).

Histology Examination

Mouse liver tissues were fixed in fixed in 4% v/v phosphate-buffered formaldehyde, embedded in paraffin, sectioned, and stained with hematoxylin and eosin (H&E) according to our described previously (Bai et al., 2019; Wang et al. 2020a; Wang et al. 2021a). Lipid droplets were visualized by Oil Red O (Solarbio Life Science, China) staining. The NAFLD activity score (NAS) and the intensity of Oil Red O analysis were also based on the method according to our previous researches (Wang et al., 2020a).

Cell Culture and Treatment

The mouse normal hepatic cell line AML12 (obtained from the Shanghai Bank of Cell Lines) was routinely cultured in Dulbecco's Modified Eagle's Medium (DMEM/F-12) containing 10% fetal bovine serum (FBS). Human hepatoma HepG2 cell line (obtained from the Shanghai Bank of Cell Lines) was cultured in Dulbecco's Modified Eagle's Medium (DMEM) containing 10% fetal bovine serum (FBS). To establish a hepatic lipid accumulation model using AML12 cells, we used 0.4 mM palmitic acid (PA) after starving in serum-free DMEM for 24 h.

Immunofluorescence Staining

To visualize the expression and localization of SREBP1 and SREBP2 in cells, AML12 cells were incubated with anti-SREBP1 (1:100) or anti-SREBP2 (1:100) antibody at 4°C overnight. Then the cells were stained by Alexa Fluor 488 secondary antibody for 1 h at room temperature. The fluorescence was visualized by a SUNNY RX50 fluorescence microscope. An average score of the immunofluorescence was calculated as described previously (Wang et al. 2018; Wang et al. 2019).

C. elegans Strain and Treatment

Wild-type N2 *C. elegans* and aak-2 (ok524) X mutant strains were obtained from the Caenorhabditis Genetics Center. All worms were fed on *Escherichia coli* OP50 lawn and raised at 20°C on nematode growth medium (NGM) agar plates. The yielding eggs were hatched in M9 buffer overnight at 20°C to obtain the age-synchronized L1 worms. The synchronized populations were further incubated on NGM plates which were pretreated with 100 μM Lim (dissolved in M9 buffer) for 6 days.

In vitro Lipid Accumulation

Lipid accumulation in hepatocytes or *C. elegans* was visualized by Oil Red O staining or quantified by commercial kits (TC and TG), according to the manufacturer's instructions.

ADP/ATP Ratio Measurement

Cells were treated with different concentrations of limonin dissolved in DMSO for 2 h. ADP/ATP ratio was measured by ADP/ATP Ratio Assay Kit (#ab65313, Abcam, Burlingame, CA) according to the manufacturer's instruction.

Plasmids and Transfection

AMPK-DN was cloned from pMIGR-AMPK-KD (Addgene, Cat #27296) into pcDNA-3xFlag plasmid. The fragment of AMPK γ 2

TABLE 1 | The primers used in this study for real time PCR.

Description	Sense primer (5'→3')	Antisense primer (5'→3')
<i>Fasn</i>	GGAGGTGGTGATAGCCGGTAT	TGGGTAATCCATAGAGCCCAG
<i>Scd1</i>	TTCTTGCGATACACTCTGGTGC	CGGGATTGAATGTTCTTGTCGT
<i>Acc1</i>	CTCCCGATTGATAATTGGGTCTG	TCGACCTTGTTTTACTAGGTGC
<i>Hmgcr</i>	TGACCTTTCTAGAGCGAGTGCAT	CACGAGCTATATTTCCCTTACTTCA
<i>Hmgcs</i>	AGAGAGCGATGCAGGAAACTT	AAGGATGCCACATCTTTTGG
<i>Gapdh</i>	TGAGGCCGGTGCTGAGTATGT	CAGTCTCTGGGTGGCAGTGAT

was cloned from pGEM-PRKAG2 (#HG16130-G, SinoBiological, Beijing, China) and constructed into pLKO-puro FLAG plasmid. Mutagenesis was performed using the Hieff Mut™ Site-Directed Mutagenesis Kit (YEASEN, Shanghai, China) according to the manufacturer's instruction. The transfection was performed using Lipofectamine 3000 reagent from Life Technologies (Carlsbad, CA) according to our described previously (Wang et al., 2021b).

Quantitative Reverse Transcriptase-Polymerase Chain Reaction

Quantitative real-time PCR was performed as described previously (Wang et al. 2020a; Wang et al. 2020b). In brief, total RNA from tissue or cells were isolated with TRIzol (#DP424, Tiangen Biotech Co. Ltd., Beijing, China). First-strand cDNA was synthesized from 1.5 µg of RNA using reverse transcriptase kits (ThermoFisher Scientific, Waltham, MA) according to the manufacturer's instructions. After cDNA synthesis, the expressions of indicated genes were estimated by real-time PCR using the SGExcel FastSYBR Mixture (#B532955-0005, Sangon Biotech Co., Ltd., Shanghai, China) on Roche LightCycler^R 480 Quantitative PCR System (Indianapolis, United States). The PCR results of GAPDH served as internal controls. The primers used for PCR are listed in **Table 1**.

Immunoblotting

Immunoblotting was performed as described previously (Wang et al., 2020a). In brief, liver tissue or cells were extracted with 1 × sodium dodecyl sulfate (SDS). A total of 25 µg protein was loaded into a 10% SDS-PAGE gel and transferred onto a polyvinylidene fluoride (PVDF) membrane. Primary antibodies were incubated overnight, and the secondary antibodies (Cell Signaling Technology, 1:3000) were added onto the membrane. Immunoreactive bands were visualized using enhanced chemiluminescence reagents (#180-501, Tanon Biotechnology, Shanghai, China). Chemiluminescence was determined using Tanon 4200SF system (Tanon Biotechnology, Shanghai, China).

Statistical Analysis

Data is presented as mean ± SD. Statistical analyses were performed using GraphPad Prism 7.0 (GraphPad Software, La Jolla California USA). Differences between the groups were analyzed using Student's t-test or one-way ANOVA followed by Dunnett's multiple comparisons test. Significance thresholds were $p < 0.05$.

RESULTS

Lim Ameliorates Metabolic Disorder in HFD Fed Mice

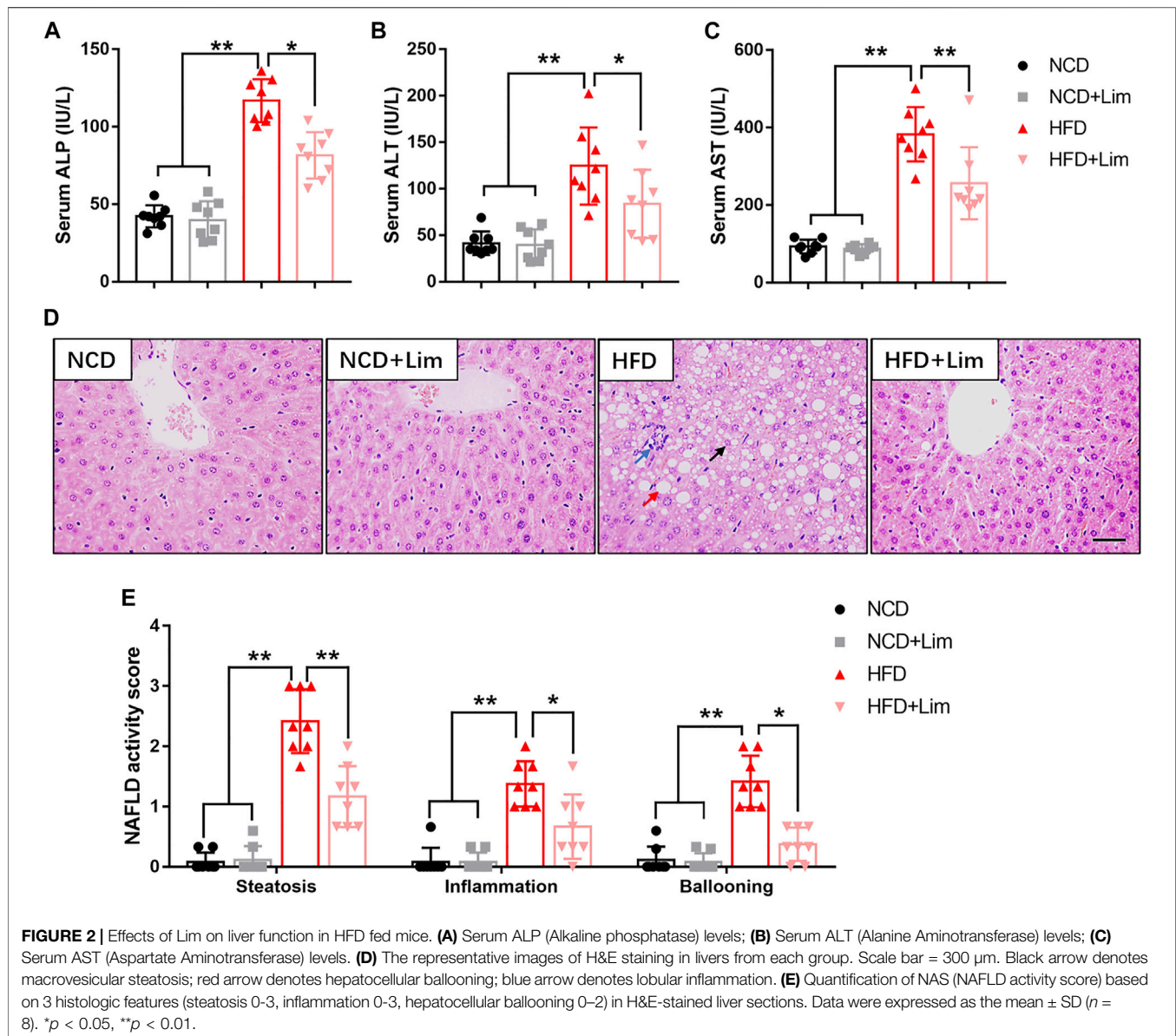
To explore the effect of Lim on metabolic disorder, we built up our animal model by feeding mice high fat diet (**Figure 1B**). Firstly, we took thorough examination of metabolic parameters in these mice. There was no difference in food intake between control and Lim treated groups no matter what kind of diet was fed (**Figure 1C**). We observed significant body weight loss in Lim treated mice fed with HFD, and there was no obvious difference in body weights in mice fed with chow diet with or without Lim administration (**Figures 1D,E**). Lim reduced the amount of epididymal and subcutaneous white adipose tissue in HFD fed mice (**Figure 1F**). Moreover, Lim treatment improved systematic insulin resistance and lowered serum lipids in HFD-fed mice (**Supplementary Figures S1, S2**). Of note, the increasing of liver weight induced by HFD feeding was also ameliorated by Lim treatment (**Figure 1G**).

Lim Improves Liver Function in HFD Fed Mice

Next, we focused on the effect of Lim on liver function under the condition of HFD feeding. As shown in **Figures 2A–C**, Lim significantly improved liver function in HFD fed mice revealed by decreased levels of ALP, ALT and AST in serum. The pathologic change of livers was determined by H&E staining and semi-quantitative analysis. We found that Lim obviously ameliorated hepatic steatosis and the infiltration of inflammatory cells in HFD fed mice (**Figures 2D,E**).

Lim Reduces Hepatic Lipid Accumulation in HFD Fed Mice

To further characterize the effect of Lim on hepatic lipid accumulation, we evaluated the lipid content of mouse liver by several approaches. We found that Lim treatment significantly reduced the content of total cholesterol (TC) and triglyceride (TG) in the livers of mice fed a high-fat diet (**Figures 3A,B**). The amount of hepatic neutral lipids determined by Oil Red O staining was also decreased in HFD induced mice after Lim administration (**Figures 3C,D**). Furthermore, the mRNA expression of genes associated with fatty acid synthesis (*Fasn*, *Scd1* and *Acc1*) and cholesterol synthesis (*Hmgcr* and *Hmgcs*) was



down-regulated by Lim in the livers of HFD fed mice (Figures 3E,F).

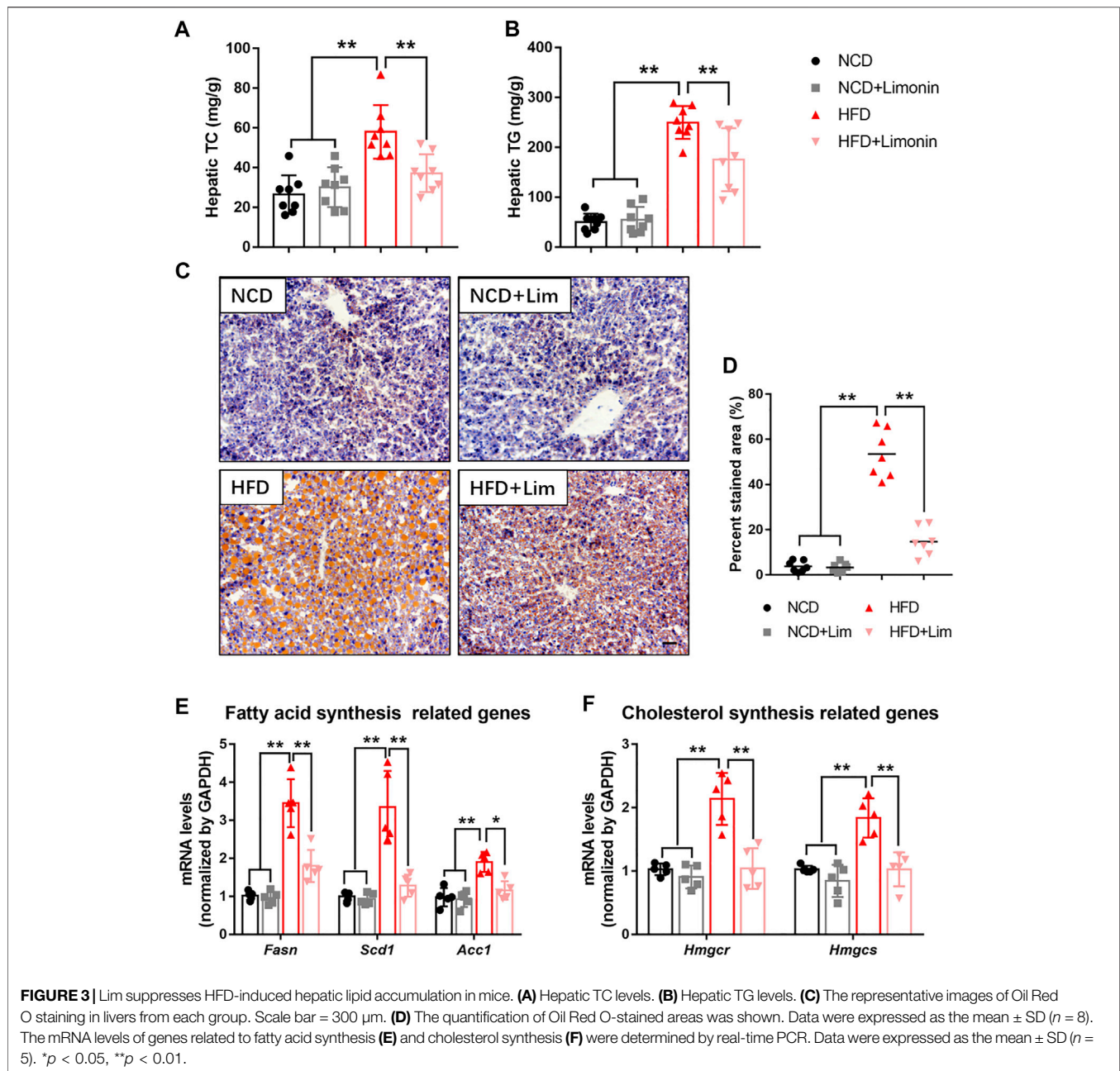
Lim Activates AMPK *in vitro* and *in vivo*

AMPK is a master regulator to keep metabolic balance in liver tissue. Previous study reported that triterpenoids can activate AMPK (Tan et al., 2008). Lim is the first tetranortriterpenoid obtained from citrus bitter principles (Roy and Saraf 2006). Therefore, we wondered if Lim could increase AMPK activity. We found that HFD feeding inhibited AMPK phosphorylation in mouse livers, while Lim treatment greatly enhanced AMPK phosphorylation, especially in HFD mice. Meanwhile, the phosphorylation of ACC, a well-known substrate of AMPK, was also increased by Lim under the condition of HFD feeding (Figures 4A,B). To determine if the activation of AMPK is a direct action of Lim on hepatocytes in the livers,

or due to its indirect effect on whole body metabolism, we treated murine hepatic cell line AML12 with Lim in different concentrations or time. As shown in Figures 4C,D, Lim induced AMPK activation and ACC phosphorylation in a dose- and time-dependent manner. To mimic HFD feeding *in vitro*, we treated AML12 cells with palmitic acid (PA). Similar with the observation *in vivo*, PA challenge decreased the phosphorylation of both AMPK and ACC, while Lim treatment re-elevated both of them in PA challenged cells (Figures 4E,F).

Lim Ameliorates Lipid Accumulation via AMPK Activation

To investigate whether the activation of AMPK is required for the reductive effect of Lim on lipid accumulation, we

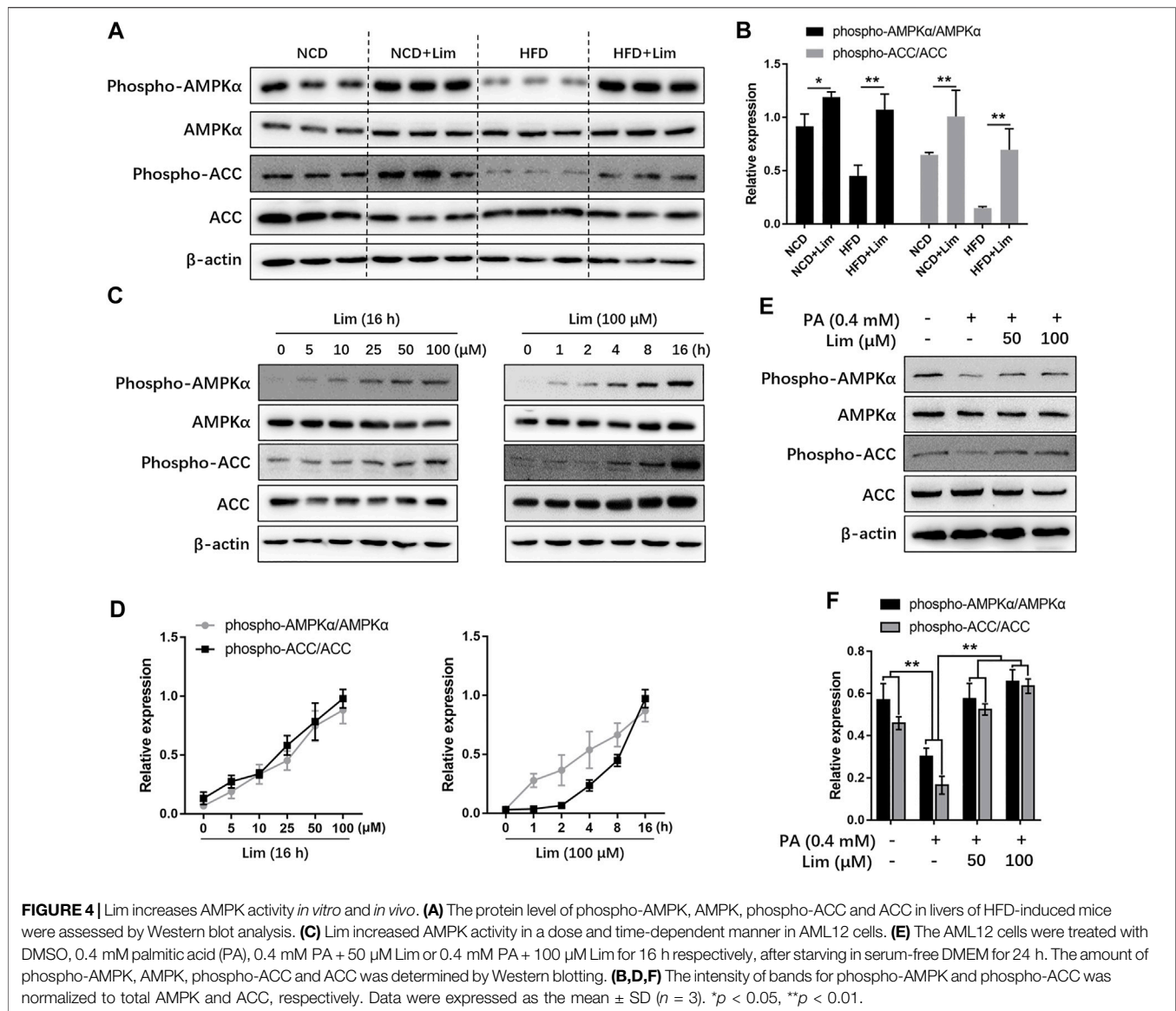


suppressed AMPK activity *in vitro* by Dorsomorphin (Compound C) or the expression of AMPK dominant negative plasmid. As shown in **Figure 5A** and **Supplementary Figure S3**, Lim significantly lowered PA induced lipid accumulation in AML12 cells, while Compound C almost completely abolished the effect. Similar with the result from Oil Red O staining, the measurement of TG and TC also showed that the administration of Compound C reversed Lim induced decrease of intracellular lipids (**Figure 5B**). In addition, Inhibition of AMPK activity by the expression of dominant

negative AMPK plasmid also greatly reversed the repressive effect of Lim on lipid accumulation (**Figures 5C,D**).

Lim Suppresses Fat Accumulation in *Caenorhabditis elegans* Through AMPK

To further confirm AMPK activation is also necessary to Lim induced reduction of lipid accumulation *in vivo*, we used genetic model from *Caenorhabditis elegans*. Lim administration reduced the amount of lipid droplets in N2 worms determined by Oil Red O staining and TC/TG measurement. In *aak-2* worms, knockout



of AMPK completely abrogated Lim induced suppression of fat accumulation (Figures 6A–C).

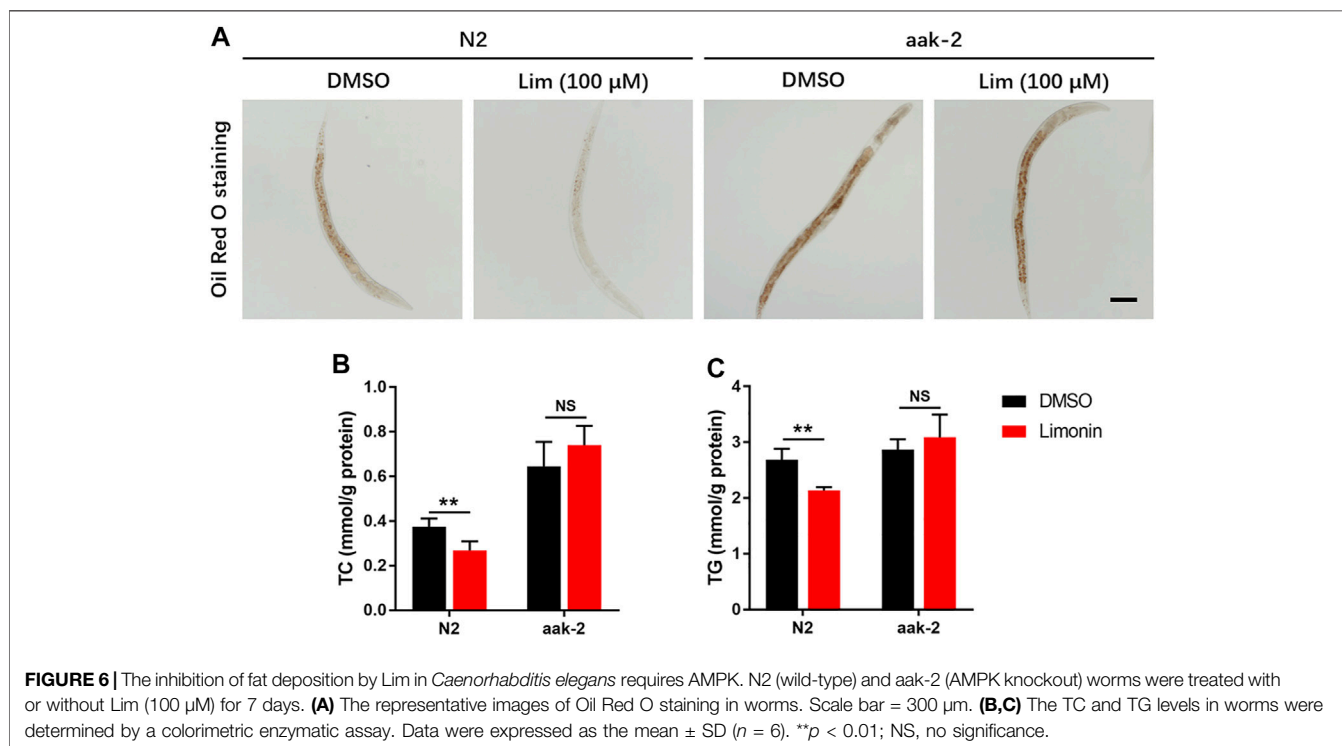
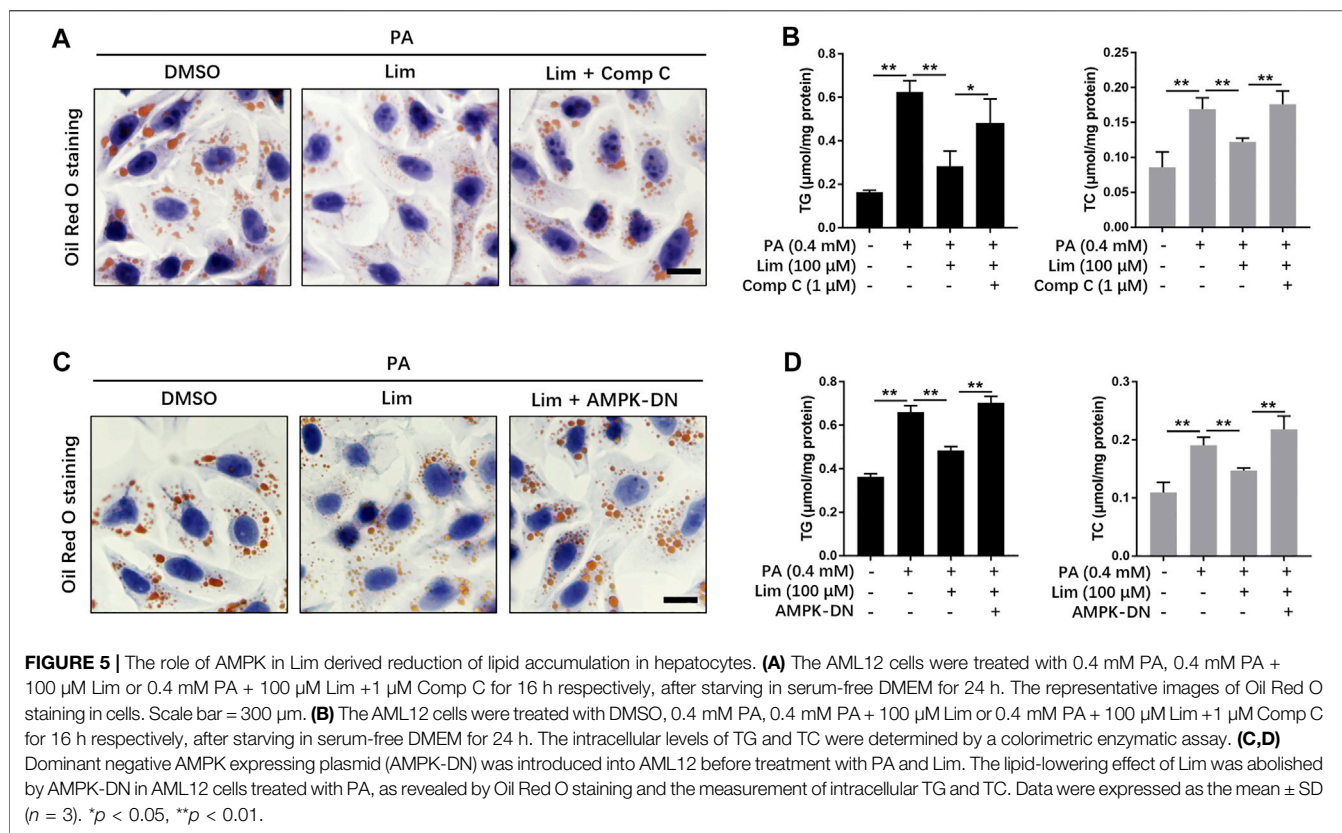
Lim Activates AMPK Through Inhibition of ATP Generation

Next, we tried to find out how Lim activated AMPK. The phosphorylation of Thr172 and the binding of AMP are considered as two major ways to activate AMPK (Herzig and Shaw, 2018). There are several alternative upstream activators and negative regulators in mammals for the regulation of AMPK activity (Jian et al., 2020). We found that neither the upstream kinases, including liver kinase B1 (LKB1), calcium/calmodulin-dependent protein kinase 2 (CAMKK2), and transforming growth factor β activated kinase 1 (TAK1), nor the negative regulators such as protein phosphatase 2A (PP2A) and PP2C

were significantly affected by Lim treatment (Figure 7A). To determine whether AMPK activation by Lim was AMP dependent, we performed the assay firstly introduced by Simon A. Hawley on *Cell Metabolism* (Hawley et al., 2010). As shown in Figures 7B,C, Lim induced AMPK activation and ACC phosphorylation in a dose-manner in WT but not in RG cells, suggesting Lim treatment inhibited ATP production. Indeed, Lim significantly increased ADP/ATP ratio measured by bioluminescent assay kit in parental 293T cells (Figure 7D).

Lim Suppresses the Transcriptional Activity of SREBP1/2 Through AMPK

Since the expression of fatty acid and cholesterol synthesis genes were down-regulated by Lim in the livers of HFD fed mice, we wondered whether Lim inhibited the expression of these genes by



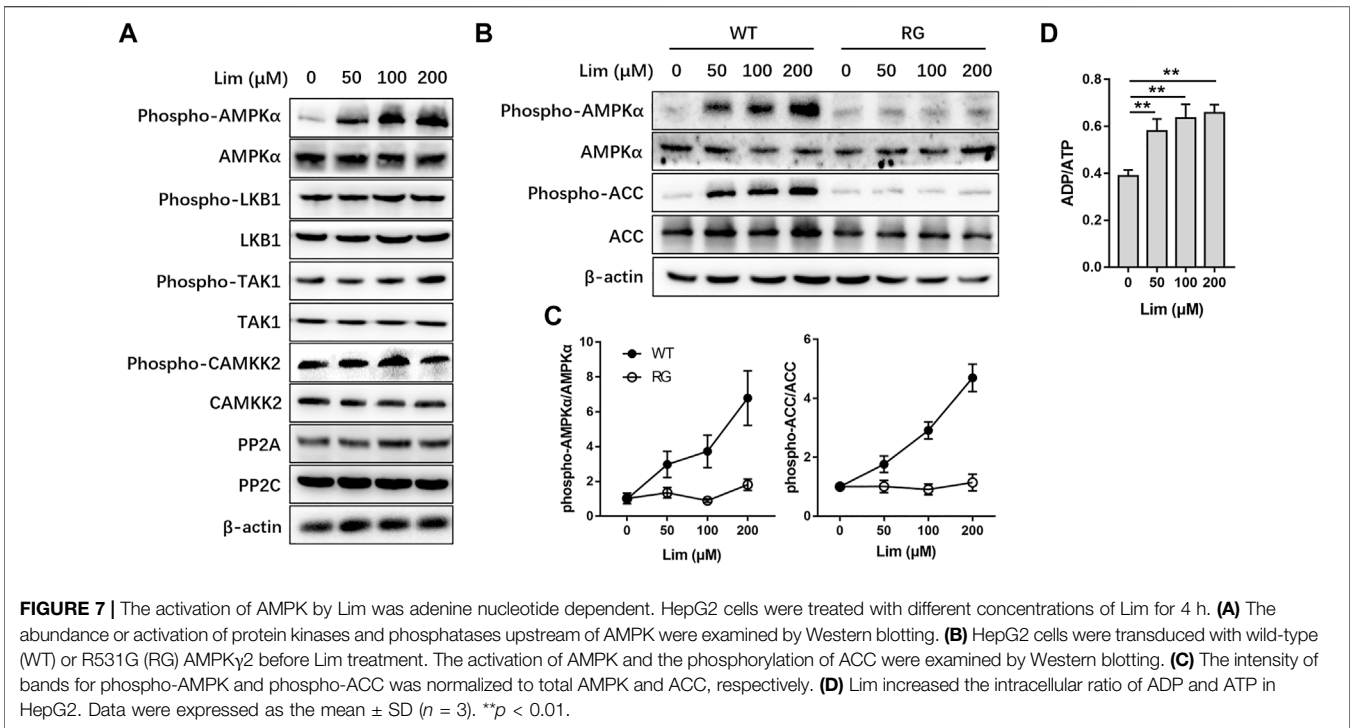


FIGURE 7 | The activation of AMPK by Lim was adenine nucleotide dependent. HepG2 cells were treated with different concentrations of Lim for 4 h. **(A)** The abundance or activation of protein kinases and phosphatases upstream of AMPK were examined by Western blotting. **(B)** HepG2 cells were transduced with wild-type (WT) or R531G (RG) AMPK γ 2 before Lim treatment. The activation of AMPK and the phosphorylation of ACC were examined by Western blotting. **(C)** The intensity of bands for phospho-AMPK and phospho-ACC was normalized to total AMPK and ACC, respectively. **(D)** Lim increased the intracellular ratio of ADP and ATP in HepG2. Data were expressed as the mean \pm SD ($n = 3$). ** $p < 0.01$.

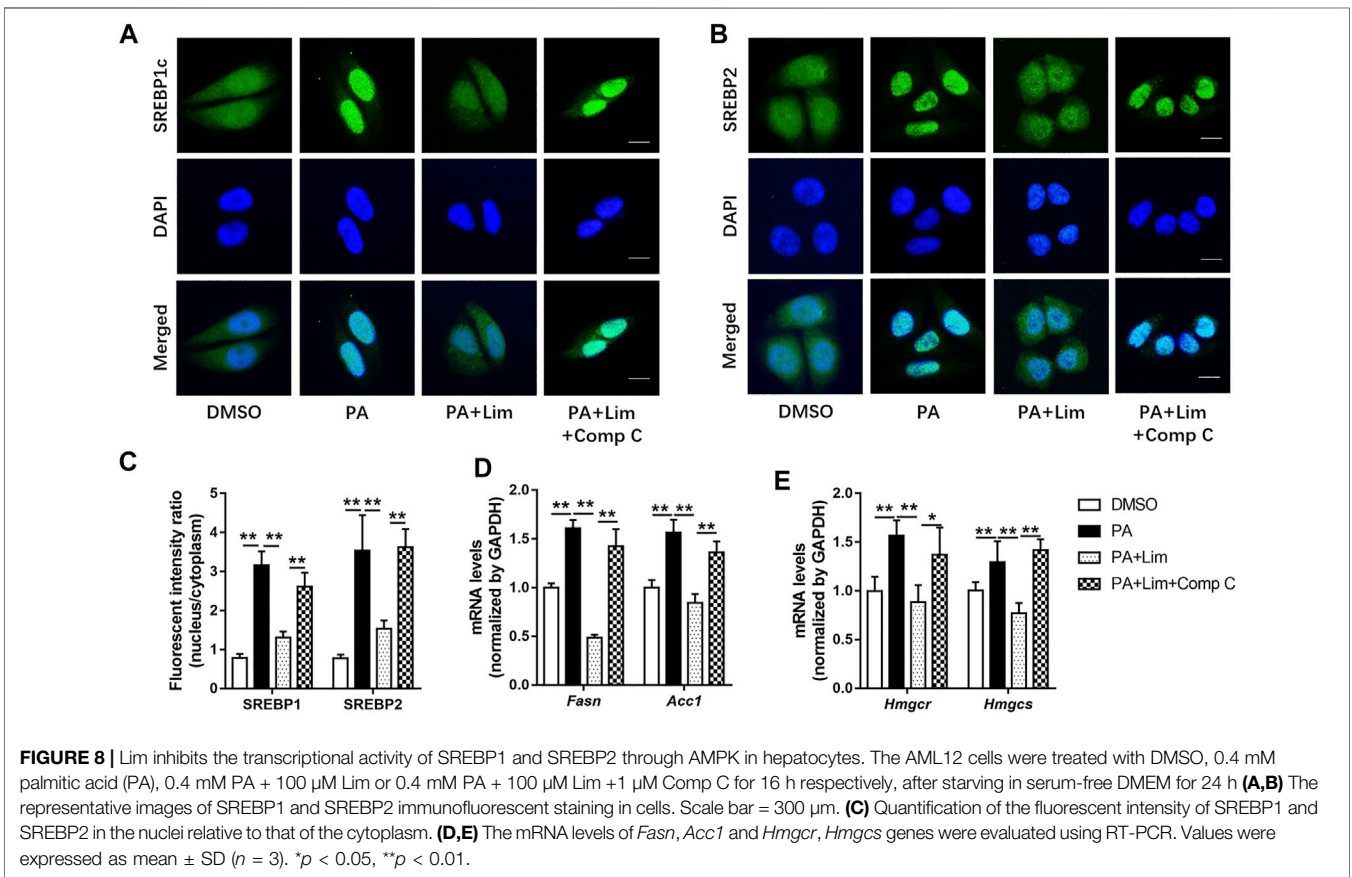


FIGURE 8 | Lim inhibits the transcriptional activity of SREBP1 and SREBP2 through AMPK in hepatocytes. The AML12 cells were treated with DMSO, 0.4 mM palmitic acid (PA), 0.4 mM PA + 100 μ M Lim or 0.4 mM PA + 100 μ M Lim + 1 μ M Comp C for 16 h respectively, after starving in serum-free DMEM for 24 h **(A,B)** The representative images of SREBP1 and SREBP2 immunofluorescent staining in cells. Scale bar = 300 μ m. **(C)** Quantification of the fluorescent intensity of SREBP1 and SREBP2 in the nuclei relative to that of the cytoplasm. **(D,E)** The mRNA levels of *Fasn*, *Acc1* and *Hmgcr*, *Hmgcs* genes were evaluated using RT-PCR. Values were expressed as mean \pm SD ($n = 3$). * $p < 0.05$, ** $p < 0.01$.

activating AMPK. To test our hypothesis, we determined the intracellular localization of SREBP1c and SREBP2, two most crucial transcriptional factors for lipid metabolism gene expression, after PA induction. As shown in **Figures 8A,B**, PA induced nuclear translocation of SREBP1c and SREBP2, while Lim treatment decreased the gathering of these two proteins in nuclei. Inhibition of AMPK by Compound C enhanced the nuclear translocation of SREBP1c and SREBP2 in Lim treated hepatocytes (**Figures 8A–C**). In addition, we examined the expression of downstream targets of SREBP1c and SREBP2. The administration of Compound C significantly re-elevated the expression of *Fasn* and *Acc1* regulated by SREBP1c, and the expression of *Hmgcr* and *Hmgcs* regulated by SREBP2 in Lim treated AML12 cells (**Figures 8D,E**).

DISCUSSION

Here, we identify, for the first time, that limonin inhibits hepatic lipid accumulation in HFD fed mice via the activation of AMPK. Limonoids are highly oxygenated triterpenoid compounds. Limonin, nomilin and obacunone are the major limonoids in Citrus (Gualdani et al., 2016). In 2011, Eri Ono's publication stated that nomilin and obacunone can activate TGR5, a bile acid membrane receptor (Ono et al., 2011). The activation of TGR5 by bile acids increases energy expenditure and attenuates diet-induced obesity in mice (Thomas et al., 2009). Eri Ono suggested that the anti-obesity and anti-hyperglycemic effects by nomilin was through activating TGR5. Furthermore, they pointed out that limonin was not a TGR5 agonist (Ono et al., 2011). Despite some studies demonstrated the effect of limonin on reducing inflammation and oxidative stress, few reported the function of limonin in the regulation of glucose and lipid metabolism, as its incapability of activating TGR5. The current study uncovered the significant role of limonin in metabolic regulation. During our previous studies, we investigated the effects of several active compounds in the citrus aurantium on NAFLD (Bai et al., 2019; Wang et al. 2020a; Wang et al. 2021a). Using cell based system, we screened natural products with regulatory effect on AMPK activity. We found that limonin can activate AMPK *in vivo* and *in vitro*. As we known, there has been no report on the activation of AMPK by limonin so far. The key regulatory roles of AMPK on glucose and lipid homeostasis further support observation that limonin can effectively ameliorate metabolic disorders including NAFLD.

A wide variety of drugs and xenobiotics has been found to activate AMPK, for example, antidiabetic drugs like metformin, phenformin and thiazolidinediones, and natural products derived from traditional medicines or foods such as berberine, quercetin, resveratrol and epigallocatechin gallate (Hawley et al., 2010). However, the underlying mechanisms by which AMPK is activated are remain unclear in many cases. Adenine nucleotides are compartmentalized between the mitochondria and cytoplasm and their diffusion is limited, they may not be uniformly distributed within the cytoplasm. The AMPK is extremely sensitive to small changes in AMP.

Subtle changes in subcellular nucleotides that may not be detectable by measuring total cellular levels would be enough to influence the activity of AMPK. The measurement of cellular contents of AMP and ADP in cell extracts will lose the information of their effects of compartmentation. Therefore, while an agent like metformin may activate AMPK without producing a detectable change in AMP, this does not prove the effect is AMP independent (Hawley et al., 2010). To determine whether or not activation of AMPK by different agents was AMP dependent, Simon A. Hawley et al. (2010) developed a test. The mutation of AMPK γ 2 subunit, R531G, causes a severe loss of binding by AMP and ATP, thus generating an AMP-insensitive AMPK complex. They expressed wild-type or R531G mutant γ 2 subunit in cells to replace the endogenous γ 1 subunit in the assay. This method was widely adopted by later researches, and was carried out in our current study. We used oligomycin and A769662 as positive controls to validate the effectiveness of the test (**Supplementary Figure S4**). Using this system, we found that Lim activated AMPK through inhibition of cellular energy metabolism and increasing ADP: ATP ratio. However, more investigation is needed in future to elaborate whether Lim increased AMPK activity through initiating/inhibiting membrane receptor associated signal pathways, uncoupling the respiratory oxidation and ATP generation in the mitochondria, or interfering glycolysis.

At the beginning of the current study, we examined the changes of systemic metabolism in mice with or without limonin treatment. Although the remission of liver pathological changes is the focus of our attention, it is obvious that limonin has a significant impact on systemic metabolic indicators, including body weight, body fat, blood sugar, blood lipids and insulin sensitivity. These indicators are affected by diverse organs such as liver, adipose tissue, skeletal muscle as well as other factors like neuroendocrine and intestinal flora. Conversely, changes in these indicators also affect the metabolic status of liver, adipose tissue, skeletal muscle and other organs. Through cell based assays *in vitro*, we determined that limonin had a direct effect on hepatic cells. Even though, we can't rule out the possibility that limonin relieved hepatic steatosis and inflammation in mice induced by high-fat diet through its joint action on the liver and other tissues such as adipose tissue. In the body, limonin may first act on the liver, and affect the metabolism of adipose tissue through secretory factors like hepatokines, thereby improving systemic metabolism. It may also act directly on adipose tissue, lowering blood lipids, thereby alleviating hepatic steatosis. This effect superimposed with its direct effect on the liver, resulting in the overall influence we observed. These speculations warrant more investigations in future.

CONCLUSION

In summary, we found that limonin activated AMPK in hepatocytes and attenuated hepatic lipid accumulation. These findings suggest limonin as potential therapeutics for MAFLD and warrant more detailed investigation for its underlying mechanisms in future.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding author.

ETHICS STATEMENT

The animal study was reviewed and approved by The Animal Care and Use Committee of Zhejiang Chinese Medical University.

AUTHOR CONTRIBUTIONS

Participated in research design: FeZ and S-wW. Conducted experiments: S-wW, TL, H-fC, HS, C-yX, L-fX, and FaZ. Performed data analysis: S-wW, FeZ, and H-fC. Wrote or contributed to the writing of the manuscript: FeZ and S-wW.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fphar.2022.833705/full#supplementary-material>

- Wang, S. W., Xu, Y., Weng, Y. Y., Fan, X. Y., Bai, Y. F., Zheng, X. Y., et al. (2018). Astilbin Ameliorates Cisplatin-Induced Nephrotoxicity through Reducing Oxidative Stress and Inflammation. *Food Chem. Toxicol.* 114, 227–236. doi:10.1016/j.fct.2018.02.041
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