**Name of Journal: *World Journal of Gastroenterology***

**Manuscript NO: 52092**

**Manuscript Type: ORIGINAL ARTICLE**

# *Basic Study*

# LB100 ameliorates nonalcoholic fatty liver disease *via* the AMPK/Sirt1 pathway

Chen XY *et al*. LB100 ameliorates NAFLD

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# Author contributions: Chen XY and Cai CZ designed and performed the study, conducted the statistical analysis and wrote the paper; Yu ML, Feng ZM, Zhang YW, Liu PH and Zeng H performed the study and provided guidance during revision; Yu CH supervised the study and provided consultation during the entire study.

**Institutional animal care and use committee statement:** All experiments were conducted with approval of the First Afﬁliated Hospital of Zhejiang University Institutional Animal Care and Use Committee.

**Conflict-of-interest statement**: Authors have no conflicts of interest to disclose.

**Data sharing statement:** No additional data are available.

**ARRIVE guidelines statement**: Authors have read the ARRIVE guidelines and prepared and revised the manuscript according to the ARRIVE guidelines.

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**Manuscript source:** Unsolicited manuscript

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**Received:** October 16, 2019

**Peer-review started:** October 16, 2019

**First decision:** November 4, 2019

**Revised:** November 10, 2019

**Accepted:** November 23, 2019

**Article in press:** November 23, 2019

**Published online:** December 7, 2019

# Abstract

BACKGROUND

It is well known that nonalcoholic fatty liver disease (NAFLD) is associated with insulin resistance (IR). LB100, a serine/threonine protein phosphatase 2A (PP2A) inhibitor, is closely related to IR. However, there is little data regarding its direct influence on NAFLD.

***AIM***

To elucidate the effect and underlying mechanism of LB100 in NAFLD.

METHODS

After 10 wk of high fat diet (HFD) feeding, male C57BL/6 mice were injected intraperitoneally with vehicle or LB100 for an additional 6 wk (three times a week). The L02 cell line was treated with LB100 and free fatty acids (FFAs) for 24 h. Hematoxylin and eosin and oil red O staining were performed for histological examination. Western blot analysis was used to detect the protein expression of Sirtuin 1 (Sirt1), total and phosphorylated AMP-activated protein kinase α (AMPKα), and the proteins involved in lipogenesis and fatty acid oxidation. The mRNA levels were determined by qPCR. Pharmacological inhibition of AMPK was performed to further examine the exact mechanism of LB100 in NAFLD.

RESULTS

LB100 significantly ameliorated HFD-induced obesity, hepatic lipid accumulation and hepatic injury in mice. In addition, LB100 significantly downregulated the protein levels of acetyl-CoA carboxylase, sterol regulatory element-binding protein 1 and its lipogenesis target genes, including stearoyl-CoA desaturase-1 and fatty acid synthase, and upregulated the levels of proteins involved in fatty acid β-oxidation, such as peroxisome proliferator-activated receptor α (PPARα), peroxisome proliferator-activated receptor gamma coactivator-1α (PGC-1α), carnitine palmitoyltransferase 1α, acyl-CoA oxidase 1 and uncoupling protein 2, as well as the upstream mediators Sirt1 and AMPKα in the livers of HFD-fed mice. In *vitro*, LB100 alleviated FFA-induced lipid accumulation in L02 cells through the AMPK/Sirt1 signaling pathway. Further studies showed that the curative effect of LB100 on lipid accumulation was abolished by inhibiting AMPKα in L02 cells.

CONCLUSION

PP2A inhibition by LB100 significantly ameliorates hepatic steatosis by regulating hepatic lipogenesis and fatty acid oxidation via the AMPK/Sirt1 pathway. LB100 may be a potential therapeutic agent for NAFLD.

**Key words:** LB100; Nonalcoholic fatty liver disease; Serine/threonine-protein phosphatase 2A; Lipid metabolism; AMP-activated protein kinase α; Sirtuin 1

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Core tip: Nonalcoholic fatty liver disease (NAFLD) is one of the most common chronic liver diseases worldwide, which greatly increases the medical and economic burden. We aimed to investigate the effect and underlying mechanism of LB100 in lipid accumulation during NAFLD development in mice fed a high fat diet and L02 cells treated with free fatty acids. Our study provided, for the first time, in vivo and in vitro evidence that LB100 can effectively inhibit hepatic lipogenesis via the AMPK/Sirt1 pathway and could be a therapeutic strategy for NAFLD.

**Citation:** Chen XY, Cai CZ, Yu ML, Feng ZM, Zhang YW, Liu PH, Zeng H, Yu CH. LB100 ameliorates nonalcoholic fatty liver disease *via* the AMPK/Sirt1 pathway. *World J Gastroenterol* 2019; 25(45): 6607-6618

**URL:** https://www.wjgnet.com/1007-9327/full/v25/i45/6607.htm

**DOI:** https://dx.doi.org/10.3748/wjg.v25.i45.6607

**INTRODUCTION**

Nonalcoholic fatty liver disease (NAFLD) includes a series of hepatic metabolic disorders characterized by excessive hepatic fat accumulation without a history of significant alcohol consumption[1]. It ranges from simple steatosis to nonalcoholic steatohepatitis (NASH), irreversible fibrosis, cirrhosis, and eventually hepatocellular carcinoma (HCC)[2]. NAFLD is one of the most common chronic liver diseases worldwide, with a global prevalence of 25.2% and a prevalence of 29.2% in China[3,4], which greatly increases the medical and economic burden. However, its exact pathogenesis remains poorly understood. Lifestyle modification is advocated for treating patients with NAFLD, but its efficacy is limited. Optional medications include vitamin E, pioglitazone and pentoxifylline, but there is a lack of Food and Drug Administration-approved treatments[5]. Therefore, there is an urgent need to develop effective drug therapies for NAFLD.

AMP-activated protein kinase (AMPK) is a monitor of cellular energy status; once activated, it inhibits various anabolic pathways, stimulates catabolic pathways, suppresses ATP consumption, and increases ATP production to restore energy homeostasis[6,7]. AMPK enhances Sirtuin 1 (Sirt1) activity by increasing cellular NAD+ levels, further leading to the deacetylation and activity regulation of downstream Sirt1 targets such as Peroxisome proliferator-activated receptor gamma coactivator-1α (PGC1α)[8]. Numerous studies have found that AMPK and Sirt1 are closely related to lipid metabolism and activate each other in a finely tuned network[8,9]. Phosphorylated AMPK can directly target Acetyl-CoA carboxylase(ACC), phosphorylating and inactivating it[10]. AMPK activation also reduces the transcriptional activation of sterol regulatory element-binding protein 1c (SREBP-1c), which is a critical transcription factor in the regulation of lipogenic genes, including *ACC*, fatty acid synthase(*FAS*) andstearoyl-CoA desaturase-1 (*SCD1*)[11,12]. In addition, AMPK has been shown to be closely related to fatty acid β oxidation[13,14]. In summary, the AMPK-Sirt1 axis plays an important role in lipid metabolism.

LB100 (Supplementary Figure 1A) is a first-line serine/threonine protein phosphatase 2A (PP2A) small molecule inhibitor that is water-soluble and was initially designed to increase the sensitivity of solid tumors to chemotherapy[15]. Emerging evidence suggests that overactivation of PP2A is closely related to elevated insulin resistance (IR) in the liver[16,17]. PP2A regulates the phosphorylation of AMPK by dephosphorylating Thr-172, a residue that increases kinase activity when phosphorylated[18,19]. The direct interaction between PP2A and AMPK was confirmed by co-immunoprecipitation and co-localization experiments[19]. Therefore, we hypothesized that the effect of LB100 on lipid metabolism was at least partly mediated through the AMPK/Sirt1 pathway.

In this study, we investigated the effect of LB100 on ameliorating hepatic lipid accumulation in both *in vivo* and *in vitro* NAFLD models and its potential interaction with the AMPK/Sirt1 pathway, which may provide a new approach for the effective treatment of NAFLD.

**MATERIALS AND METHODS**

***Animals and administration of LB100***

Male C57BL/6 [mice](https://www.sciencedirect.com/topics/pharmacology-toxicology-and-pharmaceutical-science/mouse) (6 wk, 18-22 g), purchased from B&K Laboratory Animal Corp., Ltd. (Shanghai, China) were randomly distributed into four treatment groups: Standard chow diet (SCD) + vehicle, SCD + LB100 1.5 mg/kg, HFD + vehicle, HFD + LB100 1.5 mg/kg. The mice were housed in a specific pathogen-free environment (24-26℃, relative humidity: 50%-60%) with a 12-h light/dark cycle and free access to food and water. For 10 wk, the [mice](https://www.sciencedirect.com/topics/earth-and-planetary-sciences/mice) were fed either a SCD or a HFD (60% of kilocalories as [fat](https://www.sciencedirect.com/topics/pharmacology-toxicology-and-pharmaceutical-science/fat); Product D12492, Research Diets, New Brunswick, NJ, United States). The animals were then injected intraperitoneally with the vehicle or LB100 (three times a week) dissolved in normal saline for the next 6 wk. Blood and tissue samples were stored at -80℃. All animal experiments were performed according to the guidelines approved by the Animal Care and Use Committee of the First Affiliated Hospital, College of Medicine, Zhejiang University (Permit number: 2018-842).

***Cell culture and treatments***

The normal human hepatic cell line (L02), obtained from the Chinese Academy of Science (Shanghai, China), was maintained in Dulbecco’s modified Eagle’s medium (high glucose) supplemented with 10% fetal bovine serum and 1% antibiotics at 37℃ with 95% humidified air/5% CO2. L02 cells were exposed to a mixture of free fatty acids (FFAs; oleate acid and palmitate acid, final ratio 2:1; Sigma-Aldrich, St. Louis, MO, United States) at a final concentration of 1 mmol/L with or without LB100 (Selleckchem; 6 μmol/L) for 24 h. To investigate the effect of the AMPK signaling pathway on LB100-regulated lipid metabolism, the cells were pretreated with the AMPK inhibitor compound C (10 μmol/L) for 2 h and treated with LB100 supplemented with FFAs for 24 h.

***Cell viability***

L02 cells were seeded in 96-well plates at a density of 5 × 103 cells/well and cultured overnight. Subsequently, the cells were pretreated with various concentrations of LB100 (100 nmol/L–100 µmol/L) for 24 h at 37℃. Cell viability was estimated using the cell counting kit-8 colorimetric assay (Dojindo Molecular Technologies, Inc., Kumamoto, Japan).

***Western blot analysis***

Total proteins were extracted from cells and liver tissues using RIPA buffer (Applygen Technologies Inc., Beijing, China) supplemented with protease and phosphatase inhibitors (Sigma). Equal amounts of protein were separated by 8%-10% SDS-PAGE followed by transfer to PVDF membranes (Millipore, Inc., Darmstadt, Germany). The membranes were blocked with 5% nonfat dry milk in TBST, followed by incubation overnight with the following primary antibodies: anti-PP2A subunit C (2259), anti-p-AMPKα (2535), anti-AMPKα (2603), anti-Sirt1 (8469), anti-GAPDH (2118), anti-SCD1 (2438), anti-PGC1α (2178), anti-carnitine palmitoyltransferase 1α (CPT1α) (12252), anti-uncoupling protein 2 (UCP2) (89326) (Cell Signaling Technology, Beverly, MA, USA), anti-PP2A α+β (methyl L309) (ab66597), anti-p-ACC (ab31931), anti-ACC (ab45174), anti-FAS (ab128856), anti-SREBP1 (ab3259), and anti-PPARα (ab8934) (Abcam, Cambridge, UK). Proteins were visualized with an enhanced ECL kit (Fudebio, Hangzhou, China).

***RNA isolation and real-time PCR***

Total RNA was extracted from cells or livers with RNA plus (Takara, Dalian, China), and cDNA synthesis was performed with a One Step PrimeScript® RT-PCR kit (Takara, Japan) using 2.5 μg total RNA. Real-time PCR analysis was carried out using the SYBR Premix-Ex Tag Kit (Takara) on an ABI Prism 7500 sequence detection system (Applied Biosystems, Foster City, CA, United States). The primer sequences are listed in Supplementary Table 1.

***Glucose tolerance tests and insulin tolerance tests***

Glucose tolerance tests (GTTs) and insulin tolerance tests (ITTs) were performed after feeding the mice for 14 or 15 wk with the HFD, respectively. For GTTs, the mice were injected intraperitoneally with glucose solution (1 mg/g body weight) after a 16-h fast. For ITTs, the mice were injected intraperitoneally with insulin solution (0.75 mU/g body weight) after a 6-h fast. Blood glucose levels were measured from the tail tip using a Freestyle brand glucometer (LifeScan, Shanghai, China).

***PP2A phosphatase activity assay***

After the designated LB100 treatment, L02 cells or mouse livers were lysed in the previously described RIPA lysis buffer. Supernatants containing 50 μg of total cellular proteins were assayed with the PP2A assay kit (Millipore) according to the manufacturer's instructions. The PP2A activity data are presented as the percentage of relative PP2A activity compared with that of the control.

***Hepatic and cellular triglyceride assay***

Liver tissue homogenate and cells were collected for intrahepatic and intracellular triglyceride (TG) determination using a commercial kit (Applygen Technologies Inc.) according the manufacturer’s instructions. TG values were normalized to total protein content.

***Histological analysis and Oil Red O staining***

Liver sections were fixed in 10% formalin overnight, embedded in paraffin, sectioned and then stained with hematoxylin and eosin (H and E) for histological examination. For the detection of lipid accumulation, frozen liver sections and cell slides were stained with oil red O according to a standard protocol (Nanjing Jiancheng Bioengineering Institute, Nanjing, China). Finally, liver sections and cell slides were imaged at 200 × magnification (Olympus, Tokyo, Japan). The data shown are from one representative experiment of three independent repeats.

***Statistical analysis***

The statistical methods used in this study were reviewed by Yi Zhang from the Institute of Statistics, School of Mathematical Sciences, Zhejiang University of China. Student’s *t*-test or two-way ANOVA was used in the statistical analysis. All statistical analyses were performed with SPSS 23 (IBM, Chicago, IL, United States). *P* < 0.05 was considered statistically significant. Data are presented as the mean ± standard deviation. All experiments were repeated at least three times independently.

**RESULTS**

***LB100 alleviates hepatic steatosis and enhances IR in HFD-fed mice***

Mice supplemented with LB100 for 6 wk (HFD+LB100 mice) showed significant reductions in body and liver weights compared with those of mice fed the HFD (HFD mice) (Figure 1A). With regard to liver appearance, HFD mice had markedly larger and paler livers than those of HFD+LB100 mice (Figure 1B). In accordance with H and E staining, oil red O staining and hepatic TG analysis showed a dramatic decrease in lipid levels in HFD+LB100 mice compared with those in HFD mice, whereas no difference was observed in LB100-treated SCD-fed mice (SCD+LB100 mice) and SCD-fed mice (SCD mice) (Figure 1C-E). LB100 also attenuated HFD-induced liver injury with a marked decrease in serum alanine aminotransferase and aspartate aminotransferase levels (Figure 1F). However, LB100 did not significantly decrease serum total cholesterol and TG levels in HFD mice (Supplementary Figure 2A and B). HFD mice showed elevated fasting insulin levels compared with those of HFD+LB100 mice (Figure 1G). In addition, we performed GTTs and ITTs to determine if glucose homeostasis was regulated by LB100 and found that glucose homeostasis and insulin sensitivity were improved in HFD+LB100 mice (Figure 1H-I). In addition, we evaluated liver fibrosis by staining with both Sirius red and Masson's trichrome, and found there were almost no signs of fibrosis (Supplementary Figure 3A and B). These findings demonstrate that LB100 enhances insulin sensitivity and ameliorates body weight, hepatic steatosis and liver injury in HFD mice.

***LB100 inhibits PP2A activity and activates the AMPK/Sirt1 pathway in vivo***

As expected, fatty acids increased liver PP2A activity, while LB100 reduced liver PP2A activity but did not affect protein levels of the PP2A catalytic subunit C (Figure 2A). In addition, the level of PP2A methylation in the livers of HFD mice was increased, while LB100 downregulated PP2A methylation levels (Figure 2A). As shown in Figure 2B, protein expression of Sirt1 and the ratio of P-AMPK (Thr172)/AMPK were markedly increased in HFD+LB100 mice. LB100 treatment significantly downregulated the levels of proteins involved in fatty acid synthesis (Figure 2C). In contrast, LB100 increased the expression of proteins involved in fatty acid β-oxidation, such as PPARα, PGC-1α, CPT1α, ACOX-1 and UCP-2, in the livers of HFD mice (Figure 2D). Moreover, we found that LB100 treatment significantly upregulated the [mRNA](https://www.sciencedirect.com/topics/pharmacology-toxicology-and-pharmaceutical-science/messenger-rna) levels of *SREBP1, FAS, SCD1* and *CPT1α* (Figure 2E). These results indicate that LB100 regulates lipid metabolism by inhibiting PP2A activity and activating the AMPK/Sirt1 pathway *in vivo*.

***LB100 inhibits PP2A activity and decreases FFA-induced lipid accumulation in hepatocytes***

We also investigated the effect of LB100 on FFA-induced lipid deposition *in vitro*. The optimal concentration of LB100 treatment was 6 μmol/L, as evaluated by CCK-8 assays (Supplementary Figure 1B). Consistent with the *in vivo* results, fatty acids induced overactivation of PP2A, while LB100 impaired PP2A activity (Figure 3A). LB100 attenuated the FFA-induced intracellular TG accumulation in a dose-dependent manner and attenuated lipid accumulation, as shown by oil red O staining (Figure 3B and C). These findings reveal that LB100 alleviates FFA-induced lipid deposition in hepatocytes.

***LB100 activates the AMPK/Sirt1 pathway in vitro***

Consistent with the *in vivo* findings, LB100 upregulated the expression of Sirt1 and p-AMPKα and significantly decreased the protein levels of SREBP1 and its target genes, including *SCD1, FAS* and *ACC,* in FFA-treated L02 cells (Figure 4A and B). LB100 also induced PPARα, PGC-1α, CPT1α, ACOX-1 and UCP-2expression *in vitro* (Figure 4C). As expected, LB100 significantly downregulated the [mRNA](https://www.sciencedirect.com/topics/pharmacology-toxicology-and-pharmaceutical-science/messenger-rna) levels of *SREBP1, FAS, ACC1* and *SCD1* (Figure 4D). These results indicate that LB100 ameliorates hepatic lipid accumulation *in vitro* by activating the AMPK/Sirt1 signaling pathway.

***Inhibition of AMPK*** ***eliminates the effect of LB100 on reducing hepatic steatosis***

We further examined whether AMPK plays a key role in LB100-mediated alleviation of hepatic steatosis. As predicted, the inhibitory effect of LB100 on hepatic lipid accumulation was eliminated by preincubation with the *AMPK* inhibitor compound C. Cellular TG analysis showed a dramatic increase in lipid level with compound C use (Figure 5A). Compound C treatment counteracted the activation of the AMPK/Sirt1 signaling pathway by LB100 (Figure 5B) and significantly upregulated the protein levels of *ACC* and some other proteins involved in fatty acid synthesis (Figure 5C). Inhibition of AMPK also counteracted the LB100-induced increase in lipid oxidation protein levels, such as PPARα, CPT1α and UCP2 (Figure 5D). However, there was no significant difference in the protein levels of PGC1α and ACOX-1. (Supplementary Figure 4A). In addition, compound C abolished the LB100-mediated reduction in the [mRNA](https://www.sciencedirect.com/topics/pharmacology-toxicology-and-pharmaceutical-science/messenger-rna) levels of *SREBP1, FAS, ACC1* and *SCD1* (Figure 5E). Taken together, these data suggest that *AMPK* is vital for the role of LB100 in lipid metabolism.

**DISCUSSION**

In this study, we demonstrated that LB100 can play an important role in liver lipid metabolism to prevent HFD-induced obesity, hepatic steatosis and IR in a NAFLD mouse model. LB100 reduced hepatic lipogenesis and promoted fatty acid β-oxidation *via* the AMPK/Sirt1 pathway in HFD-fed mice, although no similar changes were observed in SCD-fed mice. These findings provide molecular evidence supporting LB100 as a promising therapeutic strategy for NAFLD.

NAFLD is characterized by fat accumulation without excessive alcohol consumption and has complex potential mechanisms. At present, a widely recognized pathogenesis of NAFLD is the double-hit theory: The first “hit” is IR, which induces an increase in peripheral fat decomposition, abnormal liver lipid metabolism and hyperinsulinemia, leading to liver steatosis; then the second "hit" of endoplasmic reticulum stress occurs, leading to the development of hepatocyte inflammation, necrosis and fibrosis, further promoting the progression of NAFLD[20,21]. Therefore, inhibition of de novo lipogenesis, enhancement of fatty liver β-oxidation and improvement in IR are promising strategies for the prevention and treatment of NAFLD.

A series of studies have demonstrated that FFAs induce overactivation of PP2A both in mice and in cultured cell lines, and PP2A is a negative regulator of insulin signaling pathways[16,22]. Moreover, inhibition of PP2A protects against HFD-induced obesity and IR in mice[16,17]. LB100 is a derivative of the natural product cantharidin, which shows no obvious systemic toxicity and could be used as a pharmacologic inhibitor of PP2A[15]. However, the underlying mechanism by which LB100 inhibits hepatic steatosis remains unclear. In this study, we first tested the activity of PP2A by measuring the methylation status of the catalytic subunit of PP2A and using immunoprecipitation activity assays, and then we confirmed FFA-induced PP2A overactivation both *in vivo* and *in vitro*. After treatment with LB100, the activity of PP2A decreased with amelioration of lipid accumulation both *in vivo* and *in vitro*. In addition, LB100 improved glucose metabolism and [insulin sensitivity](https://www.sciencedirect.com/topics/medicine-and-dentistry/insulin-sensitivity) in HFD-fed mice. These results indicate that LB100 plays an important role in the development of IR and NAFLD.

Then, we further examined the potential mechanism of LB100 in lipid metabolism. Liangpunsakul *et al*[23] demonstrated, for the first time, that AMPK and PP2A can interact directly by immunoprecipitation. PP2A negatively regulates the phosphorylation state of *AMPK* by dephosphorylating Thr-172, as reported by Joseph *et al*[19]. We also found that LB100 increased the protein expression of P-AMPK (Thr172)/AMPK and Sirt1 both *in vivo* and *in vitro*. The results of the present study show that PP2A has essential roles in the regulation of autophagy. Marta Varela-Rey’s study showed that blocking PP2A activity restored autophagy flux in hepatocytes and ameliorated liver steatosis[24]. Some studies have described that inhibition of PP2A by okadaic acid suppresses autophagy in hepatocytes[25,26]. Our study, in which LB100 works through the activation of the AMPK/Sirt1 pathway, adds an exciting new mechanism through which LB100 exerts its beneficial effects. Undoubtedly, it is interesting to perform further studies to establish whether LB100 regulates lipid metabolism through the regulation of autophagy.

Recently, it was found that lipid accumulation in the liver inhibits AMPK activation and affects ACC expression, accelerating fatty acid synthesis and the development of NAFLD[27,28]. *AMPK* and *Sirt1* are closely related to lipid metabolism and activate each other in a finely tuned network[8,9]. Treatments that enhance AMPK/Sirt1 expression to inhibit ACC activity and to increase lipolysis and β-oxidation can improve NAFLD in HFD-fed mice[29,30]. Sirt1 regulates lipid homeostasis through multiple nutrient sensors such as SREBP1, PGC-1α and PPARα[31,32]. In this study, as expected, LB100 reduced the expression of adipogenic genes such as *SREBP1, FAS, ACC* and *SCD1* both at the protein and mRNA levels and increased the protein expression of fatty acid β-oxidation genes, including *PPARα, PGC1α, CPT1α,* *ACOX1* and *UCP2*. Finally, the use of the AMPK inhibitor compound C largely counteracted the effect of LB100 on the attenuation of fatty accumulation *in vitro*. Compound C significantly upregulated *SREBP1, FAS, ACC* and *SCD1,* at both the protein and mRNA levels, and downregulated the protein expression of PPARα, CPT1α and UCP2. However, there was no significant difference in the protein levels of PGC1α and ACOX1 after compound C treatment, which we consider to be a negative feedback response; the regulation of these two proteins may also be through another pathway. All of these findings indicate that AMPK signaling is a critical metabolic cue involved in the process by which LB100 alleviates lipid deposition both *in vivo* and *in vitro*.

However, there are some limitations to this study. First, the function of PP2A is extensive and complex, and AMPK may not be the only target gene regulated by LB100; therefore, further research is required to clarify the detailed downstream mechanism. Second, compound C treatment failed to neutralize the increased expression of PGC1α and ACOX1, and the regulatory mechanisms require further investigation.

In summary, our study provided, for the first time, *in vivo* and *in vitro* evidence that LB100 can effectively inhibit hepatic lipogenesis *via* the AMPK/Sirt1pathway and could be a therapeutic strategy for NAFLD.

**Article Highlights**

***Research background***

Nowadays, no pharmacological therapy is approved for nonalcoholic fatty liver disease (NAFLD). Recent studies have shown that serine/threonine protein phosphatase 2A (PP2A) is closely related to obesity and insulin resistance. LB100 is a water-soluble PP2A small molecule inhibitor. Thus, we hypothesize that LB100 can ameliorate hepatic lipid accumulation in fatty liver.

***Research motivation***

Increasing evidence indicates that PP2A regulates the AMPK, which is a monitor of cellular energy status.AMPK and Sirtuin 1 (Sirt1) are closely related to lipid metabolism and activate each other in a finely tuned network. However, as a first-line PP2A inhibitor, there is little data regarding the influence of LB100 on NAFLD and its underlying mechanism. Thus, studies on the potential effect of LB100 on NAFLD are urgently required.

***Research objectives***

To elucidate the effect and underlying mechanism of LB100 in NAFLD.

***Research methods***

This research was performed using C57BL/6 mice fed a high fat diet (HFD) for 16 wk and L02 cells stimulated with free fatty acids (FFAs) for 24 h to establish *in vivo* and *in vitro* models of hepatic steatosis. Mice were injected intraperitoneally with vehicle or LB100 (1.5 mg/kg, three times a week) and L02 cells were treated with LB100 (6 μmol/L) to determine the effect of LB100 on NAFLD.

***Research results***

LB100 significantly ameliorated HFD-induced obesity, hepatic lipid accumulation and hepatic injury in mice accompanied by activation of the AMPK/Sirt1 signaling pathway. Similar results were observed in L02 cells stimulated with FFAs. Further studies showed that the curative effect of LB100 on lipid deposition was abolished by pharmacological inhibition of AMPK in L02 cells.

***Research conclusions***

PP2A inhibition by LB100 significantly ameliorates hepatic steatosis by regulating hepatic lipogenesis and fatty acid oxidation *via* the AMPK/Sirt1 pathway.

***Research perspectives***

LB100 is a promising therapeutic strategy for NAFLD. Further clinical application should be considered.

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**P-Reviewer:** Di Leo A, Jamali R, Tarantino G, Treeprasertsuk S

**S-Editor:** Tang JZ **L-Editor:** Webster JR **E-Editor:** Ma YJ

**Specialty type:** Gastroenterology and hepatology

**Country of origin:** China

**Peer-review report classification**

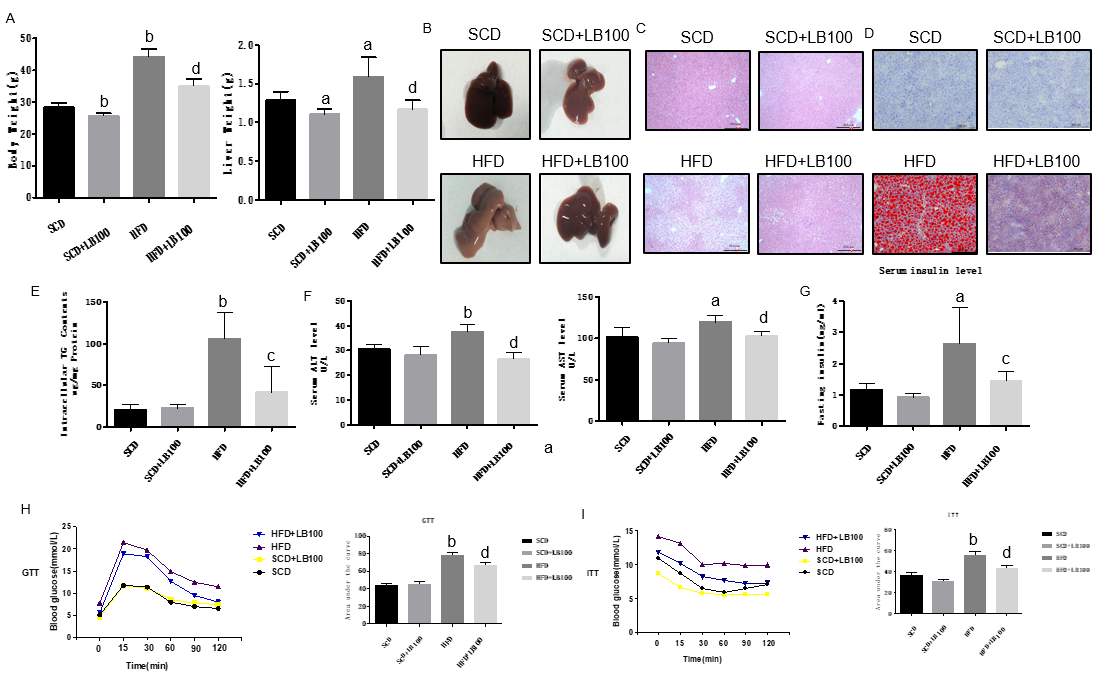
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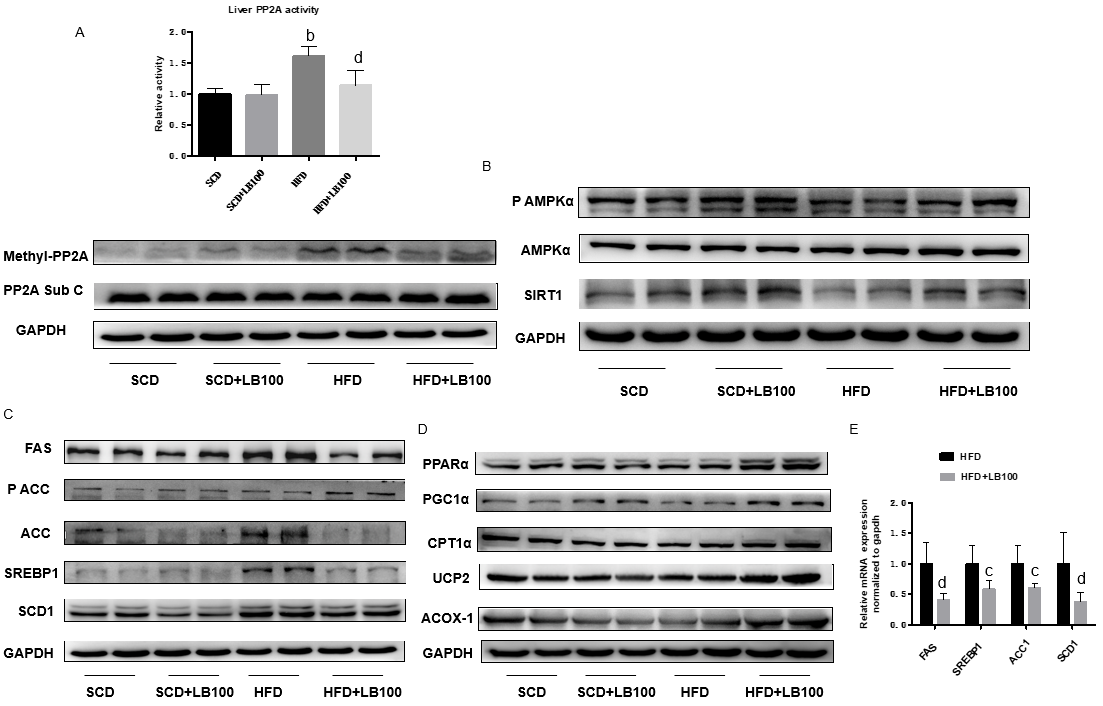
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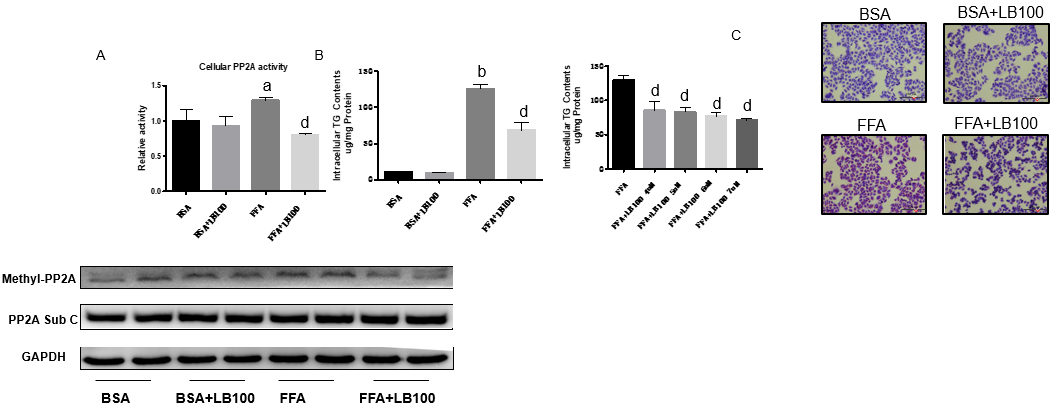
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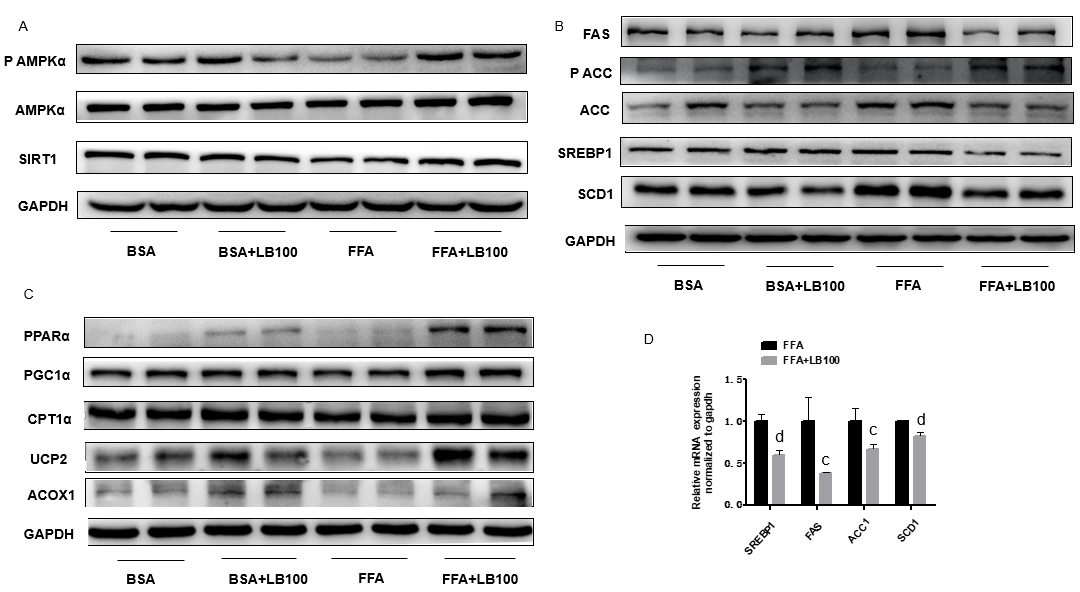
**Figure 1 LB100 reduces body weight, ameliorates hepatic steatosis and attenuates liver injury in high fat diet-fed mice.** A: Body weight and liver weight on the final day; B: Macroscopic view of livers from the four groups of mice; C: Representative liver histological section images of the four groups of mice stained with hematoxylin and eosin; D: Representative liver histological section images of the four groups of mice stained with oil red O; E: Triglyceride content of livers in the four groups; F: Serum levels of alanine aminotransferase and aspartate aminotransferase; G: Serum levels of insulin; H: Glucose tolerance test; I: Insulin tolerance test of mice fed with standard chow diet (SCD) or high fat diet (HFD) for 14 or 15 wk. Scale bar in panel represents 200 μm. Data are presented as the mean ± standard deviation. a*P* < 0.05, b*P* < 0.01 *vs* SCD mice; c*P* < 0.05, d*P* < 0.01 *vs* HFD mice. HFD: High fat diet; TG: Triglyceride; ALT: Alanine aminotransferase; AST: Aspartate aminotransferase; GTT: Glucose tolerance test; ITT: Insulin tolerance test; SCD: Standard chow diet.



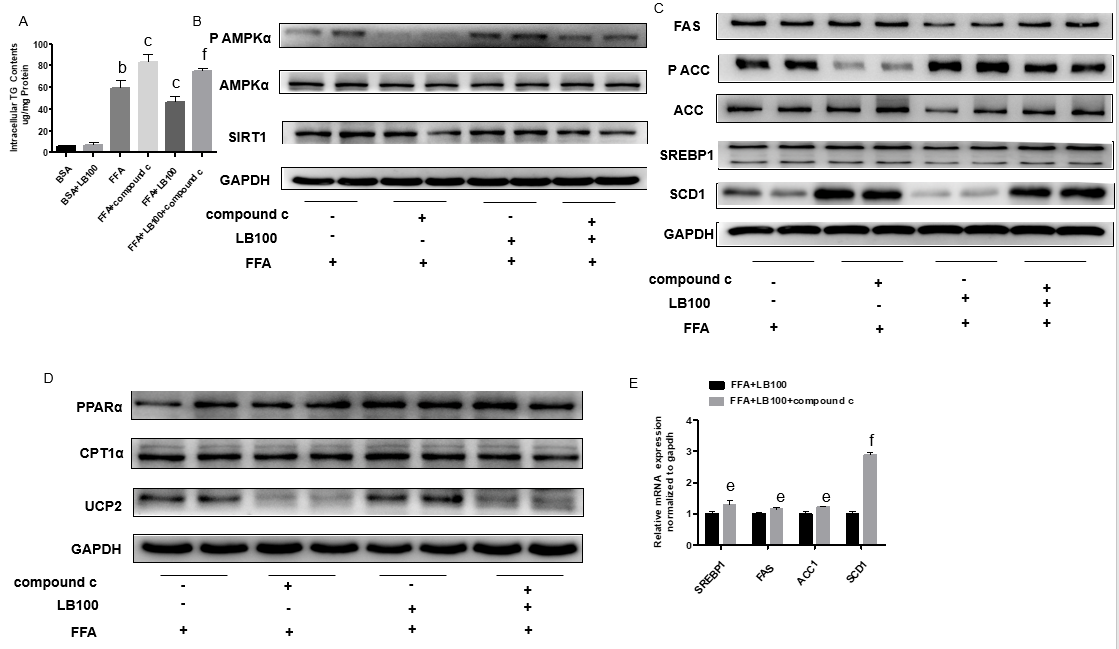
**Figure 2 LB100 inhibits serine/threonine protein phosphatase 2A activity, upregulates the expression of proteins involved in fatty acid β-oxidation, downregulates the protein levels of lipogenesis genes, and activates the AMPK/Sirt1 signaling pathway in the liver tissues of high fat diet-fed mice.** A: Serine/threonine protein phosphatase2A (PP2A) activity tested by a commercial assay (upper), protein expression of methylated PP2A and PP2A C subunit (lower) of mouse livers from the four groups; B: Western blot analysis of phosphorylated AMP-activated protein kinase α, phosphorylated AMP-activated protein kinase α, and Sirtuin 1; C: Proteins involved in lipid synthesis, such as fatty acid synthase (FAS), acetyl-CoA carboxylase (ACC), sterol regulatory element-binding protein 1 (SREBP1) and stearoyl-CoA desaturase-1 (SCD1); D: Proteins involved in fatty acid β-oxidation, such as peroxisome proliferator-activated receptor α, peroxisome proliferator-activated receptor gamma coactivator-1α, carnitine palmitoyltransferase 1α, uncoupling protein 2, and acyl-CoA oxidase 1; E: Gene expression levels of SREBP1, FAS, ACC1 and SCD1 were determined using real-time PCR. Three independent experiments were analyzed, and the data are presented as the mean ± standard deviation. a*P* < 0.05, b*P* < 0.01 *vs* standard chow diet mice; c*P* < 0.05, d*P* < 0.01 *vs* HFD mice. PP2A: Serine/threonine protein phosphatase2A; P AMPK α: Phosphorylated AMP-activated protein kinase α; Sirt1: Sirtuin 1; FAS: Fatty acid synthase; ACC: Acetyl-CoA carboxylase; SREBP1: Sterol regulatory element-binding protein 1; SCD1: Stearoyl-CoA desaturase-1; PPARα: Peroxisome proliferator-activated receptor α; PGC1α: Peroxisome proliferator-activated receptor gamma coactivator-1α; CPT1α: Carnitine palmitoyltransferase 1α; UCP2: Uncoupling protein 2; ACOX1: Acyl-CoA oxidase 1; SCD: Standard chow diet; HFD: High fat diet.



**Figure 3 LB100 inhibits serine/threonine protein phosphatase 2A activity and decreases free fatty acid-induced lipid accumulation in L02 cells.** Free fatty acid (1 mmol/L)-stimulated L02 cells were treated with LB100 (6 μmol/L) or vehicle for 24 h. A: Serine/threonine protein phosphatase 2A (PP2A) activity tested by a commercial assay (upper), protein expression of methylated PP2A and PP2A C subunit (lower) of L02 cells in the different groups; B: Intracellular triglyceride content in L02 cells; C: Representative image of oil red O staining of L02 cells in the different groups. Scale bar in panel represents 100 μm. Data are presented as the mean ± standard deviation. a*P* < 0.05, b*P* < 0.01 *vs* bovine serum albumin group; c*P* < 0.05, d*P* < 0.01 *vs* FFA group. FFA: Free fatty acid; PP2A: Serine/threonine protein phosphatase 2A; TG: Triglyceride; BSA: Bovine serum albumin.



**Figure 4 LB100 downregulates the protein levels of lipogenesis genes, upregulates the expression of proteins involved in fatty acid β-oxidation, and activates the AMPK/Sirt1 signaling pathway in L02 cells.** A: Protein expression levels of phosphorylated AMP-activated protein kinase α, AMPKα and Sirtuin 1 were detected by western blot analysis; B: Western blot analysis of proteins involved in lipid synthesis, such as fatty acid synthase (FAS), acetyl-CoA carboxylase (ACC), sterol regulatory element-binding protein 1 (SREBP1), and stearoyl-CoA desaturase-1 (SCD1); C: Proteins involved in fatty acid β-oxidation, such as peroxisome proliferator-activated receptor α, peroxisome proliferator-activated receptor gamma coactivator-1α, carnitine palmitoyltransferase 1α, uncoupling protein 2 and acyl-CoA oxidase 1; D: The mRNA levels of SREBP1 and its lipogenesis target genes, including FAS, ACC1 and SCD1, were determined using real-time PCR. Three independent experiments were analyzed, and the data are presented as the mean ± standard deviation. a*P* < 0.05, b*P* < 0.01 *vs* bovine serum albumin group; c*P* < 0.05, d*P* < 0.01 *vs* FFA group. P AMPK α: Phosphorylated AMP-activated protein kinase α; Sirt1: Sirtuin 1; FAS: Fatty acid synthase; ACC: Acetyl-CoA carboxylase; SREBP1: Sterol regulatory element-binding protein 1; SCD1: Stearoyl-CoA desaturase-1; PPARα: Peroxisome proliferator-activated receptor α; PGC1α: Peroxisome proliferator-activated receptor gamma coactivator-1α; CPT1α: Carnitine palmitoyltransferase 1α; UCP2: Uncoupling protein 2; ACOX1: Acyl-CoA oxidase 1; BSA: Bovine serum albumin; FFA: Free fatty acid.



**Figure 5 Inhibition of AMP-activated protein kinase α abolishes the effects of LB100 on the alleviation of hepatic steatosis in L02 cells.** After pretreatment with the AMP-activated protein kinase α (AMPK α) inhibitor, compound C for 2 h, L02 cells were incubated in normal medium or medium containing free fatty acid (FFA) with or without LB100 for 24 h. A: Intracellular triglyceride (TG) content in L02 cells; B: phosphorylated AMPK α (P-AMPKα), AMPKα, and Sirtuin 1 protein expression was detected by western blot analysis; C: Western blot analysis of proteins involved in lipid synthesis, such as fatty acid synthase (FAS), acetyl-CoA carboxylase (ACC), sterol regulatory element-binding protein 1 (SREBP1), and stearoyl-CoA desaturase-1 (SCD1); D: proteins involved in fatty acid β-oxidation, such as peroxisome proliferator-activated receptor α, carnitine palmitoyltransferase 1α, and uncoupling protein 2; E: Gene expression levels of SREBP1, FAS, ACC1, and SCD1 were determined using real-time PCR. Three independent experiments were analyzed, and the data are presented as the mean ± standard deviation. a*P* < 0.05, b*P* < 0.01 *vs* bovine serum albumin group; c*P* < 0.05, d*P* < 0.01 *vs* FFA group; e*P* < 0.05, f*P* < 0.01 *vs* FFA+LB100 group. P AMPK α: Phosphorylated AMP-activated protein kinase α; Sirt1: Sirtuin 1; TG: Triglyceride; FFA: Free fatty acid; FAS: Fatty acid synthase; ACC: Acetyl-CoA carboxylase; SREBP1: Sterol regulatory element-binding protein 1; SCD1: Stearoyl-CoA desaturase-1; PPARα: Peroxisome proliferator-activated receptor α; CPT1α: Carnitine palmitoyltransferase 1α; UCP2: Uncoupling protein 2; BSA: Bovine serum albumin.