Schisandrin B ameliorates non-alcoholic liver disease through anti-inflammation activation in diabetic mice

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Abstract
Type 2 diabetes mellitus (T2DM) is a metabolic risk factor associated with non-alcoholic liver disease (NAFLD). Schisandrin B (Sch B) is a promising agent for NAFLD. However, the actions of Sch B on diabetes-associated NAFLD and the underlying mechanisms are not characterized. This study aimed to assess whether Sch B has beneficial effects on T2DM-associated NAFLD. Sch B (50 mg/kg, gavage) was administrated to C57BL/KSj db/db mice for 2 weeks. Body weight, liver weight, blood glucose, and insulin resistance were measured. Serum lipid level and liver function were detected using the biochemistry analyzer. Quantitative Real-Time PCR assay was used to evaluate mRNA levels of lipid metabolism genes. Terminal-deoxynucleotidyl Transferase Mediated Nick End Labeling (TUNEL) staining was performed to measure apoptosis in the liver. Pathological analysis and immunohistochemistry assessment were used to analyze hepatic steatosis and inflammatory infiltration. Sch B supplementation significantly decrease body weight, related liver weight, blood glucose, and insulin resistance in db/db mice. Sch B obviously corrected NAFLD phenotypes including lipid deposition, steatohepatitis, and high levels of hepatic enzymes and serum lipid. In addition, mRNA levels of Sterol response element-bind protein 1c (SREBP-1c), fatty acid synthetase (Fasn), and acetyl-CoA carboxylase (ACC) were markedly downregulated by Sch B treatment. TUNEL-positive cells were also decreased by Sch B. Furthermore, Sch B inhibited the Kupffer cells, IL-1β, and TNF-α infiltration to the liver. Sch B ameliorated insulin resistance and lipid accumulation under high glucose conditions, which was partly associated with its inhibition of apoptosis and anti-inflammatory actions.

KEYWORDS
inflammatory infiltration, insulin resistance, non-alcoholic fatty liver disease, Schisandrin B, type 2 diabetes mellitus

Ruojia Ma and Yike Zhan contributed equally to this study.

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1 | INTRODUCTION

Non-alcoholic fatty liver disease (NAFLD) corresponds to the presence of hepatic steatosis and inflammatory infiltration in the absence of heavy alcohol consumption. It is commonly believed that NAFLD is strongly correlated with clinic features of metabolic syndrome (Filippatos et al., 2021; Hazer et al., 2020). The prevalence of NAFLD dramatically rising that is in the range of 11.5%–46% in the general population, indicative of the magnitude of the medical and the economic burden globally (Antunes et al., 2021). Nearly 80% patients with type 2 diabetes mellitus (DM), obesity, insulin resistance, or hyperlipidemia have NAFLD (Filippatos et al., 2021; Hazer et al., 2020; Leoni et al., 2018). Type 2 DM, characterized by insulin resistance, promotes lipolysis in peripheral tissue and results in release of free fatty acids which could be store in the liver. This is one of the critical causes for development of hepatic steatosis (Shields et al., 2009). Current data show that DM is an independent risk factor for hepatocellular carcinoma (Streba et al., 2015). Therefore, early drug intervention treatment of NAFLD patient with concerning complications is important for reducing morbidity.

Traditional Chinese medicinal herbs have a long history in using of safeguarding health that may offer a potential manner for prevention and cure of NAFLD. Schisandrin B (Sch B), an active dibenzooctadiene lignan, is abundant in the traditional Chinese medicine of fruit of Schisandra chinensis (FSC) (Nasser et al., 2020). It has been well recognized that Sch B presents various of biological functions including antihyperlipidemic, antioxidant, anti-endoplasmic reticulum stress, as well as anti-inflammation (Chu et al., 2011; Zhang et al., 2019). It has been reported that Sch B decreases apoptotic neurons and inflammatory signaling molecules in ischemic stroke rats (Fan et al., 2020). Recently, the protective effect of Sch B on metabolic diseases has also been largely demonstrated both in vitro and in vivo (Chu et al., 2011; Mak & Ko, 1997; Pan et al., 2008). For example, Sch B dose-dependently inhibits steatosis in L02 hepatocytes induced by free fatty acids (FFAs), in part through the suppression of sterol response element-bind protein-1 (SREBP-1) and adipose differentiation-related protein (Chu et al., 2011). Paralleling the results showed in vitro study, administrated with Sch B at the dose of 50–200 mg/kg reduces hepatic total cholesterol (TC) and triglyceride (TG) in cholesterol/bile salt induced hypercholesterolaemia mice model (Pan et al., 2008). Similarly, in streptozotocin-induced diabetic rats, Sch B protects against carbon tetrachloride-mediated hepatotoxicity (Mak & Ko, 1997). Although, the role of Sch B in hepatic steatosis has been evaluated, the effective action and specific mechanisms of this lignan underlying SZDM-related NAFLD are poorly understood.

The pathogenesis of NAFLD has been described based on the “multiple hit” theory, which includes insulin resistance, inflammation, apoptosis, and even genetic factors (Di Sessa et al., 2019). The critical hallmark of NAFLD is chronic inflammation which results from the expression of pro-inflammatory cytokines stimulated by overload of free fatty acids (FFAs) in the liver (Cusi, 2012). It is well-established that lipid deposition in the liver can induce IL-1β and TNF-α expression, which in turn promotes inflammatory cells infiltration, and activates downstream signaling pathways, thus leading to the development of insulin resistance and NAFLD (Tilg, 2010). Clinic evidence demonstrate that circulating level of TNF-α is highly related to the degree of liver fibrosis in non-alcoholic steatohepatitis (NASH) patients (Lesmana et al., 2009). It has been demonstrated that Sch B suppressed the upregulated protein levels of TNF-α, IL-1β, MMP-2, and MMP-9 in focal ischemic cerebral (Lee et al., 2012). It is consistent with the evidence that Sch B protects against dextran sulfate sodium-induced inflammatory bowel damage in mice via NF-κB and MAPK signaling (Liu et al., 2015). Therefore, Sch B may be promising in anti-inflammatory therapy for NAFLD.

In this study, we used db/db diabetic mice with Sch B supplementation for 2 weeks. We planned to explore the role of Sch B on diabetic-associated NAFLD and inflammatory mechanism in vivo.

2 | MATERIALS AND METHODS

2.1 | Animals

Male, 10-week-old C57BL/KSJ db/m mice and C57BL/KSJ db/db mice were purchased from the Laboratory Animal Center of Hangzhou Medical College (Hangzhou, China). All animals allowed to acclimate for 2 weeks before the experiments. The mice were bred and housed in an air-conditioned environment with 12 h light–dark cycle, and were provided with a standard chow. The animals were free access to food and water in Hangzhou Medical College Animal Core. All studies involving animals were performed in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals, and approved by the Ethics Committee of Laboratory Animal Care and Welfare, Hangzhou Medical College (NO. 20200196).

2.2 | Animal model and sampling

Blood samples adopted from the tail of mice were monitored using a commercial glucose monitoring system (MAJOR, Taiwan). Db/db mice, whose blood glucose levels greater than 11 mmol/L were considered as diabetic mice. The mice were randomly divided into four groups (n = 6): db/m mice group (control); db/m + Sch B group with 50 mg/kg Sch B (Selleck, Shanghai, China) for subsequent 2 weeks; db/db group; and db/db + Sch B group (50 mg/kg, 2 weeks). The dose and period were chosen according to before report with minor modification (Kwan et al., 2015). The Sch B was dissolved with DMSO (DMSO final concentration < 0.5%), and each mouse was administrated 0.2 ml solution. The db/m group and db/db group was given an equal volume of DMSO solution. It was freshly configured each time. Body weight and blood glucose were recorded after drug treatment of 2-week.

At the end of administration, all mice were euthanized with phenobarbital sodium (40 mg/kg, i.p.). Blood was collected from mice eyes. Samples were still standing for 1 h at room
The analysis of microvesicular steatosis used a scale of 0 according to before described method (Kleiner et al., 2005). The livers were semiquantitatively analyzed by a blinded histopathologist according to previously reported (Zhou et al., 2016). Mice were subjected to intraperitoneal injection of insulin (2 U/kg), and blood was harvested from the tail at 0, 15, 30, 60, and 90 min, respectively. For ITT, mice were deprived of chow overnight before ITT assay, and were fasted for 4 h prior to ITT assay. The isolated RNA (2 μg) was then reversed-transcribed to cDNA using the PrimeScript™ RT Reagent Kit (Takara). Quantitative Real-Time PCR assay was implemented with a SYBR Premix-Ex Tag Kit (Takara) and an ABI PRISM 7700 Sequence Detection System (Applied Biosystems, Carlsbad). All samples were measured in duplicate. The mRNA level comparison was calculated using the 2^(-ΔΔCt) method.

2.3 | Metabolic parameters and inflammatory cytokines determination

The concentrations of serum insulin, IL-1β and TNF-α were measured with ELISA kits (Insulin: SHIBAYAGI, Shibukawa, Japan; IL-1β and TNF-α: R&D Systems, MN) as previously described (Jiang et al., 2017). In addition, serum alanine aminotransferase (ALT) and aspartate aminotransferase (AST) levels were determined using an automatic biochemistry analyzer (Backman, Miami, FL) according to the manufacturer’s instruction. TC and TG both in serum and tissues were quantified with commercial kits (Cayman Chemical, Ann Arbor, MI).

2.4 | Glucose tolerance test and insulin tolerance test

GTT and ITT were performed to evaluate the insulin resistance according to previously reported (Zhou et al., 2016). Mice were deprived of chow overnight before GTT assay, and were fasted for 4 h prior to ITT assay. For GTT, mice received intraperitoneal injection of glucose (2 g/kg), and blood was harvested from the tail at 0, 15, 30, 60, and 90 min, respectively. For ITT, mice were subjected to intraperitoneal injection of insulin (2 U/kg), and blood were collected at 0, 15, 30, 60, and 90 min, respectively. The values of blood glucose were measured with glucose monitoring system.

2.5 | Histological analysis

For the pathology examination, the fixed liver tissues were embedded in paraffin and cut into a thin section (5 μm). Then samples were stained with hematoxylin and eosin (H&E) for histology score. To visualize lipid deposition, the frozen liver section was used to Oil Red O staining. The samples were cut into 5 μm thin sections, and were stained with Oil Red O reagent for 10 min. Images were taken at 200× using a microscope (Leica Microsystems, Wetzlar, Germany).

The severity of steatosis, ballooning, and lobular inflammation in the livers were semiquantitatively analyzed by a blinded histopathologist according to before described method (Kleiner et al., 2005). The analysis of microvesicular steatosis used a scale of 0–3, where 0 referred to less than 5% of steatosis, 1 referred to less than 33% of steatosis, 2 referred to less than 66% of steatosis, 3 referred to more than 66% of steatosis. The evaluation of ballooning used a scale of 0–2: 0 referred to no sign of hepatocyte ballooning, 1 referred to several ballooning cells in the liver, 2 referred to mass ballooning cells. In addition, the evaluation criterion of infiltration of the lobular inflammatory cells was as follow: 0 referred to none of the lobular inflammatory cells, 1 referred to 0–2 inflammatory cells, 2 referred to 2–4 inflammatory cells, and 3 referred to more than 4 inflammatory cells. The sum of the above scores were defined as a NAFLD activity score, which 0–2 was identified as no symptom for NASH, 3–4 was identified as uncertain, more than 5 was identified as likely NASH.

2.6 | Isolation of RNA and quantitative real-time PCR

Total RNA of isolated hepatic tissues was purified using the RNAiso Plus (Takara, Dalian, China) according to the manufacturer’s protocol. The isolated RNA (2 μg) was then reversed-transcribed to cDNA using the PrimeScript™ RT Reagent Kit (Takara). Quantitative Real-Time PCR assay was implemented with a SYBR Premix-Ex Tag Kit (Takara) and an ABI PRISM 7700 Sequence Detection System (Applied Biosystems, Carlsbad). All samples were measured in duplicate. The mRNA level comparison was calculated using the 2^(-ΔΔCt) method.

2.7 | Terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) assay

To examine the apoptotic cells in the liver tissues, a TUNEL assay kit (Wuhan Boster Biological Technology, Wuhan, China) was applied. The procedure was performed according to the manufacturer’s instructions. Briefly, the slides were incubated with 1% TUNEL reaction mixture for 1 h at room temperature, and were rinsed three times with phosphate buffer saline (PBS). Images were taken at high magnification (200×) with a microscope (Leica Microsystems). TUNEL-positive cells in three randomly areas were quantitated in each slide.

2.8 | Immunohistochemical staining

Samples of liver tissues were excised, and fixed in 10% formalin for 24 h at room temperature. Liver tissues were embedded in paraffin, cut into 5 μm thin sections, and subsequently deparaffinization, rehydration, antigen retrieval and 5% normal goat serum blocking for 4 h at room temperature. The slides were incubated in specific primary antibodies for 4 h, and followed by PBS washing for three times. Then, slides were incubated with horseradish peroxidase-conjugated secondary antibodies for 2 h, and visualized using substrate diaminobenzidine. The following primary antibodies were used: CD68 (ab125212, Abcam, 1:800 dilution), F4/80 (LS-C96373-100, Lifespan, 1:800 dilution), IL-1β (SRP9033,
Sigma, 1:800 dilution), TNF-α (ab6671, Abcam, 1:1000 dilution). One sample from each mouse was detected, and three high-power fields at 200× were analyzed for each slide using a microscope (Leica). The number of positive cells were counted.

2.9 | Statistical analyses

All values are expressed as mean ± SEM with six mice each group to ensure their reliability. Statistical analysis was performed using
GraphPad Software Prism-8. Data of two groups were analyzed with Student's t-test, and three and four groups were analyzed with Newman–Keuls multiple comparison test. In each test, the conventional analysis of variance (F test) was applied to the ranks instead of the original data. The statistical significance was considered as $P$ less than .05.

3 | RESULTS

3.1 | Improvements of body weight, relative liver weight, and insulin resistance by Sch B treatment in db/db mice

As expected, body weight in db/db mice was significantly increased compared with db/m mice (56.92 ± 0.38 g vs. 25.48 ± 0.29 g, $P < .05$; Figure 1a). Although Sch B administrated did not change the body weight in db/m mice, it significantly reduced the body weight gain in db/db mice (54.17 ± 0.53 g vs. 56.92 ± 0.38 g, $P < .05$; Figure 1a). Similarly, the percentages of liver weight to body weight were higher in db/db mice than in the db/m mice (4.14 ± 0.03% vs. 3.70 ± 0.01%, $P < .05$; Figure 1b). Oral administration of Sch B did not alter the relative liver weight in db/m mice, but it significantly decreased this value in db/db mice (4.04 ± 0.02% vs. 4.14 ± 0.03%, $P < .05$; Figure 1b). Furthermore, db/db diabetic mice displayed greatly high blood glucose and serum insulin level, which were suppressed by Sch B (Blood glucose: 12.92 ± 5.88 vs. 16.24 ± 6.83 mmol/L, $P < .05$; Figure 1c; Insulin: 6.93 ± 0.38 vs. 8.57 ± 0.22 ng/ml, $P < .05$; Figure 1d). There were no significant differences of blood glucose and insulin concentrations in db/m mice with or without Sch B treatment ($P > .05$; Figure 1c,d). GTT and ITT assays were applied to further detect insulin resistance. As show in Figure 1e, db/db diabetic mice had a higher blood glucose level in GTT assay, while the blood glucose was partly inhibited by Sch B administration ($P < .05$). Calculation of area under curve (AUC) verified this result ($P < .05$; Figure 1f). In line with these results, Sch B administration also improved the injury of insulin sensitivity in db/db mice ($P < .05$; Figure 1g,h). There were no significant differences of both GTT and ITT assays in db/m mice with or without Sch B treatment ($P > .05$; Figure 1e–h).

**FIGURE 2** Improvements of hepatic pathology and hepatic function by Sch B treatment in db/db mice. (a–d) representative images of hepatic H&E staining (magnification, 200×). The black arrow pointed to the lipid droplet and asterisk was used to draw inflammatory cells. Scale bar, 100 μm. Histopathological analysis of steatosis (e), hepatocyte ballooning (f), lobular inflammation (g), and the total NAFLD activity (h). Changes of serum ALT (i), and AST (j) were detected. The data are presented as mean ± SEM, n = 6, *$P < .05$ versus db/m, # $P < .05$ versus db/db.
3.2 Effects of Sch B treatment on hepatic pathology and hepatic function in db/db mice

HE staining and histology scores were used to analyze the steatosis and steatohepatitis in diabetic mice. As illustrated in Figure 2a–d, lipid droplets extensively accumulated in the liver tissues of db/db mice, but that Sch B decreased this accumulation. To further evaluate the histological injury in the liver, the scores were marked by a blinded histopathologist. Steatosis, hepatocyte ballooning, and inflammation scores were remarkably increased in db/db mice (P < .05; Figure 2e–g), consequently, NAFLD activity score was elevated compared with db/m mice (P < .05; Figure 2h). Sch B treatment reversed these abnormalities in db/db mice (P < .05). In addition, db/db diabetic mice displayed obvious liver damage reflected by elevations of serum ALT and AST (ALT: 74.5 ± 2.8 U/L vs. 24.2 ± 2.1 U/L, P < .05; Figure 2i; AST: 157.3 ± 4.6 U/L vs. 55.0 ± 3.3 U/L, P < .05; Figure 2j), which were restored by Sch B treatment (ALT: 54.3 ± 2.2 U/L vs. 74.5 ± 2.8 U/L, P < .05; Figure 2i; AST: 104.5 ± 3.5 U/L vs. 157.3 ± 4.6 U/L, P < .05; Figure 2j). There were no significant differences of both scores of hepatic pathology and hepatic function in db/m mice with or without Sch B administration (P > .05; Figure 2e–j).

3.3 Effects of Sch B treatment on hepatic steatosis in db/db mice

Oil Red O staining showed amount of lipid deposited in the liver in diabetic mice compared with db/m mice (Figure 3a–d). Oral administration of Sch B was largely attenuated hepatic steatosis in db/db mice. Furthermore, serum and liver tissues levels of TC and TG were significantly increased in db/db mice compared with db/m mice (TC in serum: 115.8 ± 2.1 vs. 56.8 ± 1.6 mg/dL, P < .05; Figure 3e; TG in serum: 116.8 ± 2.5 vs. 75.5 ± 2.3 mg/dL, P < .05; Figure 3f; TC in liver: 6.23 ± 0.10 vs. 2.53 ± 0.09 mg/g, P < .05; Figure 3g; TG in liver: 9.06 ± 0.21 vs. 5.09 ± 0.13 mg/g, P < .05; Figure 3h), which were effectively reduced by Sch B in db/db mice (TC in serum: 75.6 ± 2.0 vs. 115.8 ± 2.1 mg/dL, P < .05; Figure 3e; TG in serum: 89.1 ± 2.0 vs. 116.8 ± 2.5 mg/dL, P < .05; Figure 3f; TC in liver: 6.42 ± 0.18 vs. 9.06 ± 0.21 mg/g, P < .05; Figure 3g; TG in liver: 6.42 ± 0.18 vs. 9.06 ± 0.21 mg/g, P < .05; Figure 3h). These results were supported by an amelioration of mRNA levels of Sterol response element-bind protein 1c (SREBP-1c), fatty acid synthetase (Fasn), and acetyl-CoA carboxylase (ACC) in db/db mice receiving Sch B (SREBP-1c: 1.47 ± 0.02 vs. 3.02 ± 0.04, P < .05; Fasn: 1.41 ± 0.02 vs. 1.61 ± 0.02, P < .05; ACC: 1.40 ± 0.02 vs. 1.62 ± 0.02, P < .05).

**FIGURE 3** Attenuation of hepatic steatosis by Sch B treatment in db/db mice. (a–d) Representative images of oil red O staining (magnification, 200×). Scale bar, 100 μm. Levels of total cholesterol in serum (e), triglyceride in serum (f), total cholesterol in livers (g), and triglyceride in livers (h) were measured. The data are presented as mean ± SEM, n = 6, *P < .05 versus db/m, #P < .05 versus db/db.
± 0.03, \(P < .05\); ACC: 1.64 ± 0.04 vs. 2.27 ± 0.03, \(P < .05\); Figure 3i).

There were no significant differences on hepatic steatosis, serum lipid concentrations, and mRNA levels of SREBP-1c, Fasn, and ACC between db/m mice with Sch B group and db/m mice without Sch B group (\(P > .05\)).

3.4 | Sch B suppresses hepatocellular apoptosis in db/db mice

To further evaluate the influence of Sch B on liver injury in diabetic mice, TUNEL staining was used to detect the level of hepatocellular apoptosis. In db/db mice group, the liver tissues appeared significant apoptosis, reflected by more TUNEL-positive nuclei than db/m mice group (\(P < .05\); Figure 4a,b,d). As expected, db/db mice with Sch B group obviously suppressed TUNEL-positive cells when compared to db/db mice group (\(P < .05\); Figure 4b–d).

3.5 | Sch B alleviates inflammatory infiltration in liver tissues in db/db mice

Inflammation is involved in diabetic and lipid metabolism disorders (Kleiner et al., 2005), therefore, inflammatory infiltration in liver tissues were measured. Immunohistochemical staining for CD68 and F4/80 showed that accumulated Kupffer cells were markedly increased in liver tissues of db/db mice compared with db/m mice, while Sch B treatment significantly reduced the number of Kupffer cells in liver in db/db mice (\(P < .05\); Figure 5a–f,m). In line with these results, immunohistochemical staining for IL-1\(\beta\) and TNF-\(\alpha\) indicated that the levels of inflammatory cytokines were obviously upregulated in db/db mice compared with db/m mice (\(P < .05\); Figure 5g–m). In addition, concentrations of IL-1\(\beta\) and TNF-\(\alpha\) were also significantly increased in the liver tissues of db/db mice (\(P < .05\); Figure 5n). There was also a significant downregulation of IL-1\(\beta\) and TNF-\(\alpha\) expression in liver tissues by Sch B treatment (\(P < .05\); Figure 5g–n).

4 | DISCUSSION

The principal findings arose from the current study. First, the supplementation of Sch B significantly lowered blood glucose and improved insulin resistance in diabetic mice. Second, Sch B remarkably ameliorated steatosis, steatohepatitis and apoptosis in the liver in db/db mice. Furthermore, oral supplementation of Sch B downregulated Kupffer cells infiltrating and IL-1\(\beta\) and TNF-\(\alpha\) expressions in the liver.

It is well-established that NAFLD is the manifestation of metabolic syndrome in the liver, therefore, it is closely associated with obesity, dyslipidemia, insulin resistance, and T2DM (Filippatos et al., 2021; Hazer, 2020). The insulin resistance contributes to the progress of NAFLD through promoting the storage of FFAs in the liver (Shields et al., 2009). It has been reported that liver stellate cells incubated with high glucose or insulin overstimulated indirectly caused hepatic fibrosis (Paradis, 2001). NAFLD prevalence is dramatically
increased in insulin resistance population (Brunt, 2005). So far, lifestyle modification is regarded as the first-line interventions for NAFLD (Younossi et al., 2018). There is limited in terms of drug therapeutic options for diabetes mellitus-related NAFLD.

The Sch B is deemed to be beneficial for health, partly due to a broad spectrum of biological and pharmacological activities (Nasser et al., 2020). Recently, this compound has been identified to directly bind to peroxisome proliferator activator receptor gamma (PPARG) and protect against hepatic fibrosis through a network analysis (Xing et al., 2018). Likewise, the hepatoprotection afforded by Sch B is considered to be attributed to regulation of the hepatic (glutathione) GSH antioxidant enzymes (Ip et al., 1995). Moreover, Sch B alleviated

**FIGURE 5** Alleviation of inflammatory infiltration by Sch B treatment in the liver tissues of db/db mice. Liver sections were stained with CD68 (a-c), F4/80 (d-f), IL-1β (g-i), or TNF-α (j-l) to monitor inflammatory infiltration (magnification, 200×). Scale bar, 100 μm. (m) The bar graph showed that Sch B treatment significantly downregulated the infiltration of inflammatory cells and inflammatory cytokines in the liver tissues. (n) Concentrations of IL-1β and TNF-α in the liver tissues were measured using ELISA kits. The data are presented as mean ± SEM, n = 6, *P < .05 versus db/m, #P < .05 versus db/db.
Inflammatory adipokines such as IL-1β further deteriorating liver disease (Katsarou et al., 2020). Several pro-inflammatory Kupffer activation that triggers the inflammatory activation which (Katsarou et al., 2020; Lefere & Tacke, 2019). Excessive lipid accumulation is most critical mechanism related to the pathogenesis of NAFLD. Despite, both mRNA and protein levels of SREBP-1c and Fasn were upregulated in the liver of type 1 diabetes mellitus rats (Wang et al., 2013). In line with these results, we found that Sch B weakened mRNA expressions of SREBP-1c, Fasn, and ACC in the liver tissues of db/db mice. This result suggests that the effects of Sch B on hepatic steatosis under diabetic conditions may partly through inhibition of these lipid metabolism-related genes.

Growing evidence suggests that inflammation seems to be the most critical mechanism related to the pathogenesis of NAFLD, in db/db mice associating with suppression of the inflammatory response. (Katsarou et al., 2020). Hence, suppression of SREBP-1c obviously reversed hepatic steatosis in obese mice (Sekiya et al., 2003). Besides, both mRNA and protein levels of SREBP-1c and Fasn were upregulated in the liver of type 1 diabetes mellitus rats (Wang et al., 2013). In line with these results, we found that Sch B weakened mRNA expressions of SREBP-1c, Fasn, and ACC in the liver tissues of db/db mice. This result suggests that the effects of Sch B on hepatic steatosis and hepatic dysfunction in db/db mice.

5 | CONCLUSION
In summary, this study investigated the pharmacological effects of Sch B in type 2 diabetic mice, revealing that inhibition of liver inflammation by Sch B contributed to the amelioration in insulin resistance and lipid accumulation. Therefore, the inhibition of apoptosis and anti-inflammation activity of Sch B may be a promising target for prevention of T2DM-related NAFLD.

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CONFLICT OF INTEREST
The authors declare no conflicts of interests.

DATA AVAILABILITY STATEMENT
The data that support the findings of this study are available on request from the corresponding author. The data are not publicly available due to privacy or ethical restrictions.

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