Drug Repurposing in Alternative Medicine: Sochehwan, a Polyherbal Traditional Korean Digestant, Protects against Alcoholic Steatohepatitis by Regulating Cytochrome P450 2E1 Expression

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Abstract: Sochehwan (SCH) is an herbal prescription from traditional oriental medicine and is currently used to treat digestive ailments. In a previous study, SCH was found to have the potential to attenuate metabolic syndrome (MetS) by activating AMPK and downstream signaling. From the view of drug repurposing, the efficacy of SCH on alcoholic liver injury is implied in classic medical texts but is yet to be proven. C57BL/6J mice were pre-treated with SCH orally for 5 days and challenged by providing a pair-fed Lieber DeCarli diet containing alcohol for 20 days. Hepatic enzyme and triglyceride levels and endoplasmic reticulum (ER) stress-related markers were analyzed. Moreover, mitogen-activated protein kinases (MAPKs) and cytochrome P450 2E1 (CYP2E1) levels were determined. CYP2E1-transfected HepG2 cells were used to test the cytoprotective efficacy of SCH against the adverse effects of alcohol in vitro. In mice, SCH administration notably reduced hepatic enzyme activity and neural lipid levels. Furthermore, ER-stress markers and MAPK phosphorylation were reduced due to ROS suppression, which was attributed to decreased CYP2E1 expression in liver tissue. In addition, SCH successfully protected CYP2E1-transfected HepG2 cells against ethanol. Our findings suggest SCH attenuated alcohol-induced liver injury by inhibiting CYP2E1 expression and indicate drug repurposing should be considered as a valuable option for drug development in traditional herbal medicines.

Keywords: drug repurposing; Sochehwan; alcoholic liver injury; alcoholic liver disease; ER stress; CYP2E1; MAPKs; inflammation

1. Introduction

Excessive alcohol consumption is considered a major etiology of liver-related diseases [1]. Alcoholic liver disease (ALD) (also referred to as alcohol-related liver disease, ARLD) is widely acknowledged by hepatologists to be a pandemic and presents serious economic burdens and health issues worldwide [2,3]. ALD includes a wide-ranging spectrum of alcohol-induced liver damage from simple steatosis, to alcoholic hepatitis, and hepatic cirrhosis and fibrosis [4].

As a member of the classical cytochrome P450 family (CYP 450), CYP2E1 is regarded as a major source of alcohol- or drug-induced reactive oxygen species (ROS) in the liver [5]. CYP2E1 is directly involved in alcohol metabolism and converts ethanol into acetaldehyde [6]. Ethanol breakdown by cytochrome P450 is referred to as the microsomal ethanol oxidizing system (MEOS), and this system generates ROS as a by-product, which causes a
transient intracellular imbalance in redox status [7]. Furthermore, CYP2E1 causes significantly higher NADPH activity than other members of the cytochrome P450 family and can have deleterious effects on mitochondrial function [8].

Numerous authors have suggested that the inhibition of CYP2E1 should be considered a core pharmaceutical target to address the harmful effects of xenobiotic exposure [9,10]. The precise pathophysiological roles of specific CYP450 isoforms in alcoholic steatohepatitis remain unclear; however, alterations in the expression of CYP2E1 have been implicated in liver diseases [11]. In addition, CYP2E1-related endoplasmic reticulum (ER) stress and the cellular inflammatory response with signaling molecules are emerging as potent aggravating factors of steatohepatitis [12].

Drug repurposing, also referred to as drug repositioning, is now recognized as common practice in developing new agents in pharmaceutical industries [13,14], and is also applicable in liver-related diseases [15]. Based on the well-studied pharmacokinetic profiles and safety data from clinical trials, it provides an innovative means of discovering new indications for existing drugs, which can save cost, time, and resources [16,17]. Until fairly recently, drug repurposing of herbal medicine was barely discussed. However, interest in herbal prescriptions as sources of novel natural-product drugs has increased dramatically [18].

SCH is a centuries-old traditional herbal digestant prescription consisting of *Pharbitis Semen*, *Trogopterori Faeces*, and *Cyperi Rhi zona*, and is currently commercialized as an over-the-counter drug for its efficacy and safety [19]. Although the prescription has been well-recognized for its efficacy on digestive disorders, we further expected it has an effect on a wide range of diseases associated with diet.

In a previous study, we showed that SCH has multifaceted effects on metabolic syndromes by activating the AMP-activated protein kinase (AMPK) and its downstream signaling in the hepatic steatosis model in vitro (concentration at 50 µg/mL) and in vivo (administrated 200 mg/kg/day) [19]. Pathological marker analysis in an NAFLD mouse model revealed reduced hepatic enzyme activities and serum pro-inflammatory markers, which implied that SCH might be active against steatohepatitis. Moreover, the description of SCH in classic medical texts implies its efficacy against alcohol-related digestive disorders [20]. However, the efficacy and mode of action of SCH on alcoholic steatohepatitis have yet to be investigated.

We investigated the efficacy of SCH on alcoholic steatohepatitis using an in vitro CYP2E1-transfected HepG2 cell model and a pair-fed alcohol-induced liver injury mouse model (in vivo). Various mechanisms and significant markers of SCH on alcohol metabolism via CYP family members, antioxidant enzymes, and endoplasmic reticulum stress (ER stress) were explored. By scrutinizing the repurposing process and data we produced, we review the application of drug repurposing in traditional herbal medicine.

2. Materials and Methods

2.1. Antibodies

Antibodies against c-Jun N-terminal kinases (JNK, no. 9258S), p-JNK (no. 9251S), p38 mitogen-activated protein kinases (p38, no. 9212S), p-p38 (no. 9211S), interleukin-6 (IL-6, no. 12153S), and tumor necrosis factor alpha (TNF-α, no. 3707S) were obtained from Cell Signaling Technology (Beverly, MA, USA). Antibodies specific for cytochrome P450 2E1 (CYP2E1), inositol-requiring enzyme 1 alpha (IRE-1α, no. sc-390960), C/EBP-homologous protein (CHOP, no. sc-7351), activating transcription factor 6 alpha (ATF6α, no. sc-166659), extracellular signal-regulated kinase (ERK, no. sc-94), and p-ERK (no. sc-7383) were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). The β-Actin antibody was from Santa Cruz Biotechnology, and horseradish peroxidase (HRP)-conjugated secondary antibodies were supplied by Santa Cruz Biotechnology.
2.2. Preparation of SCH Sample

All herbs were purchased from Humanherb (Gyeongsangbuk-do, Korea). *Pharbitis Semen*, *Trogopterorum Faeces*, and *Cyperi Rhizoma* (2:1:1, w/w) were extracted in distilled water for 3 h at 95 °C by reflux extraction, as previously described [19]. The hot water extract obtained was filtered twice through 8 μm-pore-size Whatman filter paper, rotary evaporated, and freeze-dried to obtain lyophilized SCH (yield 15.53%, referred to hereafter as SCH). SCH was stored at −20 °C until required.

2.3. Animals

Eight-week-old male C57BL/6J mice (n = 24) were purchased from Orientbio (Gyeonggi-do, South Korea) and fed a Lieber-DeCarli (LDC) liquid diet (Supplementary Table S1). Mice were housed in a temperature- and humidity-controlled environment under a 12 h dark/light cycle and given water ad libitum. After 4 days of acclimatization, mice were randomized into four groups of six mice: Group 1’s (pair-fed, PF) diet did not contain alcohol, but mice were pair-fed the same calories of the LDC diet that mice in the alcohol-induced group consumed during the experimental period. Mice in group 2 (alcohol-fed, AF), group 3 (alcohol-fed with SCH, AF+SCH), and group 4 (alcohol-fed with silymarin, AF+PC) received the LDC diet containing ethanol (5%) for 3 weeks with an ethanol binge (on the 15th and 21st days of the experiment, 10% and 20%, w/w, respectively) (Supplementary Figure S1). Groups 3 and 4 were administered SCH (100 mg/kg body weight) or silymarin (50 mg/kg body weight), respectively, once a day for the first and last 5 days of the experimental period by gavage. Bodyweights were checked daily from the beginning of the feeding period. After sacrifice on experimental day 21, whole blood was collected from abdominal aortas, and livers were harvested under zoletil anesthesia, as previously described [21]. Animal experiments were conducted in accordance with the guidelines issued by Dongguk University Ethics Committee (Approval No. IACUC-2017-026-1), which also approved the study protocols before study commencement.

2.4. Serum Biochemistry

Aspartate aminotransferase (AST), alanine aminotransferase (ALT), and triglyceride (TG) were measured in mouse sera. Serum hepatic enzymes were assayed using a commercial ALT, AST, and TG colorimetric test kit (Asan Pharmaceutical, Seoul, Korea) according to the manufacturer’s protocols with slight modification [9]. Optical densities (OD) were measured using a microplate reader (VersaMax, Molecular Devices, CA, USA).

2.5. mRNA Isolation and Real-Time Polymerase Chain Reaction (qPCR)

Total mRNA was extracted from mouse liver tissues using the Trizol reagent (Thermo Scientific, Waltham, MA, USA) according to the manufacturer’s instructions. mRNA quantities and integrities were checked to ensure sample qualities. The relative expression of CYP2E1 mRNA was determined by quantitative real-time PCR (qPCR). Briefly, reverse transcription was conducted using AccuPower RT PreMix (Bioneer, Daejeon, Korea) and oligo dT (18) primers (Invitrogen, Carlsbad, CA, USA). The amplification of cDNA was performed using target-specific primers and a LightCycler 480 PCR system (Roche Applied Science, Basel, Switzerland) and the following program: Initial denaturation 10 min at 95 °C followed by 45 cycles of denaturation at 95 °C for 10 s, annealing at 52 °C for 20 s, and extension at for 72 °C for 20 s. The following primers were used: CYP2E1 sense 5′-CCAGTCCAGGACCTCATTT-3′ and antisense 5′-CACCTTTGCACAAGTCTCT-3′, β-actin sense 5′-GCAAGTGGCTTCTAGCGGAC-3′, and antisense 5′-AAGAAAGGTTGTTAAAAACGCCACG-3′ were used as the internal control. Results were normalized using Ct values of β-actin gene expression.

2.6. Cell Culture and CYP2E1 Transfection

HepG2 human hepatocellular carcinoma cells were obtained from the Korea Cell Line Bank (KCLB, Seoul, Korea) and cultured in Dulbecco’s Modified Eagle Medium
(DMEM) (Hyclone, Logan, UT, USA) supplemented with 10% fetal bovine serum (FBS) (Invitrogen, Carlsbad, CA, USA), 100 U/ml penicillin, and streptomycin. Cells were plated and incubated in a humidified 5% CO$_2$ environment at 37 °C. For CYP2E1 gene transfection, HepG2 cells were seeded at $1 \times 10^6$ cells/well on 6-well plates, and 24 h later, the culture medium was replaced with FBS-free opti-MEM, and cells were transfected with the human CYP2E1 plasmid containing the T7 promoter and the myc-DDK tag (OriGene Technologies Inc., Rockville, MD, USA) using the Lipofectamine 3000 (Invitrogen) transfection reagent according to the manufacturer’s instructions. In brief, 96 µL of OPTI-MEM (Invitrogen) including 4 µL of the transfection reagent and 2 µg of plasmid DNA were added to each culture well of 6-well plates and incubated at 37 °C for 6 h. After transfection, the culture medium was replaced, and SCH and different concentrations (0–500 mM) of pure ethanol were treated for 72 h (media were changed every 24 h). CYP2E1-transfected HepG2 cells did not show any morphological changes and no significant cytotoxicity was observed.

2.7. Western Blot Analysis

Total proteins were isolated using the PRO-PREP tissue protein extraction buffer (iNtRON Biotechnology, Burlington, MA, USA) and quantified using the BCA protein assay kit (Thermo Scientific). Briefly, samples were washed with ice-cold DPBS and lysed using a protein extraction buffer containing a protease inhibitor and phosphatase inhibitor cocktail (GenDEPOT, Barker, TX, USA). Equal amounts of proteins (25 µg/lanes) mixed with the Lane Marker Reducing sample buffer (Thermo Scientific) were loaded onto 10% SDS-PAGE, separated, and transferred to polyvinylidene difluoride (PVDF) membranes (GE Healthcare, Buckinghamshire, BK). Membranes were blocked with 5% skim milk in Tris-buffered saline contacting 0.1% Tween (TBST) for 1 h, incubated overnight at 4 °C with primary antibodies (1:1000) diluted in cold TBST containing 3% skim milk, washed with TBST three times, and incubated with horseradish peroxidase (HRP)-conjugated secondary antibodies (1:2000 dilutions in TBST containing 1% skim milk) at room temperature for 1 h. Blots were detected using a Fusion Solo imaging system (Vilber Lourmat, Marne-la-Vallée, France), and results were visualized using a SuperSignal West Pico PLUS chemiluminescent substrate (Thermo Scientific) and normalized using Bio-1D software (Vilber Lourmat) versus β-actin.

2.8. Statistical Analyses

The results were analyzed using Graph Pad Prism version 5.0 (Graph Pad, La Jolla, CA, USA). Standard curves were plotted using Excel (Microsoft, Redmond, WA, USA). Analysis of variance and One-Way ANOVA with Dunnett’s multiple comparison tests were used to determine the significances of differences. Results are presented as means ± SDs, and statistical significance was accepted for $p$ values < 0.05.

3. Results

3.1. Body Weights and Serum Analysis

At sacrifice day, no difference was observed between mean body weights in the three alcohol-administered groups (AF, AF+SCH, AF+PC) and PF group (Figure 1a). However, the mean liver weight/body weight ratio of AF mice was significantly greater than in the PF group, and this ratio was markedly reduced in the AF+SCH and AF+PC groups (Figure 1b). As shown in Figure 1c,d, SCH significantly decreased plasma glutamic oxaloacetic transaminase (GOT) and glutamic pyruvic transaminase (GPT) by 4.13 and 3.75 IU/L, respectively (Figure 1c,d). Serum TG levels were also significantly lower in the alcohol+SCH group than in the AF+SCH group (Figure 1e). Serum GOT, GPT, and TG were also lower in the AF+PC group than in the alcohol group.
Figure 1. SCH protected mouse livers from alcohol-induced hepatic injury. (a) Weekly body weight changes in the study groups normal pair-fed controls (PF), alcohol-fed group (AF), alcohol-fed and SCH administered group (AF+SCH), and alcohol-fed mice and silymarin administered group (AF+PC). (b) Liver weight/body weight ratio. (c) GOT, (d) GPT, (e) TG levels in mouse serum. Results are presented as means ± SDs. ### \( p < 0.001 \) versus non-alcohol fed controls, and * \( p < 0.05 \), ** \( p < 0.01 \), *** \( p < 0.001 \) versus the alcohol-fed group.

3.2. Reduction of Endoplasmic Reticulum (ER) Stress

IRE1, an important ER stress sensor, was significantly induced in the liver in the alcohol group \( (p < 0.005) \) and was reduced by SCH administration more so than by silymarin (Figure 2a,b). To investigate the anti-apoptotic effect of SCH, we performed immunoblot analysis for the cell signals ATF6 and CHOP. Densitometry showed the ethanol diet increased these protein levels by 3.77- and 2.84-fold compared with non-alcohol-fed controls (Figure 2c,d). Silymarin treatment significantly reduced these increases \( (p < 0.005 \) and \( p < 0.01 \), respectively). Similarly, SCH administration also remarkably decreased these alcohol-induced increases in ATF6 and CHOP protein levels \( (p < 0.01 \) and \( p < 0.05 \), respectively).

3.3. Regulation of Antioxidative Enzyme Expression

Antioxidative enzyme gene expressions of catalase (CAT), superoxide dismutase (SOD), and glutathione peroxidase (GPx) were evaluated in mouse livers. The expressions of all three genes were significantly upregulated in the three alcohol-administered groups (Figure 3a–d). However, their expressions were significantly lower in the AF+SCH and AF+PC groups than in the alcohol group, indicating that ROS levels were relatively resolved in these groups.
Figure 2. SCH inhibited endoplasmic reticulum stress in mice with alcohol-induced hepatic injury. (a) Representative immunoblot images of IRE-1, ATF6, and CHOP. (b) Inositol-requiring enzyme 1 (IRE-1) expressions levels, (c) activating transcription factor6 (ATF6) levels, and (d) C/EBP-homologous protein (CHOP) levels as determined by Western blot. Band intensities were measured by densitometry and normalized versus β-actin. Results are presented as the means ± SDs of three different experiments. # p < 0.05, ## p < 0.01, ### p < 0.001 versus non-alcohol fed controls, and * p < 0.05, ** p < 0.01, *** p < 0.001 versus alcohol fed mice.

Figure 3. SCH upregulated antioxidant enzyme expressions in mice with alcohol-induced hepatic injury. (a) The expressions of SOD1, (b) SOD2, (c) CAT, and (d) GPx were determined by qPCR. Relative gene expression levels were normalized versus β-actin. Results are presented as means ± SDs. # p < 0.05, ## p < 0.01, ### p < 0.001 versus non-alcohol fed controls, and * p < 0.05, ** p < 0.01 versus the alcohol fed group.
3.4. Regulation of MAPK Signals and Cytokine Production

The MAPK signaling pathway was investigated in liver tissues to investigate the effect of SCH on alcohol-induced steatohepatitis. MAPKs, including ERK, JNK, and p38, were significantly \( p < 0.005 \) phosphorylated in ethanol-fed mice, which indicated stress-induced MAPK activation was associated with hepatic damage. As shown in Figure 4a, silymarin treatment successfully blocked MAPK phosphorylation to the normal control level. Moreover, alcohol-induced ERK, JNK, and p38 phosphorylation were significantly suppressed by SCH or silymarin (Figure 4b–d).

Subsequently, we found SCH prevented ethanol-induced pro-inflammatory cytokine production. Ethanol-fed mice showed 4.10- and 2.07-fold increases in IL-6 and TNF-\( \alpha \) expression as compared with normal controls, and SCH significantly \( p < 0.005 \) reduced this IL-6 increased to 0.21-fold, whereas silymarin reduced it to 0.46-fold (Figure 5b). Furthermore, densitometry showed SCH dramatically reduced alcohol-induced TNF-\( \alpha \) production, whereas silymarin had little effect (Figure 5c).

3.5. Reduction of CYP2E1 Expression in Mouse Liver Tissue

To investigate the mechanism responsible for the effect of SCH, we performed immunoblot analysis and qPCR for the CYP2E1 protein and mRNA. CYP2E1 protein levels were 45.36-fold higher in the alcohol-fed group than in normal controls \( p < 0.005 \). SCH administration significantly \( p < 0.05 \) reduced these CYP2E1 protein and mRNA level increases (Figure 6a–c), and silymarin also significantly reduced these increases.
Figure 5. SCH inhibited pro-inflammatory cytokine production in alcohol-fed mice. (a) Representative immunoblot images of IL-6 and TNFα levels. (b) IL-6 and (c) TNFα levels as determined by Western blot. Band intensities were measured by densitometry and normalized versus β-actin. Results are presented as the means ± SDs of three different experiments. ### p < 0.001 versus non-alcohol fed controls, and * p < 0.05, *** p < 0.001 versus the alcohol group.

Figure 6. SCH inhibited induction of the CYP2E1 gene in alcohol-fed mice. (a) Cytochrome P450 2E1 (CYP2E1) levels as determined by Western blot, and band intensities determined by densitometry and normalized versus β-actin. (b) CYP2E1 gene expression as determined by real-time PCR. (c) CYP2E1 levels were determined by Western blot, and band intensities were measured by densitometry and normalized versus β-actin. Results are presented as the means ± SDs of three different experiments. ## p < 0.01, ### p < 0.001 versus normal controls, and * p < 0.05, ** p < 0.01 versus the alcohol-fed group.
3.6. Reduction of Ethanol-Induced Injury in CYP2E1-Transfected HepG2 Cells

To evaluate the protective effect of SCH against ethanol, we transfected the CYP2E1 plasmid into hepatocellular carcinoma HepG2 cells and exposed them to ethanol concentrations in culture media. CYP2E1-transfected HepG2 cells proliferated less than non-transfected intact cells, which was attributed to the toxicity of liposome as a transfection vehicle. In addition, the comparison between CYP2E1-transfected HepG2 cells with different ethanol concentrations demonstrated that ethanol at a concentration of 500 mM significantly decreased cell viability. However, when these CYP2E1-transfected HepG2 cells were co-treated with ethanol (500 mM) and SCH, the cell toxicity of the ethanol binge was removed, at concentrations of 25 and 50 µg/mL (p < 0.05) (Figure 7b).

Figure 7. SCH protected CYP2E1-transfected HepG2 cells from ethanol. (a) Cell viabilities of non-transfected HepG2 cells and CYP2E1-transfected HepG2 cells exposed to different concentrations of ethanol (0-500 mM) for 72 h. (b) Cell viability of CYP2E1-transfected HepG2 cells in the presence of ethanol (500 mM) with or without SCH for 72 h. Results are presented as the means ± SDs of three different experiments. # p < 0.05, ## p < 0.01, ### p < 0.001 versus non-treated intact control group (a and b), and * p < 0.05, ** p < 0.01 versus CYP2E1-transfected HepG2 cells (a) and CYP2E1-transfected HepG2 cells treated with 500 mM ethanol (b).

4. Discussion

In the present study, we investigated the effect of SCH on alcoholic steatohepatitis with various molecular markers or signaling pathways as its mechanisms. The NIAAA model used in this study is a representative alcoholic liver injury model involving chronic ethanol feeding and binge ethanol administration and induces the major hallmarks of human ALD [22].

GOT and GPT activity levels (widely used parameters of hepatic damage [23]) were determined in mice sera after sacrifice, and not surprisingly, the results obtained indicated alcohol feeding markedly increased hepatic enzyme activities (p < 0.001), which may reflect the development of alcoholic hepatitis in murine models [24]. Notable changes in the liver weight/body weight ratio and increases in serum triglyceride also implied steatotic changes in ethanol-fed mice [25,26]. As SCH administration was found to successfully block these changes, we further tried to deduce a reason for the effect regarding the pathological mechanism of alcoholic steatohepatitis.

Steatosis related to chronic alcohol consumption is caused by dysregulated lipid metabolism [27]. In brief, ethanol exposure reduces AMPK, sirtuin1 (SIRT1), and peroxisome proliferator activated receptor alpha (PPARα) activities and upregulates the expressions and activities of sterol regulatory element binding protein-1c (SREBP-1c), acetyl-CoA carboxylase 1 (ACCI), and peroxisome proliferator activated receptor gamma (PPARγ), which promote lipogenesis [27]. A previous study conducted on SCH showed SCH can activate AMPK and modulate its downstream signals, including ACC and PPARγ, in the liver [19], which may partially explain the suppression of alcohol-induced hepatosteatosis by SCH.
ER stress is induced during ethanol metabolism and is related to the accumulation of unfolded/misfolded proteins [28]. Under prolonged or severe ER stress, the unfolded protein response (UPR) can become cytotoxic, and even induce apoptosis, rather than cytoprotective [29]. Inositol-requiring enzyme 1α (IRE1α), the C/EBP-homologous protein (CHOP), and activating transcription factor 6 (ATF6) are predominant and unique signaling transduction pathways of UPR [30] and can detect abnormal conditions in the ER lumen and transmit signals to activate nuclear transcription factors [31]. We observed SCH administration reduce alcohol-induced IRE1α, CHOP, and ATF6 upregulations.

Alcohol-induced ER stress had a pro-inflammatory effect by activating the NF-κB and JNK signaling pathways [28]. IRE1α activates apoptotic signaling kinase-1 (ASK-1), and thus, activates downstream factors such as JNK (Jun-N-terminal kinase) and p38 MAPK (p38 mitogen-activated protein kinase), which enhance apoptosis in the liver. Moreover, activation of the MAPK pathway induces transcription factor activation [32], which in turn promotes macrophage infiltration and inflammatory cytokine and chemokine production, which lead to liver damage [33]. However, SCH inhibited the alcohol-induced phosphorylation of JNK, ERK, and P38 in the MAPK pathway in alcohol-fed mice.

Alcohol-induced liver disease is fundamentally associated with a state of excessive oxidative stress [34]. The metabolism of alcohol by alcohol dehydrogenase (ADH) and cytochrome P450 2E1 (CYP2E1) leads to the formation of acetaldehyde [6]. CYP2E1 metabolizes and activates ethanol to produce more reactive, toxic products, and thus, increases ROS production, especially the production of H2O2 and the superoxide anion (O2−), which causes cell damage [35]. ROS, a by-product formed during this process, can also activate MAPK family members, and therefore, contribute to the development of ALD [36]. However, SCH inhibited oxidative stress and ROS production in liver tissue by reducing CYP2E1 expression, and thereby, suppressed ER stress and reduced lipid accumulation and the inflammatory response, which suggests SCH might alleviate or prevent alcoholic liver injury or steatohepatitis.

Studies conducted by our team consistently suggest that the application of drug repurposing to traditional herbal medicine provides a potent means of discovering new natural-product drugs because these medicines represent a huge resource of medically interesting natural products with multi-target characteristics [37,38]. There are even more possibilities of new drug developments in herbal concoctions, with numerous candidates and accumulated experience of medical usages. Furthermore, the application of up-to-date methodologies of drug repurposing such as network pharmacology and molecular docking simulation provides a means of rapidly obtaining more objective, practical, and valuable results.

5. Conclusions

In summary, SCH did not affect the body weights in alcohol-fed mice but significantly reduced liver weights to the normal control level. SCH administration reduced alcohol-induced upregulations of ER-stress markers (IRE1α, ATF6, and CHOP), which indicated it reduced alcohol-induced ER stress. Furthermore, SCH inhibited alcohol-induced MAPK signaling and inflammatory responses. We believe that these effects of SCH were caused by its suppressive effect on CYP2E1 expression. Our findings suggest SCH might have therapeutic effects on metabolic syndromes and alcoholic liver injury.

Supplementary Materials: The following are available online at https://www.mdpi.com/article/10.3390/pr9101760/s1, Figure S1: The schematic diagram of animal experimental design of alcohol-induced steatohepatitis in C57BL/6J mice. Table S1: Composition of the standard Lieber-DeCarli liquid diet used in this study.

Author Contributions: G.-R.Y. performed the animal surgery and wrote the manuscript. S.-J.L., D.-W.L. and H.K. reviewed the study design and results. J.-E.K. designed the experiments and analyzed the data. W.-H.P. supervised the project and contributed to the final draft of the paper. All authors have read and agreed to the published version of the manuscript.
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