

Total C-21 steroidal glycosides, isolated from the root tuber of *Cynanchum auriculatum* Royle ex Wight, attenuate hydrogen peroxide-induced oxidative injury and inflammation in L02 cells

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Abstract. Oxidative stress plays an important role in the pathology of liver disorders. Total C-21 steroidal glycosides (TCSGs), isolated from the root tuber of *Cynanchum auriculatum* Royle ex Wight, have been reported to exert numerous effects, including liver protective and antioxidant effects. In order to investigate the potential mechanisms underlying the protective effects of TCSGs on liver function, the present study used the human normal liver cell line, L02, to evaluate the effects of TCSGs on hydrogen peroxide (H₂O₂)-induced oxidative injury and inflammatory responses. The L02 cells were pretreated with various concentrations of TCSGs, followed by exposure to 1.5 mM H₂O₂. Cell viability was determined by a 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay. The levels of alanine aminotransferase (ALT), aspartate aminotransferase (AST), lactate dehydrogenase (LDH) and nitric oxide (NO) were measured using colorimetric assays. The activities of superoxide dismutase (SOD), catalase (CAT), glutathione peroxidase (GSH-Px) and the production of malondialdehyde (MDA) were also determined. Intracellular reactive oxygen species (ROS) levels were detected using a fluorescent probe.

H₂O₂-induced oxidative toxicity was attenuated following treatment with TCSGs, as indicated by the increase in cell viability, the decreased levels of ALT, AST, LDH, NO, MDA and ROS, and the increased activities of SOD, CAT and GSH-Px. To further explore the possible mechanisms of action of TCSGs, the nuclear factor erythroid 2-related factor 2 (Nrf2) and nuclear factor-κB (NF-κB) pathways were examined. The results revealed that treatment with TCSGs markedly induced Nrf2 nuclear translocation and upregulated the expression of heme oxygenase-1 (HO-1) in the L02 cells damaged by H₂O₂. In addition, pretreatment with TCSGs inhibited the NF-κB signaling pathway by blocking the degradation of the inhibitor of nuclear factor κBα (IκBα), thereby reducing the expression and nuclear translocation of NF-κB, as well as reducing the expression of tumor necrosis factor-α (TNF-α), interleukin-6 (IL-6), inducible nitric oxide synthase (iNOS) and cyclooxygenase 2 (COX-2). On the whole, the findings of this study demonstrate that TCSGs can protect L02 cells against H₂O₂-induced oxidative toxicity and inflammatory injury by increasing the expression of Nrf2 and HO-1, mediated by the NF-κB signaling pathway.

Introduction

Cynanchum auriculatum (*C. auriculatum*) Royle ex Wight, a member of the *Asclepiadaceae* family, is widely distributed in China. The root tuber of *C. auriculatum* Royle ex Wight, a well-known traditional Chinese herbal medicine known as 'Baishouwu' has been used as a local tonic and medicine for >1,000 years since the Tang Dynasty in China (1). Modern phytochemical and pharmacological studies have demonstrated that C-21 steroidal glycosides are the major active components of Baishouwu (2,3). The total C-21 steroidal glycosides (TCSGs), isolated from Baishouwu, possess various pharmacological activities, including antitumor (4-11), aging-attenuating (12), free radical-scavenging (13), immunity-enhancing (14), depression-reducing (15) and fungus-suppressing (16) activities. Recently, it has been reported that the C-21 steroidal

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glycosides isolated from Baishouwu exhibit notable hepatoprotective effects *in vivo* (17). However, the underlying mechanisms remain largely unknown.

Pathological and experimental evidence has suggested that multiple mechanisms of hepatic injury are implicated in oxidative damage, inflammation, the dysfunction of intracellular targets and the innate immune system (18-20). It has been well established that cell oxidative stress damage induced by reactive oxygen species (ROS) is a principal mechanism of hepatic injury. When there is an imbalance in the levels of intracellular oxidative factors and antioxidants, oxidative stress can result in a disruption in redox signaling and cellular injury (21,22). An increasing body of evidence has indicated that superoxide anion and hydrogen peroxide (H_2O_2) are associated with various pathological diseases, such as viral hepatitis (23), alcoholic hepatitis (24) and non-alcoholic fatty liver diseases (NAFLD) (25). H_2O_2 -induced hepatic injury is a common cell model for investigating the potential hepatoprotective activity (26). Lipid peroxidation is one of the significant causes of H_2O_2 -induced hepatic injury and can be monitored by detecting the content of intracellular malondialdehyde (MDA). The disruption of the hepatic antioxidant defense system is characterized by increased MDA and/or altered enzymatic antioxidants, including superoxide dismutase (SOD), catalase (CAT) and glutathione peroxidase (GSH-Px). The superoxide radical (O_2^-) is an oxygen free radical that damages the body, which is then converted to O_2 and H_2O_2 by the action of SOD and detoxified to water by CAT or GSH-Px. The activities of these antioxidants have been used to evaluate oxidative stress levels in cells (27). Excessive ROS levels induced by H_2O_2 disrupt the balance between ROS production and the antioxidant defense system.

It has been reported that excessive ROS levels induced by H_2O_2 can induce nuclear factor (NF)- κ B activation, and subsequently increase NF- κ B p65 subunit nuclear translocation (28). The activation of NF- κ B is enhanced by increasing the degradation and phosphorylation of inhibitor of nuclear factor- κ B (I κ B), which further modulates the hepatic injury by regulating proinflammatory cytokine production, such as tumor necrosis factor (TNF)- α and interleukin (IL)-6, and the expression of inflammatory mediators including inducible nitric oxide synthase (iNOS) and cyclooxygenase (COX)-2. Nitric oxide (NO) plays a critical role in hepatic injury caused by H_2O_2 . H_2O_2 exposure generates excessive levels of NO via the activation of iNOS, thereby leading to hepatic tissue damage (29).

In addition, a previous study revealed that the nuclear factor erythroid 2-related factor 2 (Nrf2) signaling pathway played an important role in the intracellular defense against oxidative stress (30). The nuclear transcription factor Nrf2 binds to antioxidant response elements (AREs) in order to activate antioxidant genes that are involved in the elimination of free radicals by promoting the expression of antioxidant enzymes, such as SOD and CAT (31). The role of the Nrf2/ARE signaling pathway in liver disease pathogenesis and its possible application as a underlying therapeutic target to block and treat chronic hepatitis, and alcoholic as well as non-alcoholic hepatic injury, have been extensively investigated (32).

The critical roles of oxidative stress and inflammation in the initiation and progression of liver injury have attracted

increasing levels of attention (33). Previous studies have reported that H_2O_2 -derived ROS production induced oxidative stress and promoted the generation of proinflammatory cytokines via the Nrf2 and NF- κ B signaling pathway (34-37). Thus, redox-sensitive transcription factors, such as Nrf2 and NF- κ B, are essential transcription factors that modulate an array of antioxidant responses and proinflammatory gene expression in the liver (38).

Natural products derived from herbal medicines are attracting increasing attention as alternative treatment options for the prevention and treatment of liver diseases. TCSGs, the most important active components of Baishouwu, are commonly considered to be strong antioxidants due to their high free radical scavenging properties. Recent studies have revealed that TCSGs affect cellular inflammatory stress and inhibit tumors by regulating carcinogenesis-associated processes (3,9,10,39). In addition, it has been demonstrated that TCSGs exert hepatoprotective effects against liver injury induced by carbon tetrachloride *in vivo* (17). To date, few studies have provided conclusive results, particularly with regard to the molecular mechanism of TCSGs in protecting hepatocytes against oxidative damage and inflammatory response.

The aim of the present study was to explore the possible protective effects of TCSGs on human liver cells against H_2O_2 -induced oxidative damage by detecting oxidative stress indicators and inflammatory markers. More specifically, the potential molecular mechanisms of action of TCSGs were investigated by examining antioxidant and inflammatory signaling pathways. These results may elucidate the underlying protective mechanisms of TCSGs in liver injury.

Materials and methods

Plant material and preparation of TCSGs. The peeled root tubers of *C. auriculatum* were collected from Binhai County (Jiangsu, China) in December, 2015. The materials were identified and authenticated by Professor Shi-Hui Qian (Jiangsu Province Academy of Traditional Chinese Medicine, Jiangsu, China). A voucher specimen (no. WFC-20151225) was deposited at the Department of Natural Product Chemistry, Jiangsu Province Academy of Traditional Chinese Medicine (Nanjing, China). Chemical reference substances, namely caudatin-2,6-dideoxy-3-O-methyl- β -D-cymaropyranoside (no. 1), wilfoside C_1N (no. 2), caudatin (no. 3), wilfoside K_1N (no. 4), wilfoside C_1G (no. 5), cynauricuoside A (no. 6), cynauricuoside C (no. 7) and auriculoside IV (no. 8) (Fig. 1), were isolated from *C. auriculatum* (40) and identified based on infrared, ultraviolet, mass spectrometry (MS) and nuclear magnetic resonance spectroscopic analyses (data not shown). The purity of these compounds was determined to be >98% by high-performance liquid chromatography (HPLC) analysis. The root tubers were cut into small sections and boiled in 95% ethanol (1:10) 2 times, 2 h each time, filtered through gauze and concentrated under reduced pressure to produce the ethanol extract. The ethanolic extract was extracted with ethyl acetate 3 times and then merged and evaporated to obtain the TCSG extract, as previously described (41). The main chemical components of the TCSGs were identified and determined by the ultra high-performance liquid chromatography triple quadrupole tandem mass spectrometry

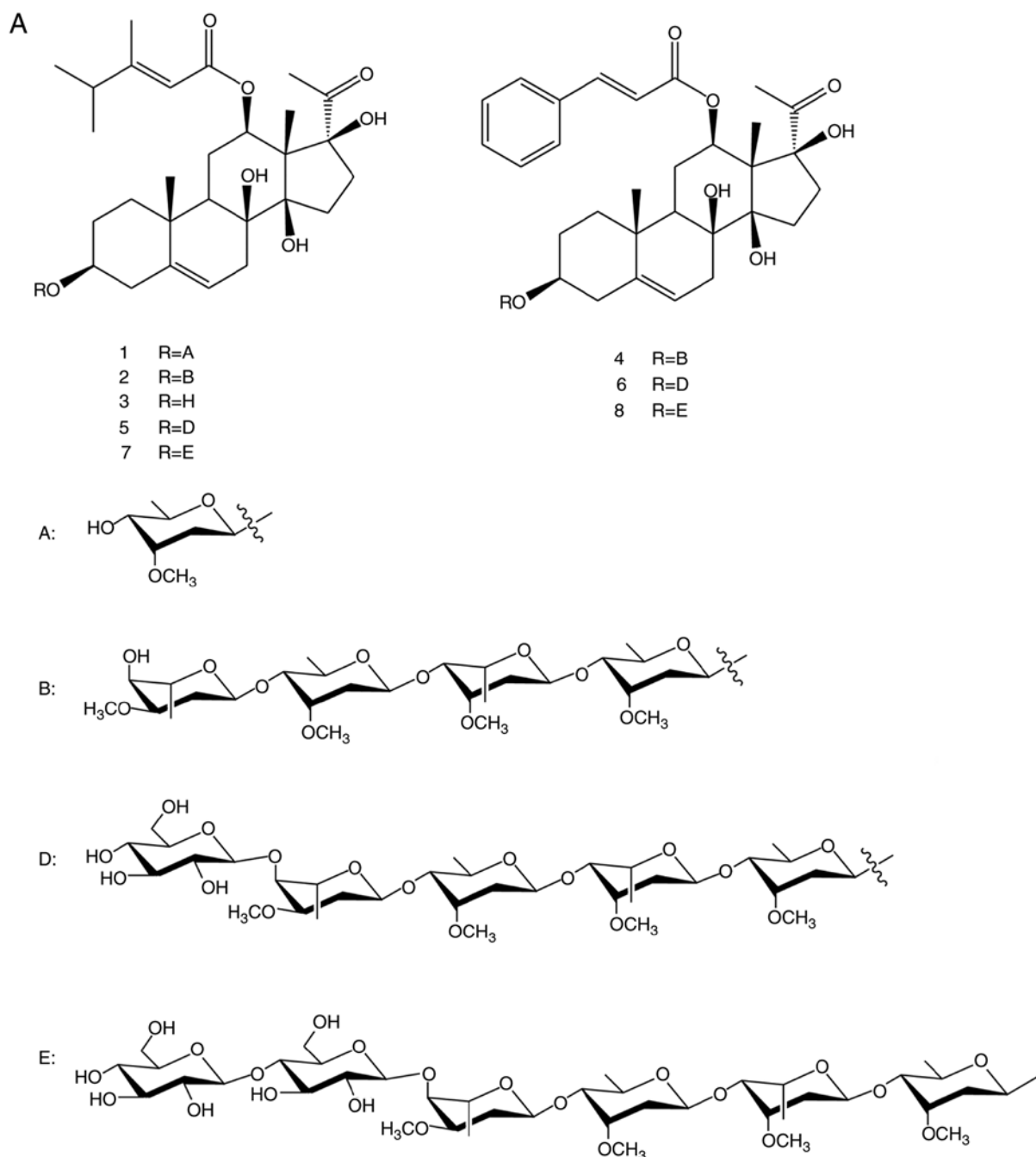


Figure 1. (A) Chemical structures of the 8 compounds.

method according to previously reported protocols (41). An Acquity UHPLC BEH C_{18} column (2.1x100 mm, 1.7 μ m) was used for separations. The mobile phase was composed of (A) water [0.1% (v/v) formic acid] and (B) acetonitrile, and a linear gradient elution was used. MS was operated using an electrospray ionization (ESI) source in negative mode and the ESI-MS spectra were acquired by multiple reaction monitoring (MRM). Chlorzoxazone was used as the internal standard. A series of concentrations of standard solution were prepared for the establishment of the calibration curves. It was revealed that the TCSGs mainly contained 8 C-21 steroidal glycoside components, including caudatin-2,6-dideoxy-3-O-methyl- β -D-cymaropyranoside (20.11 mg/g), wilfoside C_1N (5.13 mg/g), caudatin (3.96 mg/g), wilfoside K_1N (7.90 mg/g),

wilfoside C_1G (73.25 mg/g), cynauricuoside A (80.16 mg/g), cynauricuoside C (7.20 mg/g) and auriculoside IV (3.07 mg/g). The chemical structures and the total ion chromatograms of the 8 compounds are presented in Fig. 1. The total glycosides content of the TCSGs was 73.5%, as detected according to the vanillin-vitriol colorimetric method (42). The standard curve was constructed using the cynauricuoside A standard.

Reagents. H_2O_2 was purchased from Nanchang Baiyun Pharmaceutical Co., Ltd. (Nanchang, China). Fetal bovine serum (FBS) was purchased from Gibco/Thermo Fisher Scientific, Inc. (Waltham, MA, USA). Alanine aminotransferase (ALT; cat. no. C009-2), aspartate aminotransferase (AST; cat. no. C010-2), lactate dehydrogenase (LDH; cat.

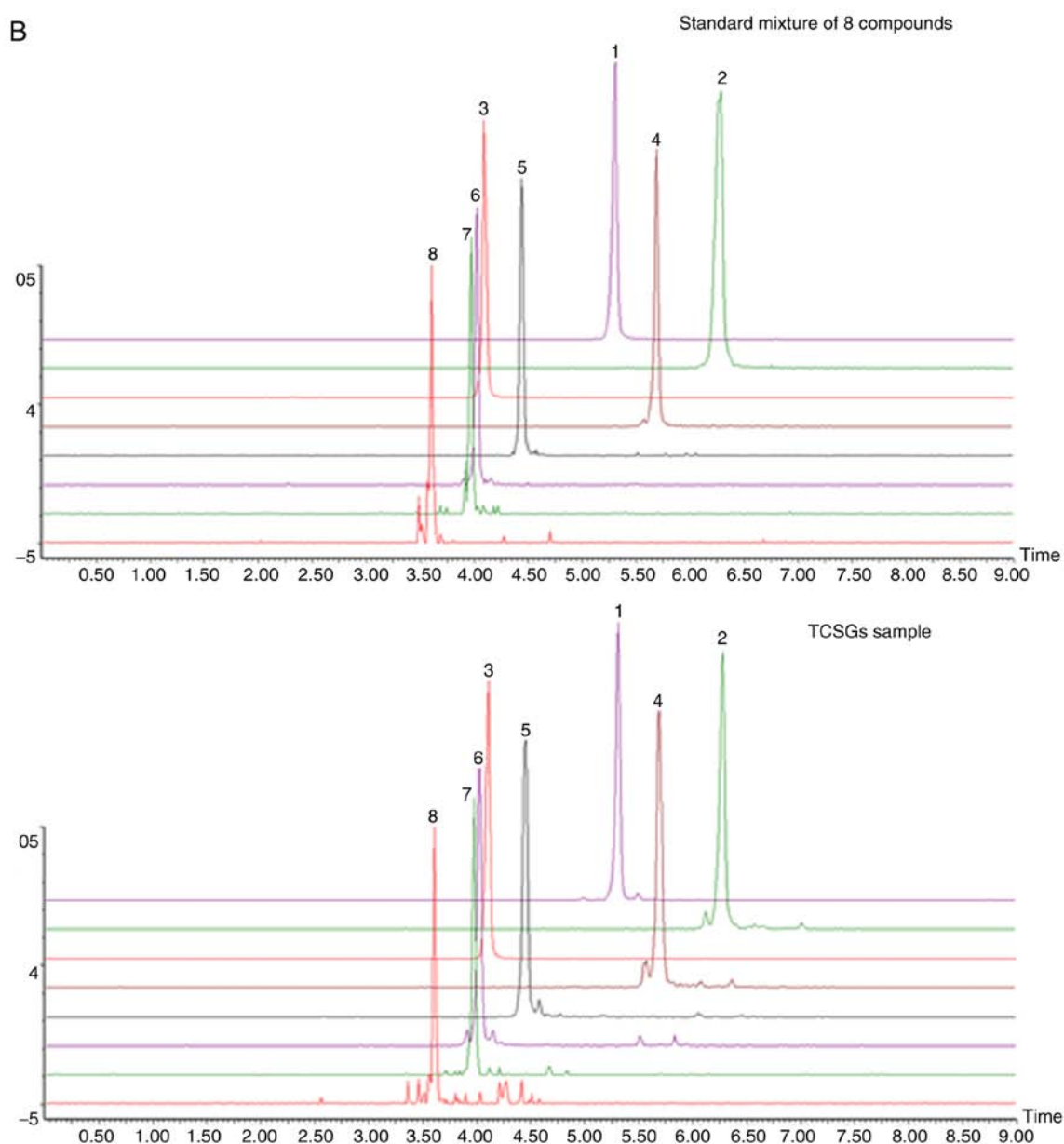


Figure 1. Continued. (B) The total ion chromatograms of 8 standard compounds and TCSGs by multiple reaction monitoring. No. 1, caudatin-2,6-dideoxy-3-O-methy- β -D-cymaropyranoside; no. 2, wilfoside C1N; no. 3, caudatin; no. 4, wilfoside K1N; no. 5, wilfoside C1G; no. 6, cynauricoside A; no. 7, cynauricoside C; no. 8, auriculoside IV. TCSGs, total C-21 steroidal glycosides.

no. C020-2) and NO (cat. no. C013-2) assay kits were obtained from Nanjing Jiancheng Bioengineering Institute (Nanjing, China). SOD (cat. no. S0101), CAT (cat. no. S0051), GSH-Px (cat. no. S0058), MDA (cat. no. S0131) and ROS (cat. no. S0033) assay kits were purchased from Beyotime Institute of Biotechnology (Shanghai, China). Dulbecco's modified Eagle's medium (DMEM), the BCA protein quantification kit and the cytoplasmic-nuclear protein extraction kit were obtained from KeyGEN Biotech Co., Ltd. (Nanjing, China). Anti-iNOS (cat. no. 13120), anti-NF- κ B (cat. no. 8242), anti-phospho-NF- κ B p65 (cat. no. 3033) and anti-GAPDH (cat. no. 5174) antibodies were purchased from Cell Signaling Technology, Inc. (Danvers, MA, USA). Anti-I κ B α (cat. no. abp57568) antibody was obtained from AmyJet Scientific (Wuhan, China). Anti-phospho-I κ B α (cat. no. orb13487) antibody was obtained from Biorbyt (Wuhan, China). Anti-COX-2

(cat. no. ab15191) and anti-Lamin B1 (cat. no. ab133741) antibodies were purchased from Abcam (Cambridge, UK). Anti-heme oxygenase-1 (HO-1; cat. no. Pb0212) antibody was purchased from Wuhan Boster Biological Technology, Ltd. (Wuhan, China). Anti-Nrf2 (cat. no. Abs120634) antibody was obtained from Absin (Shanghai, China). Goat anti-rabbit (cat. no. AP307P) and mouse IgG (cat. no. AP124P) antibodies were purchased from EMD Millipore (Billerica, MA, USA).

Cell culture. L02 cells, a normal human hepatic cell line, were obtained from KeyGEN Biotech Co., Ltd. (cat. no. KG063) and used in the subsequent experiments, as previously described (43). The L02 cells were incubated at 37°C and 5% CO₂ in DMEM supplemented with 10% FBS and 100 U/ml penicillin-streptomycin. Once cell confluence reached ~80%, the cells were passaged.

Cell viability assay. The viability of the L02 cells was determined using the 3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2-H-tetrazolium bromide (MTT) method. Firstly, the effective concentrations and treatment durations for TCSGs and H₂O₂ were determined using the cell viability data. The L02 cells at the exponential growth phase were suspended in DMEM (5x10³ cells/well in 100 μ l) and seeded into Corning disposable 96-well plates. Cell treatments were performed as follows: i) The cells were incubated with the TCSGs (0, 5, 10, 15, 20, 25, 30, 35, 40 and 45 μ g/ml) for 24, 48 and 72 h; ii) The cells were cultured for 72 h and then treated with H₂O₂ (0, 0.2, 0.4, 0.6, 0.8, 1.0, 1.2, 1.5 and 2 mM) for 24 h. The cells were then washed with PBS, and 10 μ l of MTT solution (5 mg/ml) were added to each well and the plate was incubated for 4 h. Then cell supernatants were discarded and 150 μ l dimethyl sulfoxide were added to each well. The formazan crystals in the wells were dissolved and the absorbance was read at 570 nm using the Universal Microplate Reader (Thermo Fisher Scientific, Inc.). During these experiments, the TCSG group (0 μ g/ml) or the H₂O₂ group (0 mM) was considered to be the control group (negative control) in each test, respectively. Based on the results obtained from the above-mentioned experiments, the effect of TCSGs on the viability of the L02 cells damaged by H₂O₂ was investigated as the selected protocol. There were 5 groups during this test: The control group (negative control), the H₂O₂ group (positive control), and the TCSGs I (10 μ g/ml), TCSGs II (20 μ g/ml) and TCSGs III (40 μ g/ml) groups. Following exposure to the TCSGs (0, 10, 20 and 40 μ g/ml) for 48 h, the cells were treated with or without H₂O₂ (1.5 mM) for 24 h. MTT assay was then performed as described above.

Cell morphological changes assessment. Briefly, the L02 cells (2x10⁵ cells/well) were incubated in corning disposable 6-well plates at 37°C and 5% CO₂ for 24 h. The cells were pretreated with the TCSGs (0, 10, 20 and 40 μ g/ml) for 48 h, and then stimulated with or without H₂O₂ (1.5 mM) for 24 h. Cell morphological changes were imaged under a contrast microscope (Zeiss, Axiovert 200; Zeiss GmbH, Jena, Germany).

Assay of AST, ALT and LDH activities and NO levels. The levels of NO and the activities of hepatocellular leakage enzymes were assayed using commercial assay kits according to the manufacturer's instructions, respectively. Exponential growth phase cells were cultured in corning disposable 96-well plates (5x10³ cells/well in 100 μ l) for 24 h and pretreated with the TCSGs (0, 10, 20 and 40 μ g/ml) for 48 h, followed by exposure to, or the absence of, 1.5 mM H₂O₂ for 24 h. The cell culture media were then collected for the detection of AST, ALT and LDH activities, and NO levels. The data were expressed as units per liter (U/l) and μ mol/l, respectively.

Determination of SOD, CAT and GSH-Px activities, and MDA levels. Following treatment with TCSGs and/or H₂O₂ as described above, the cells were washed and lysed with radioimmunoprecipitation assay (RIPA) buffer containing 1 mM phenylmethylsulfonyl fluoride (PMSF) and maintained on ice for 30 min. Cell lysates were collected and centrifuged at 10,000 x g for 5 min. The supernatants were used to detect the activities of SOD, CAT and GSH-Px, and the levels of MDA following the manufacturer's instructions. The protein

contents of the supernatants were assayed using a BCA protein quantification kit.

Detection of ROS levels. To evaluate the direct effects of TCSGs on H₂O₂-induced oxidative stress in L02 cells, the intracellular ROS levels were observed and analyzed via a fluorescence microplate assay and fluorescence microscopy. According to the manufacturer's instructions and as previously described (44), the intracellular ROS levels were measured using the ROS assay kit. Briefly, exponential growth phase L02 cells were pretreated with or without TCSGs for 48 h, followed by exposure to, or the absence of, 1.5 mM H₂O₂ for 24 h. Following 3 washes with PBS, the L02 cells were treated with 10 μ M of the fluorescent probe, 2',7'-dichlorodihydrofluorescein diacetate (DCFH-DA; 1:1,000), and incubated for 20 min at 37°C in the dark. Finally, the cells preloaded with DCFH-DA were imaged with a fluorescence microscope (Olympus Corporation, Tokyo, Japan), and the DCF fluorescence intensity was detected with an EnSpire Multifunctional Microplate Reader (PerkinElmer Inc., Waltham, MA, USA) under a 488 nm excitation wavelength and 525 nm emission wavelength.

RNA extraction and reverse transcription-quantitative polymerase chain reaction (RT-qPCR) detection. Total RNA was extracted from the L02 cells using TRIzol reagent (Invitrogen/Thermo Fisher Scientific, Inc.) and first-stand cDNA was obtained using the Thermo Fisher K1622 first-stand cDNA Synthesis kit (Thermo Fisher Scientific, Inc.). Gene-specific primer sequences were designed and synthesized by GenScript Inc., Nanjing, China, as previously described (45). The primer sequences utilized are presented in Table I.

The qPCR reactions were carried out in 20 μ l reaction mixtures containing 10 μ l 2X Real Time PCR Master Mix (SYBR-Green), 2 μ l primer mix (including forward and reverse primers), and 1 μ l cDNA diluted in RNase-free water. The housekeeping gene, GAPDH, was used as an internal standard. The levels of relative mRNA were analyzed using the 2^{- $\Delta\Delta$ C_q} comparative approach (46).

Western blot analysis. The L02 cells were lysed in RIPA buffer (containing 1 mM PMSF) on ice for 30 min and total proteins were collected. The cytoplasmic and nuclear protein samples were extracted using the cytoplasmic-nuclear protein extraction kits. The protein concentration was detected using the BCA protein quantification kit. Equal amounts of proteins (50 μ g/lane) were separated on 10% SDS-PAGE and electroblotted onto polyvinylidene difluoride membranes (EMD Millipore). Following blocking with 5% skim milk for 1 h, the membranes were incubated overnight with primary antibodies (anti-iNOS, 1:1,000; anti-COX-2, 1:800; anti-NF- κ B p65, 1:1,000; anti-phospho-NF- κ B p65, 1:1,000; anti-I κ B α , 1:1,000; anti-phospho-I κ B α , 1:1,000; anti-Nrf2, 1:500; anti-HO-1, 1:800; anti-Lamin B1, 1:1,000; and anti-GAPDH, 1:1,000) at 4°C. Following 3 washes, the membranes were incubated for 1 h with goat anti-rabbit IgG (1:2,000) and goat anti-mouse IgG (1:2,000) peroxidase-conjugated secondary antibodies at room temperature. The transferred proteins were visualized using an enhanced chemiluminescence detection kit (Beyotime Institute of Biotechnology, Shanghai, China) and the grayscale values

Table I. Primer sequences used for reverse transcription-quantitative polymerase chain reaction.

Gene	Sequence (5'-3')	Accession number
TNF- α	F: GTCAACCTCCTCTCTGCCATCAAG	NM_000594
TNF- α	R: CTGAGTCGGTCACCCCTTCTCCA	
IL-6	F: CTCAATATTAGAGTCTCAACCCCA	NM_000600
IL-6	R: GAAGGCGCTTGTGGAGAAGG	
iNOS	F: GGAGCCAGCTCTGCATTATC	NM_000620
iNOS	R: TTTTGTCTCCAAGGGACCAG	
COX-2	F: TGCACTACATACTTACCCACTTCAA	NM_000963
COX-2	R: CAAATGTGATCTGGATGTCAACACA	
GAPDH	F: GAAGGTCGGAGTCAACGGAT	NM_002046
GAPDH	R: CCTGGAAGATGGTGATGGG	

F, forward; R, reverse; TNF, tumor necrosis factor; IL, interleukin; iNOS, inducible nitric oxide synthase; COX-2, cyclooxygenase-2.

of each band were quantified using Tanon-5200 Analyzer software. Target protein expression levels were normalized based on the internal controls (GAPDH and Lamin B1).

Statistical analysis. All results are presented as the means \pm standard deviation, and analyzed using one-way analysis of variance followed by Tukey's post hoc test for multiple comparisons with SPSS software (version 22.0; IBM Corp., Armonk, NY, USA). All histograms were plotted using GraphPad Prism 5 software (GraphPad Software, Inc., La Jolla, CA, USA). A value of $P < 0.05$ was considered to indicate a statistically significant difference.

Results

Effect of TCSGs on H_2O_2 -induced cytotoxicity in and morphology of L02 cells. The cell viability data indicated that the TCSGs exerted no significant toxicity towards the L02 cells, even at high concentrations (40 μ g/ml) following treatment for 24 and 48 h. However, TCSGs at the concentrations exceeding 40 μ g/ml and treatment for longer than 48 h markedly decreased cell viability ($P < 0.05$; Fig. 2A). As shown in Fig. 2B, with increasing concentrations of H_2O_2 , cell viability decreased in a dose-dependent manner, and the 50% inhibitory concentration of H_2O_2 treatment was 1.49 mM, as determined by SPSS software. Based on these results, treatment with 10, 20 and 40 μ g/ml TCSGs and 1.5 mM H_2O_2 , as well as the time points of 48 h for the TCSGs and 24 h for H_2O_2 , were employed in the subsequent experiments. With this selected protocol, the effects of TCSGs on H_2O_2 -induced cytotoxicity in and morphology of the L02 cells were investigated. The results revealed that pretreatment with the TCSGs significantly increased the cell survival rate in the cells exposed to H_2O_2 (Fig. 2C). As shown in Fig. 2D, the control cells exhibited an irregular polygon morphology, with only a few cells being spindle-shaped. Following exposure to H_2O_2 (1.5 mM), the cells became smaller. However, pretreatment with the TCSGs (10, 20 and 40 μ g/ml) for 48 h, attenuated the morphological changes of the L02 cells observed with exposure to H_2O_2 .

Effect of TCSGs on the activities of ALT, AST, LDH and NO levels in L02 cells. As shown in Fig. 3, exposure to H_2O_2 led

to a significant elevation in the levels of hepatocellular leakage enzymes, including ALT, AST and LDH, which are indicative of hepatic injury. When compared with the control group, the NO levels in the group exposed to H_2O_2 were markedly increased ($P < 0.01$). Following pretreatment with the TCSGs, there was a significant attenuation in the activities of ALT, AST and LDH, and the NO levels when compared with the H_2O_2 group ($P < 0.05$, $P < 0.01$ or $P < 0.001$). The results thus suggested that TCSGs attenuated H_2O_2 -induced injury in L02 cells.

Effect of TCSGs on the activities of SOD, CAT and GSH-Px, and MDA levels in L02 cells. The effects of TCSGs on the activities of SOD, CAT and GSH-Px in the L02 cells are presented in Fig. 4A-C. H_2O_2 significantly ($P < 0.001$) decreased the activities of SOD, CAT and GSH-Px in the L02 cells. However, pretreatment with the TCSGs alleviated the inhibition of antioxidant enzyme activities in a dose-dependent manner under oxidative stress.

The effects of TCSGs on the levels of MDA in the H_2O_2 -exposed L02 cells are presented in Fig. 4D. Exposure to H_2O_2 significantly increased the levels of MDA in the L02 cells when compared with the control group ($P < 0.001$). However, pretreatment with the TCSGs dose-dependently inhibited the increase in the MDA levels when compared with the H_2O_2 group ($P < 0.001$).

Effect of TCSGs on ROS levels in L02 cells. The effects of TCSGs on the production of ROS in the H_2O_2 -exposed L02 cells are presented in Fig. 5B. Exposure to H_2O_2 significantly increased intracellular ROS levels in the L02 cells ($P < 0.001$), and this effect was markedly suppressed by the TCSGs. The results revealed that the TCSGs dose-dependently attenuated H_2O_2 -induced ROS production in the L02 cells. In addition, each group of images was recorded using a digital camera attached to an inverted fluorescence microscope. As shown in Fig. 5A, the results were in agreement with the fluorescence intensity assays.

Effect of TCSGs on H_2O_2 -induced production of inflammatory mediators and cytokines in L02 cells. iNOS and COX-2 are used as inflammatory markers and play critical roles in the

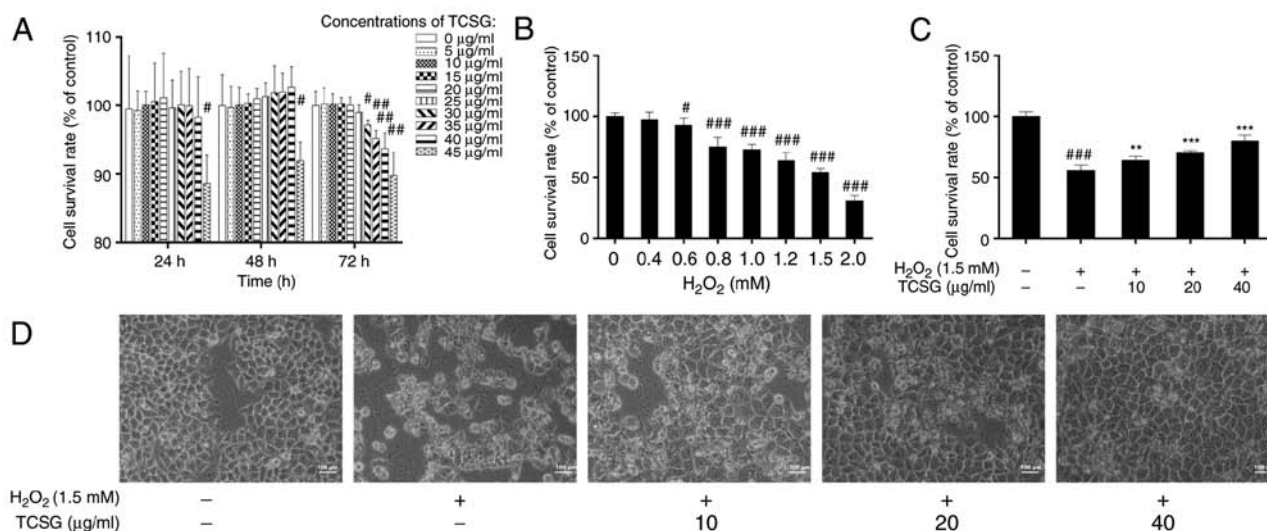


Figure 2. Effect of TCSGs and H_2O_2 on L02 cell morphological changes and cell viability. (A) Cells were treated with the TCSGs (0, 5, 10, 15, 20, 25, 30, 35, 40 and 45 $\mu\text{g/ml}$) for 24, 48 and 72 h. (B) Cells were cultured for 72 h and then exposed to H_2O_2 (0, 0.2, 0.4, 0.6, 0.8, 1.0, 1.2, 1.5 and 2 mM) for 24 h. (C) Pretreatment with TCSGs (10, 20 and 40 $\mu\text{g/ml}$) for 48 h, followed by treatment with H_2O_2 (1.5 mM) for 24 h. (D) L02 cells were cultured with TCSGs (10, 20 and 40 $\mu\text{g/ml}$) for 48 h, followed by the incubation with 1.5 mM H_2O_2 for 24 h. Cell morphology was imaged using an inverted phase contrast microscope (magnification, $\times 200$; scale bar, 100 μm). All data are presented as the means \pm standard deviation ($n=6$). $^{\#}P<0.05$, $^{##}P<0.01$ and $^{###}P<0.001$ vs. the control group (0; no treatment); $^{**}P<0.01$ and $^{***}P<0.001$ vs. H_2O_2 group. TCSGs, total C-21 steroidal glycosides; H_2O_2 , hydrogen peroxide.

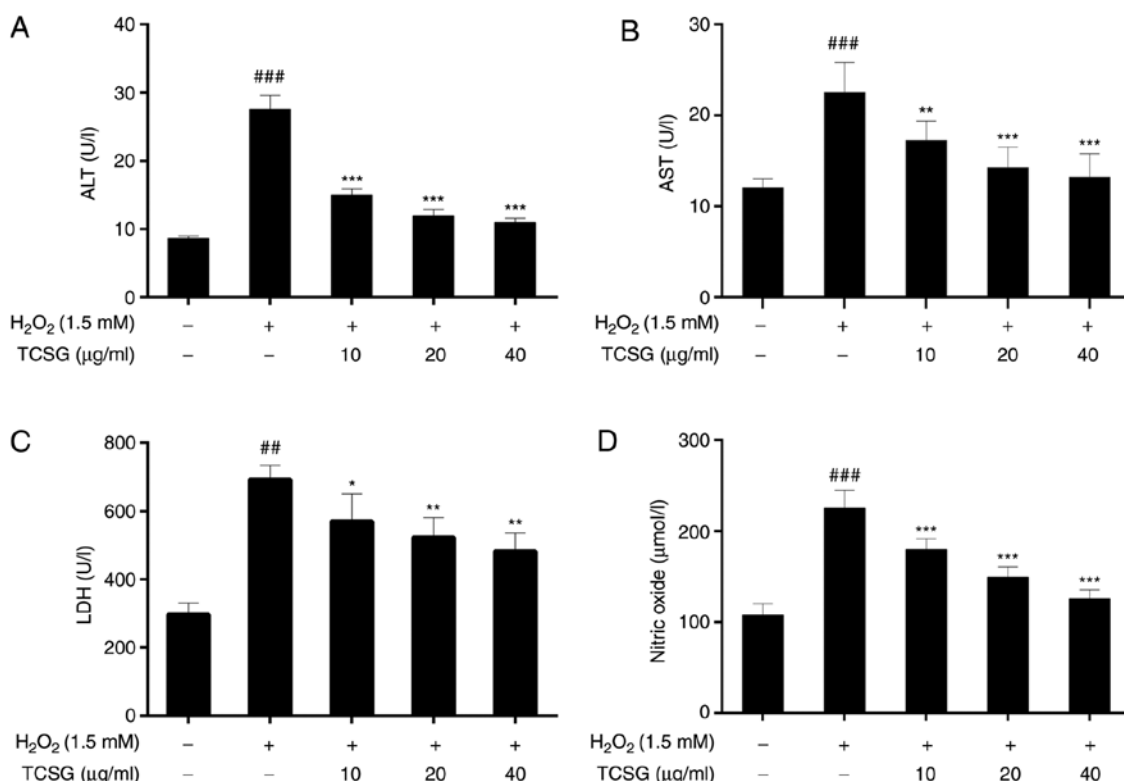


Figure 3. Effect of TCSGs on the supernatant levels of (A) ALT, (B) AST, (C) LDH and (D) NO in H_2O_2 -exposed L02 cells. Data are expressed as the means \pm standard deviation ($n=5$). $^{##}P<0.01$ and $^{###}P<0.001$ vs. the control group (no treatment); $^{\#}P<0.05$, $^{**}P<0.01$ and $^{***}P<0.001$ vs. H_2O_2 group. TCSGs, total C-21 steroidal glycosides; H_2O_2 , hydrogen peroxide; ALT, alanine aminotransferase; AST, aspartate aminotransferase; LDH, lactate dehydrogenase; NO, nitric oxide.

pathogenesis of chronic inflammation. The effects of TCSGs on the expression levels of iNOS and COX-2 in the L02 cells are presented in Fig. 6. The protein and mRNA expression levels of iNOS (Fig. 6A, B and D) and COX-2 (Fig. 6A, C and D) increased significantly in the H_2O_2 -exposed group; however,

pretreatment with the TCSGs dose-dependently inhibited the increase in the iNOS and COX-2 levels. Furthermore, the results of RT-qPCR demonstrated that the TCSGs markedly attenuated the upregulation in the mRNA expression levels of TNF- α and IL-6 (Fig. 6D) induced by H_2O_2 in the L02 cells,

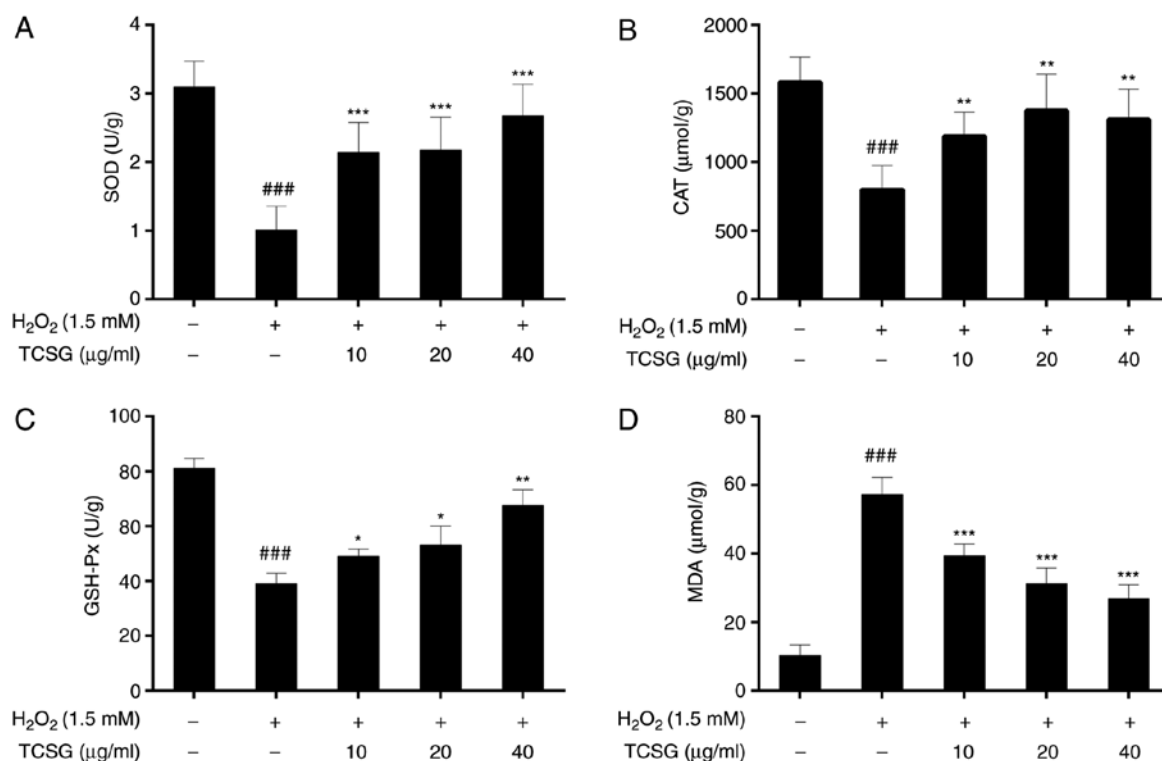


Figure 4. Effect of TCSGs on (A) SOD, (B) CAT, (C) GSH-Px and (D) MDA levels in H₂O₂-induced L02 cells. The results are presented as the means \pm standard deviation (n=5). ###P<0.001 vs. the control group (no treatment); *P<0.05, **P<0.01 and ***P<0.001 vs. H₂O₂ group. TCSGs, total C-21 steroidal glycosides; H₂O₂, hydrogen peroxide; SOD, superoxide dismutase; CAT, catalase; GSH-Px, glutathione peroxidase; MDA, malondialdehyde.

which was consistent with the iNOS and COX-2 expression levels mentioned above.

Effect of TCSGs on the Nrf2 signaling pathway in L02 cells. In order to determine the antioxidative mechanisms of action of the TCSGs, the protein expression levels of Nrf2 and HO-1, as well as the translocation of Nrf2 into the nucleus were measured by western blot analysis (Fig. 7). The results revealed that H₂O₂ significantly decreased Nrf2 expression in the cytosol and increased Nrf2 expression in the nucleus, indicating that nuclear Nrf2 translocation had occurred (Fig. 7A, B and D). H₂O₂ also downregulated the protein expression of HO-1 in the L02 cells (Fig. 7C). Pretreatment with the TCSGs resulted in the enhanced expression of Nrf2 and HO-1, suggesting that treatment with the TCSGs activated the Nrf2 signaling pathway.

Effect of TCSGs on the activation of the NF- κ B signaling pathway in L02 cells. To further elucidate the mechanisms responsible for the suppression of the inflammatory mediators by the TCSGs in the H₂O₂-exposed L02 cells, the NF- κ B signaling pathway was investigated (Fig. 8). It was revealed that pretreatment with the TCSGs dose-dependently downregulated NF- κ B p65 and phospho-NF- κ B p65 levels in the nucleus, and enhanced NF- κ B p65 expression in the cytosol in the H₂O₂-injured L02 cells (Fig. 8A-C and G). As shown in Fig. 8A and E, the TCSGs significantly inhibited the H₂O₂-induced phosphorylation of I κ B α in the L02 cells. These results indicated that the TCSGs prevented the translocation of NF- κ B by inhibiting the phosphorylation of NF- κ B, as well as the degradation and phosphorylation of I κ B α in the H₂O₂-exposed L02 cells.

Taken together, the potential molecular mechanisms of action of TCSGs in human liver cells and their protective effects against H₂O₂-induced oxidative injury and inflammation are presented in Fig. 9.

Discussion

Hepatic injury is a leading cause of hepatic fibrosis and cancer development. Several factors, including oxidative stress, inflammation and immune responses, have been considered to be involved in the pathogenesis of hepatic injury. An accumulating body of evidence has indicated that oxidative stress and inflammation play important roles in the pathogenesis of liver disease (47). High levels of oxidative stress cause cellular damage, such as lipid peroxidation (48). Inflammatory cytokines also induce oxidant formation and cause liver injury. Anti-oxidative stress-based treatments are promising therapeutic strategies for chronic liver diseases.

TCSGs, the main bioactive components of Baishouwu, have been reported to have potent free radical scavenging properties and hepatoprotective effects (13,17). In the present study, it was confirmed that TCSGs was able to ameliorate H₂O₂-induced toxicity in the L02 human hepatic cell line. The results of cell viability assay revealed that treatment with the TCSGs markedly protected the cells from damage caused by H₂O₂, and improved the cell survival rate. As shown in Fig. 2B, treatment with TCSGs at a concentration exceeding 40 μ g/ml or given for a prolonged period of time (72 h) seemed to be toxic to the cells. To the best of our knowledge, C-21 steroidal glycosides at higher concentrations may exert cytotoxic effects *in vitro* (49-52). We speculate the reasons are as follows:

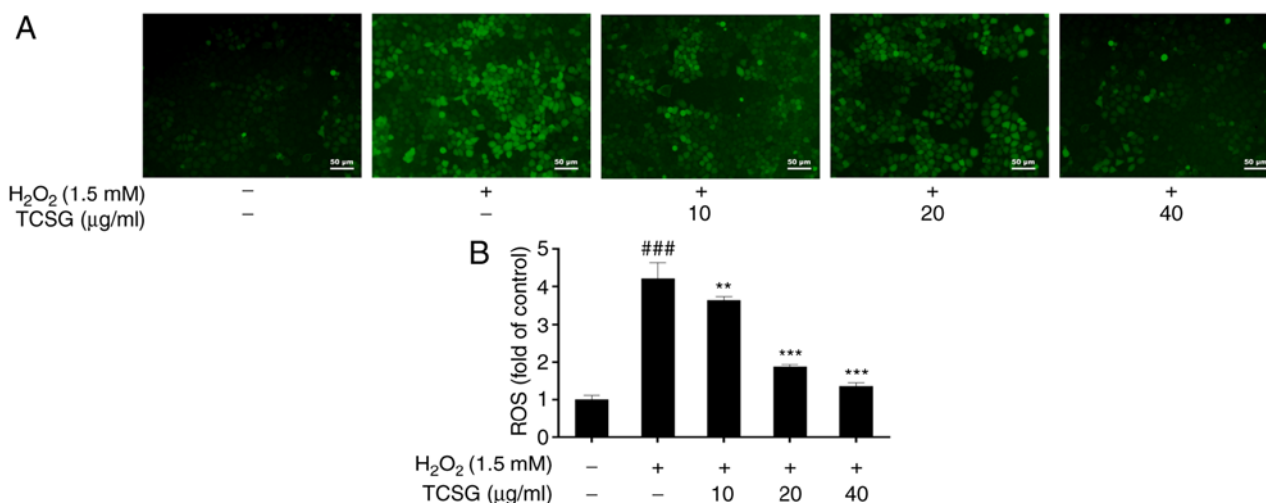


Figure 5. Effect of TCSGs on H₂O₂-induced intracellular ROS production in L02 cells. (A) Microphotographs of intracellular ROS levels in H₂O₂ stimulated L02 cells were obtained by fluorescence microscopy (magnification, x400; scale bar, 50 μm). (B) Intracellular ROS production was measured using a fluorescence microplate reader. The results are presented as the means ± standard deviation (n=4). ^{###}P<0.001 vs. the control group (no treatment); ^{**}P<0.01 and ^{***}P<0.001 vs. H₂O₂ group. TCSGs, total C-21 steroidal glycosides; H₂O₂, hydrogen peroxide; ROS, reactive oxygen species.

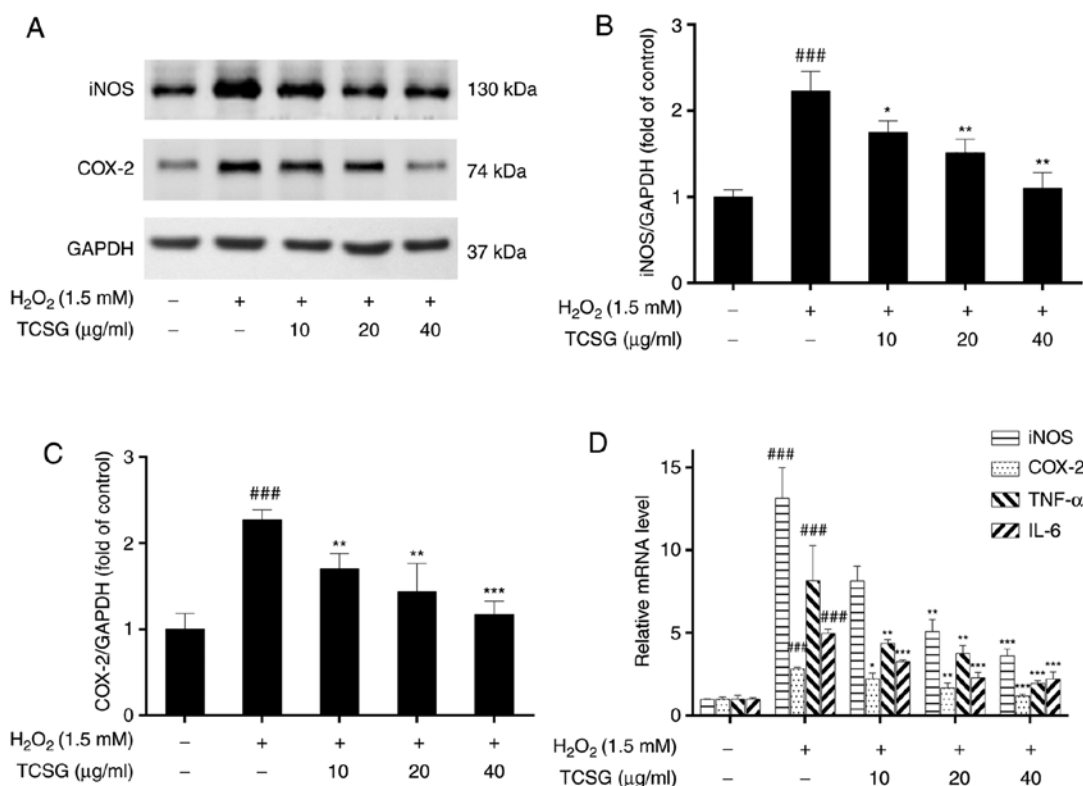


Figure 6. Effect of TCSGs on the H₂O₂-induced expression of inflammatory mediators and cytokines in L02 cells. Equal quantities of proteins (50 μg/lane) were subjected to 10% SDS-PAGE, and the protein expression levels of (A) iNOS, COX-2 and GAPDH were detected by western blot analysis in H₂O₂-exposed L02 cells. The results of (B) iNOS and (C) COX-2 protein expressions were represented. The mRNA expression levels of (D) iNOS and COX-2 were detected by RT-qPCR in the H₂O₂-exposed L02 cells. The mRNA expression levels of (D) TNF-α and IL-6 were detected by RT-qPCR in the H₂O₂-exposed L02 cells. All data are presented as the means ± standard deviation (n=3). ^{###}P<0.001 vs. the control group (no treatment); ^{*}P<0.05, ^{**}P<0.01 and ^{***}P<0.001 vs. H₂O₂ group. TCSGs, total C-21 steroidal glycosides; H₂O₂, hydrogen peroxide; RT-qPCR, reverse transcription-quantitative polymerase chain reaction; iNOS, inducible nitric oxide synthase; COX-2, cyclooxygenase-2; TNF, tumor necrosis factor; IL, interleukin.

i) Chemical structures of C-21 steroidal glycosides usually contain hydrophilic and lipophilic groups, which probably affect the cell membrane and produce a surfactant-like action (53); ii) DNA damage may be involved in the cytotoxic mechanisms of C-21 steroidal glycosides. It has been reported

that C-21 steroidal glycosides exert growth inhibitory effects on human glioma cells by triggering DNA damage (54). To date, the majority of reports on the cytotoxicity of C-21 steroidal glycosides have mainly focused on tumor cells lines, and few on normal cell lines. From this, it can be deduced that

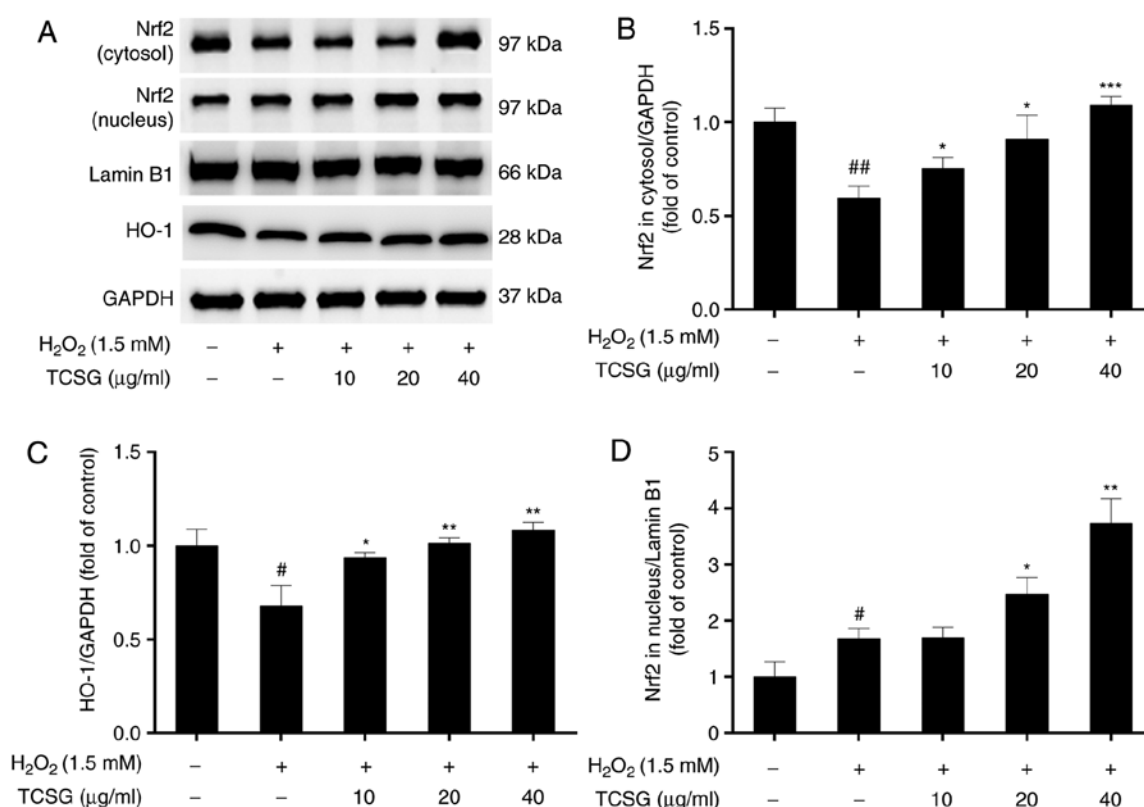


Figure 7. Effect of TCSGs on the protein expression of Nrf2 and HO-1 in L02 cells. Cells were pretreated with the TCSGs (10, 20 and 40 $\mu\text{g/ml}$) for 48 h, stimulated with H_2O_2 (1.5 mM) for 24 h. The nuclear and cytosolic protein samples were prepared to measure the (A, B and D) Nrf2 in the cytosol and nucleus, and (A and C) HO-1 expression. GAPDH and Lamin B1 served as the internal controls. Results are presented as the means \pm standard deviation ($n=3$). $^{\#}P<0.05$ and $^{##}P<0.01$ vs. the control group (no treatment); $^*P<0.05$, $^{**}P<0.01$ and $^{***}P<0.001$ vs. H_2O_2 group. TCSGs, total C-21 steroidal glycosides; H_2O_2 , hydrogen peroxide; Nrf2, nuclear factor erythroid 2-related factor 2; HO-1, heme oxygenase-1.

the cytotoxic effects of C-21 steroidal glycosides on normal cells may differ from those on cancer cell lines. Thus, in the future, we aim to carry out further investigations on the cytotoxic effects and mechanisms of action of C-21 steroidal glycosides.

Oxidative stress leads to cellular damage and cell death via the generation of ROS. Excess ROS primarily induces lipid peroxidation, as indicated by the production of MDA. MDA is a critical marker of lipid peroxidation and is one of the final products of the peroxidation of unsaturated fatty acids. The levels of MDA reflect the extent of cell oxidative damage (55). The antioxidant defense system, containing SOD, CAT, and GSH-Px, scavenges ROS to prevent cell damage in response to oxidative stress (56,57). The present study demonstrated that the enzyme activities of SOD, CAT and GSH-Px, decreased following exposure to H_2O_2 . In addition, the ROS levels and MDA levels increased in the L02 cells exposed to H_2O_2 . However, treatment with the TCSGs attenuated these effects. The results indicated that TCSGs inhibited H_2O_2 -induced oxidative damage by suppressing ROS production and lipid peroxidation, and enhancing the antioxidant defense systems.

The pro-inflammatory enzymes, iNOS and COX-2, play important roles in liver damage. The activation of iNOS induces the production of high levels of NO and further damages hepatocytes (58,59). The present study demonstrated that the TCSGs significantly inhibited COX-2 and iNOS expression, which was consistent with the reduced NO levels in

the L02 cells exposed to H_2O_2 but pretreated with the TCSGs. It has been reported that the upregulation in the mRNA levels of iNOS and COX-2 in hepatic tissues and cells is a result of the increased expression of pro-inflammatory cytokines, such as TNF- α and IL-6, and the activation of NF- κB (60,61). Cytokines, including TNF- α and IL-6 have been the focus of previous investigations evaluating the inflammatory injury of tissues or cells, as they activate the inflammatory and immune responses, exacerbating hepatic injury (62). The results of the present study revealed that the increased levels of TNF- α and IL-6 induced by H_2O_2 were markedly suppressed by pretreatment with the TCSGs. Notably, treatment with the TCSGs also markedly downregulated the expression of NF- κB p65 and p-NF- κB p65 in the nucleus, upregulated NF- κB p65 expression in the cytosol, and reduced the degradation and phosphorylation of I $\kappa\text{B}\alpha$ induced by H_2O_2 in the L-02 cells. Oxidative stress is known to activate NF- κB , leading to the induction of inflammatory genes, which in turn increases neutrophil recruitment to the liver and exacerbates oxidative stress and inflammation. Taken together, the results of the present study indicated that TCSGs exerted hepatoprotective effects against H_2O_2 -induced oxidative stress via the inhibition of the NF- κB signaling pathway and the reduction in COX-2 and iNOS levels, resulting in the reduction of inflammatory factors.

Nrf2 is a pivotal transcription factor, which can encode detoxification enzymes and antioxidant proteins to against oxidative stress (63). Nrf2 regulates the expression of various

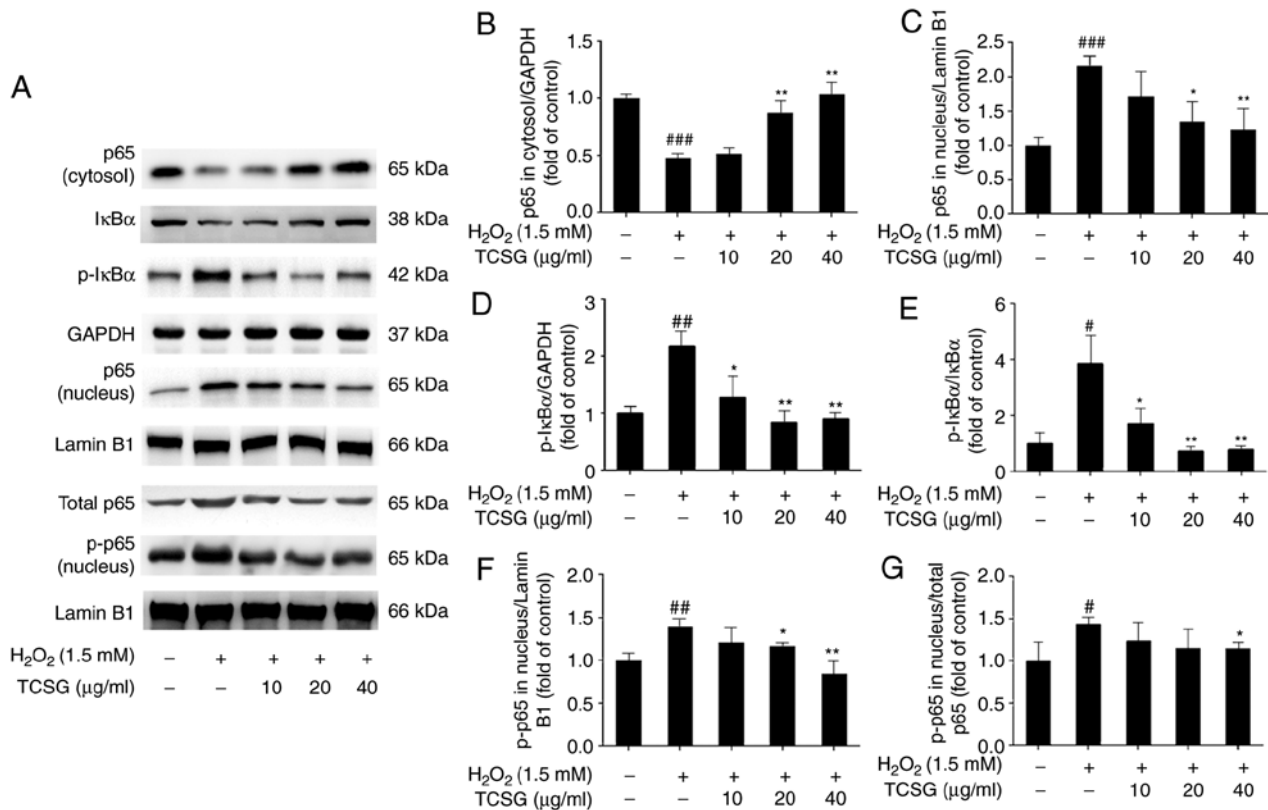


Figure 8. Effect of TCSGs on the activation of the NF- κ B signaling pathway in H_2O_2 -induced L02 cells. Cells were pretreated with the TCSGs (10, 20 and 40 μ g/ml) for 48 h, and stimulated with H_2O_2 (1.5 mM) for 24 h. The samples of nuclear and cytosolic proteins were prepared to detect the expression of (A-C) NF- κ B p65, (A, D and E) I κ B α and p-I κ B α , and (A, G and F) p-NF- κ B p65 using specific antibodies by western blot analysis, respectively. GAPDH and Lamin B1 were used as the internal controls. All results are presented as the means \pm standard deviation (n=3). *P<0.05, **P<0.01 and ***P<0.001 vs. the control group (no treatment); #P<0.05 and ##P<0.01 vs. H_2O_2 group. TCSGs, total C-21 steroidal glycosides; H_2O_2 , hydrogen peroxide; NF- κ B, nuclear factor- κ B; I κ B α , inhibitor of nuclear factor- κ B; p-, phosphorylated.

antioxidant enzymes, including CAT, GSH-Px, and SOD, by using GSH as their substrate (64). The activation of Nrf2 in cells provides an indirect strategy to enhance antioxidative capacity (65,66). Under oxidative stress conditions, Nrf2 is released from Kelch-like ECH-associated protein 1 and translocates to the nucleus; it then recognizes the ARE and regulates the expression of antioxidant enzymes including HO-1. In the present study, the antioxidant ability of TCSGs in the regulation of redox mechanisms by augmenting the expression of antioxidant genes via the Nrf2 signaling pathway, was evaluated. Following exposure to H_2O_2 , Nrf2 expression in the L02 cells decreased in the cytoplasm, but increased in the nucleus when compared with the control group, which indicated that the nuclear Nrf2 translocation and Nrf2 activation had occurred. To the best of our knowledge, the effects of H_2O_2 on Nrf2 activation are strongly dependent on H_2O_2 concentrations and the method of H_2O_2 delivery (67). Different results however, have been obtained under different experimental conditions. Ma *et al* reported that treatment with 200 μ M H_2O_2 for 30 min exerted no significant effect on the cytosolic Nrf2 expression in HepG2 cells (68). In addition, in another study, no significant influence on the expression of nuclear Nrf2 was observed in melanocyte cells exposed to 1 mM H_2O_2 for 24 h (69). Some researchers have proven that the nuclear Nrf2 translocation occurs in melanocytes or human dental pulp stem cells following exposure to H_2O_2 (36,70). Other studies have found that exposure to 300 μ M H_2O_2 for 4 h significantly

decreased the expression of Nrf2 at both the mRNA and protein level in PC12 cells (71). In this study, H_2O_2 decreased Nrf2 expression in the cytoplasm and increased it in the nucleus, which indicated that nuclear Nrf2 translocation and Nrf2 activation occurred. It was speculated that Nrf2 may have been slightly activated as an adaptive defense against oxidative stress under these conditions. In the groups treated with the TCSGs, Nrf2 expression and the translocation of Nrf2 into the nucleus were markedly increased when compared with the H_2O_2 group, indicating that there was an enhanced activation of the Nrf2 signaling pathway. Furthermore, the TCSGs markedly upregulated the expression of downstream target proteins of the Nrf2 signaling pathway, such as HO-1. HO-1 is a key rate-limiting enzyme in the degradation of heme, preventing free heme from participating in the oxidative reaction. The overexpression of HO-1 inhibited ROS generation, NF- κ B activation and nuclear translocation, and protected against oxidative stress and inflammatory injury (72). Combined with the findings of the present study, these results demonstrated that TCSGs protected L02 cells against H_2O_2 -mediated oxidative injury by activating the Nrf2 signaling pathway and inducing HO-1 expression.

In conclusion, in this study, it was demonstrated that pretreatment with the TCSGs protected the L02 cells against H_2O_2 -induced oxidative damage by increasing the expression of Nrf2 and HO-1, mediated by the NF- κ B signaling pathway. The present study also indicated that TCSGs isolated from Baishouwu

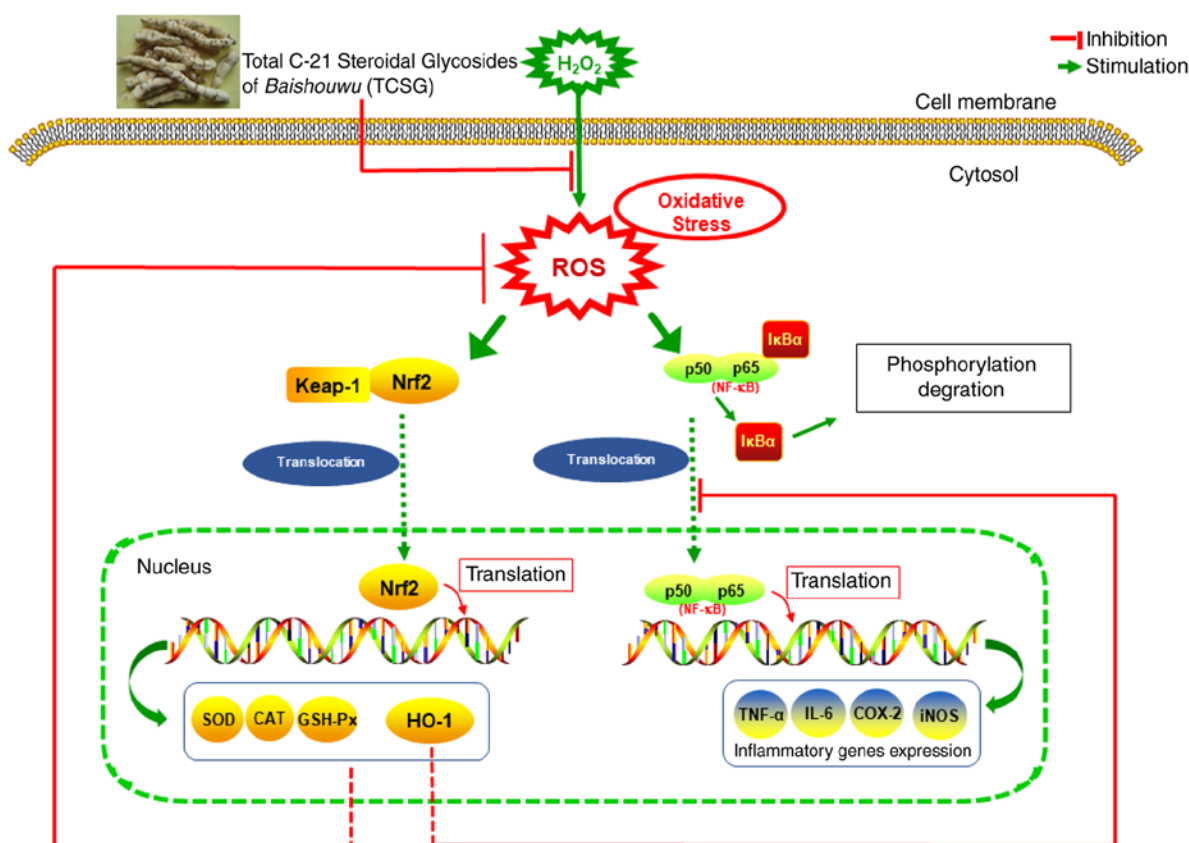


Figure 9. A schematic illustration of the potential molecular mechanism underlying the TCSGs attenuation of oxidative stress and inflammation in L02 cells via the Nrf2 and NF-κB signaling pathways. TCSGs, total C-21 steroidal glycosides; Nrf2, nuclear factor erythroid 2-related factor 2; NF-κB, nuclear factor-κB.

possessed marked anti-oxidative stress and anti-inflammatory effects against H_2O_2 -induced hepatic injury via Nrf2 activation and inhibition of the NF-κB signaling pathway (Fig. 9). Further studies on the potential effects of TCSGs on any other cell signaling pathways involved in the pathogenesis of H_2O_2 -induced hepatotoxicity are still ongoing. Based on these results, TCSGs may be proven to be a promising strategy for preventing inflammation and reducing the risk of hepatic disease.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

PW and YP designed the experiments. ZW, YWa and XM performed the experiments. XW, ZL and SQ equally

contributed in extraction and analysis of chemical constituents. YWe, LS and YD analyzed the results. ZW drafted the manuscript. All the authors read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Patient consent for publication

Not applicable.

Competing interests

The authors confirm that they have no competing interests.

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