

Effect of tanshinone IIA on oxidative stress and apoptosis in a rat model of fatty liver

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Abstract. Oxidative stress is a crucial factor associated with fatty liver disease, which raises the possibility of using antioxidants to improve liver steatosis. Tanshinone IIA (TSIIA) is a traditional Chinese medicine that has been reported to have antioxidant effects *in vitro*. The present study aimed to investigate whether TSIIA possesses antioxidant effects *in vivo* and whether TSIIA was able to improve liver steatosis. Hence, the ability of TSIIA to protect rats from liver disease was explored, particularly in regard to antioxidant activity. Rats were fed a high-lipid diet for 90 days, causing severe liver steatosis, both morphologically and biochemically. An increase in reactive oxygen species (ROS) in the liver was exhibited in addition to significantly elevated serum lipids and malondialdehyde (MDA). Furthermore, hepatocyte apoptosis was measured by Hoechst staining, reverse transcription-quantitative polymerase chain reaction and western blot analysis and an increase in hepatocyte apoptosis rate was indicated in mice on a high-fat diet. Following intraperitoneal injection of TSIIA (10 mg/kg/day), liver steatosis was significantly inhibited. In rats receiving TSIIA treatment, less ROS were indicated in the liver and significantly decreased levels of MDA ($P<0.05$) in serum were exhibited, whereas significantly increased activities of total superoxide dismutase (T-SOD) and glutathione peroxidase (GSH-PX) were observed ($P<0.05$ and $P<0.01$, respectively). In addition, the rate of hepatocyte apoptosis was significantly decreased in the TSIIA group ($P<0.01$). However, TSIIA elicited no effect on serum lipid profiles. These results suggest that TSIIA attenuates oxidative stress by decreasing ROS and MDA production and enhancing the activity of

T-SOD and GSH-PX, which may contribute to the inhibition of apoptosis and amelioration of liver steatosis.

Introduction

Oxidative stress results from an imbalance between *in vivo* oxidative and anti-oxidative effects (1). Oxidative stress is involved in the pathophysiological processes of various diseases, including arteriosclerosis, diabetes, ischemia/reperfusion injury, ethanol intoxication and liver steatosis (2). When the body encounters harmful stimuli, reactive oxygen species (ROS) are produced that may not be effectively removed by antioxidants (3). ROS may be produced by non-alcoholic fatty liver disease (NAFLD) oxidase, xanthine oxidase and the mitochondrial electron transport system (4). Increased levels of ROS have been indicated to enhance the expression of apoptotic genes (5). Furthermore, apoptosis has been indicated as a prominent pathology of liver diseases and is a well-acknowledged type of programmed cell death (6). For protection against oxidative stress-related injury, the body possesses several scavenger systems, including enzymatic and non-enzymatic antioxidants. Enzymatic antioxidants include superoxide dismutase (SOD), catalase (CAT) and the selenium dependent enzyme, glutathione peroxidase (GSH-PX) (7). As antioxidant enzymes may affect lipid peroxidation, an increase in the activities of these enzymes may delay liver steatosis. Furthermore, oxidative stress has been indicated to induce structural and functional cellular effects through damage to DNA, proteins or membrane lipids (8). With respect to hepatocytes, lipid metabolism function may be impaired.

Clinically, liver steatosis without excessive alcohol consumption is known as NAFLD, which is the most common liver condition in the world (9). Abnormal lipid metabolism, which is a high risk for atherosclerosis, is a condition that may result in various types of cardiovascular diseases (10). Previous studies have demonstrated that steatotic livers have increased susceptibility to oxidative stress and that increased ROS may trigger an apoptotic cascade (2,11). Although the scavenging mechanisms of the human body that act against excess ROS-induced cellular injury are typically sufficient, these may not always be enough to combat external stresses. Therefore, there is a pressing requirement to discover natural antioxidants (12). The dried roots of *Salvia miltiorrhiza* Bge, a perennial herb in the mint family, are used as a traditional

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Chinese medicine. In China, the herb is used as an adjunct therapy to treat particular circulatory diseases by promoting blood circulation and removing blood stasis (13). Tanshinone IIA (TSIIA) is one of the major lipophilic components isolated from *Salvia miltiorrhiza* Bge (14). Previous clinical trials and experimental studies have reported the antioxidant and anti-apoptotic properties of TSIIA (15,16). As a natural antioxidant, TSIIA is able to protect cardiac myocytes through anti-oxidative pathways, inhibit oxidation of LDL *in vitro* and reduce the serum levels of oxidized low-density lipoprotein in mice (17-19). In our previous study, we showed that TSIIA was able to ameliorate atherosclerosis by suppressing toll-like receptor 4 (TLR4) and tumor necrosis factor (TNF)- α expression, *in vitro* (20). Large-scale prospective studies have also shown that TSIIA can be beneficial for cardiovascular diseases (14-16). However, there are few studies on the specific mechanism of TSIIA activity on the liver. We have previously indicated that TSIIA is able to protect cells from apoptosis *in vitro* (21). Thus, the purpose of the present study was to evaluate the effects of TSIIA on oxidative stress and hyperlipidemia in rats with fatty livers and to further explore the mechanism of action of TSIIA.

Materials and methods

Pharmacological agents, chemicals and reagents. Sodium TIIA sulfonate was purchased from Shanghai No. 1 Biochemical Pharmaceutical Co., Ltd. (Shanghai, China). Lactate dehydrogenase (LDH), MDA, peroxidase (POD), T-SOD and creatine kinase (CK) test kits were obtained from Nanjing Jiancheng Bioengineering Institute (Nanjing, China). Total cholesterol (TC), thyroglobulin (TG), low density lipoprotein cholesterol (LDL-C), high density lipoprotein cholesterol (HDL-C) test kits were purchased from Sichuan Maker Biotechnology Co., Ltd. (Chengdu, China). TRIzol reagent and the reverse transcription-quantitative polymerase chain reaction (RT-qPCR) kit were purchased from Takara Biotechnology Co., Ltd. (Dalian, China). The BCA Protein Assay kit, SDS-PAGE Gel Preparation kit and all horseradish peroxidase-conjugated secondary antibodies were purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). RIPA lysis buffer was purchased from Beyotime Institute of Biotechnology (Shanghai, China).

Animals. A total of 90 male Sprague-Dawley rats (300 \pm 10 g) aged 6 weeks were purchased from Vital River Laboratories (Beijing, China). Rats were maintained under standard conditions of 12-h light-dark cycle (temperature, 22 \pm 1 $^{\circ}$ C; humidity, 50 \pm 5%) with free access to water. All animal studies were approved by the Local Ethics Committee for Animal Experimentation (approval number 2012-113), according to the guidelines of the Animal Committee of Liaoning University of Traditional Chinese Medicine (Shenyang, China). All efforts were made to minimize the number of animals used and their suffering throughout the study. Rats were randomly divided into a control (CON) group, a high-fat diet group (HFD) group and a TIIA treatment (TAN) group (n=30 per group). Rats in the CON group were fed a regular balanced diet, while those in the HFD group and TAN group were fed a high-fat diet (6.0% sucrose, 1.0% sodium glutamate, 5.0% yolk powder,

8.0% peanut oil, 1.5% cholesterol, 0.4% methylthiouracil, 0.2% sodium cholate and 77.9% regular balanced diet) for 3 months. Furthermore, rats in the TAN group received 1.2 ml sodium TIIA sulfonate (10 mg/kg) by intraperitoneal injection once a day in the last 2 months, while those in the control group and the HFD group received the same quantity of normal saline for the same duration of time. At 3 months, the rats were sacrificed by cervical dislocation under 10% chloral hydrate anesthesia (TCI Development Co, Ltd., Shanghai, China), and small portions of hepatic tissue were excised from the right lobe of the liver.

High-resolution ultrasound imaging. Rats were anaesthetized with isoflurane (Beijing Zhongsheng Ruitai Technology Co., Ltd., Beijing, China) (1.5-3.0%), together with continuous oxygen (1 l/min). Lipid deposition in the liver was observed using a Vevo 2100 High-Resolution Imaging System (FUJIFILM VisualSonics Inc., Toronto, Canada) according to the manufacturer's instructions.

Serum LDH, MDA, POD, T-SOD, CK and GSH-PX content. Rats were anaesthetized with 10% chloral hydrate. Aortic blood was collected and the samples were left to stand for 2 h at room temperature prior to centrifugation at 4 $^{\circ}$ C at 1,000 x g for 20 min. Serum was collected and stored at -80 $^{\circ}$ C. Serum LDH, MDA, POD, T-SOD, CK, and GSH-PX content were estimated using commercial assay kits (A020-1, A003-1, A084-3, A001-1, A032 and A005, all Nanjing Jiancheng Bioengineering Institute).

Lipid detection. A liver sample of 0.1 g was homogenized in 1 ml normal saline, supplemented with 50 ml of a 2:1 chloroform (\geq 99.7%) to methanol (\geq 99.0%) mixture and left to stand for 10 min. Subsequently, the sample was centrifuged at 4 $^{\circ}$ C at 3,000 x g for 10 min and the upper layer was removed. The remaining sample was dried using a Termovap sample concentrator, resuspended in 0.5 ml methanol (\geq 99.0%) and examined using an automatic biochemical analyzer (Toshiba Medical Systems Corp., Otawara-Shi, Japan). TC and TG levels were measured by the COD-CE-PAP and GPO-PAP methods (22), respectively. Serum TG, TC, HDL-C, LDL-C levels were determined by corresponding assay kits (0615021, 0815031, 0915051 and 0915061, all Sichuan Maker Biotechnology Co., Ltd., Sichuan, China).

Hematoxylin and eosin staining of liver tissues. Liver samples were fixed in 4% paraformaldehyde solution at room temperature for 24 h and stained with hematoxylin and eosin using standard techniques. The samples were embedded in paraffin, and 5- μ m thick paraffin sections were cut. The paraffin was removed with xylene, rehydrated through an alcohol gradient, and stained with hematoxylin and eosin and observed using light microscopy at magnification, x200. Images were captured using light microscope linked to a digital CCD camera (Olympus Corp., Tokyo, Japan).

Oil Red O staining of liver tissue. Oil Red O staining was performed following a standard protocol. Frozen liver sections (6- μ m thick) were cut and dried. Samples were washed with 50% ethanol, stained with Oil Red O for 8 min, differentiated

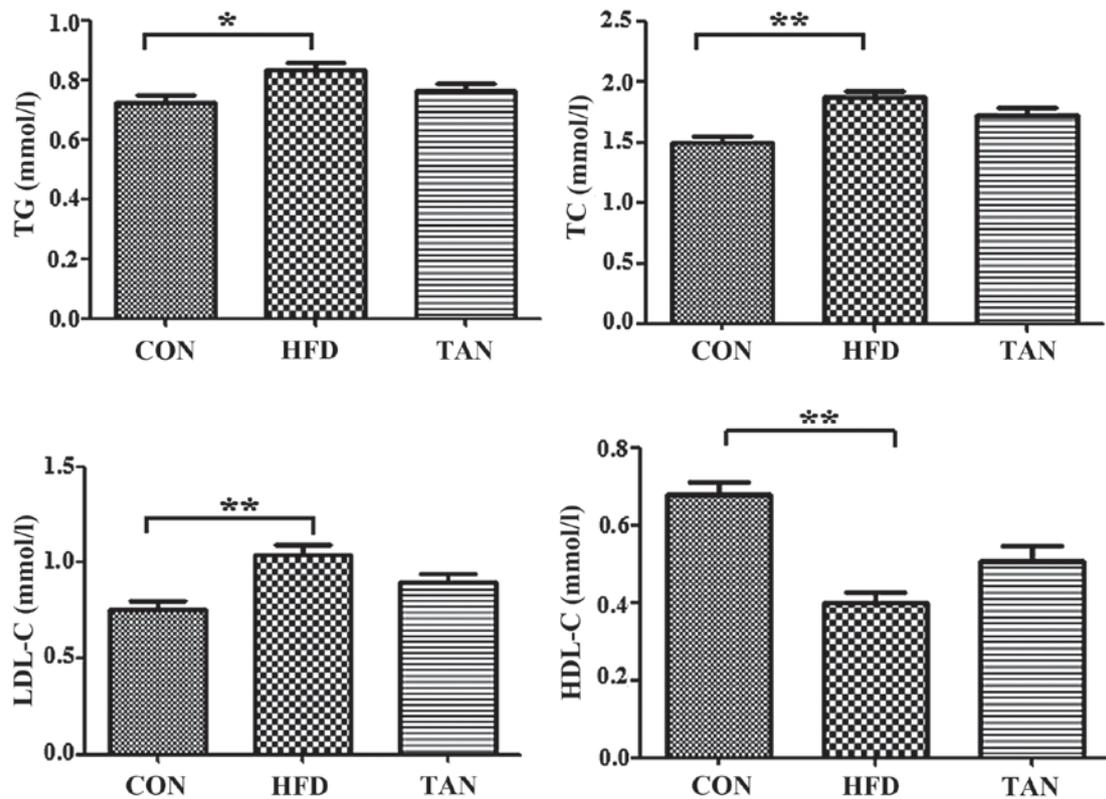


Figure 1. Serum lipid profile. TG, TC, HDL-C and LDL-C levels were determined using corresponding test kits in the CON, HFD and TAN groups. Data are presented as the mean + standard deviation (n=6). *P<0.05; **P<0.01. CON, control group; HFD, high-fat diet group; TAN, tanshinone II A treatment group; TG, thyroglobulin; TC, total cholesterol; HDL-C, high density lipoprotein cholesterol; LDL-C, low density lipoprotein cholesterol.

with 50% ethanol, rinsed with tap water, stained with hematoxylin, rinsed with tap water and mounted with glycerin jelly. Sections were subsequently examined and images were captured using a light microscope linked to a digital CCD camera at magnification, x200.

Hoechst staining. Liver samples that had been fixed with 4% paraformaldehyde were cut into 5- μ m thick sections, deparaffinized and rehydrated. The tissue was permeabilized with Triton X-100 for 15 min. Nuclei were counterstained with Hoechst 33258, according to the manufacturer's instructions using an Apoptotic cell Hoechst 33258 Detection kit (Nanjing KeyGen Biotech Co., Ltd., Nanjing, China). Slides were examined for Hoechst using a fluorescence microscope at 350 nm and magnification, x200.

RNA isolation and RT-qPCR. Total RNA was isolated from liver tissue using TRIzol reagent and reverse transcribed into cDNA using the miScript II RT kit (Qiagen GmbH, Hilden, Germany). Primers were chosen to detect the amplification of the chosen gene and were designed as follows: B cell lymphoma 2 associated protein X (Bax), forward 5'-GCAAACCTGGTGCTCAAGG-3' and reverse 5'-CGTCCCGAAGTAGGAAGG-3'; B cell lymphoma-2 (Bcl-2), forward 5'-CGGGAGAACAGGGTATGA-3' and reverse 5'-CAGGCTGGAAGAGAAGAT-3'; caspase 3, forward 5'-CTGGACTGCGGTATTGAG-3' and reverse 5'-GGGTGCGGTAGAGTAAGC-3'; and GAPDH, forward 5'-TGTGTCCGTCGTGGATCTGA-3' and reverse 5'-TTGCTGTTGAAGTCGCAGGAG-3'. The

reaction mix consisted of 2 μ l cDNA, 0.5 μ l forward and reverse primer mix (10 μ M of each), 12.5 μ l SYBR Premix Ex Taq II (Takara Biotechnology Co., Ltd.) in a final volume of 25 μ l. The following thermal conditions were used for all PCR reactions: 2 min at 95°C, followed by 40 cycles of 30 sec at 95°C and 30 sec at 60°C. RT-qPCR was performed in triplicate using SYBR Green Master Mix (Takara Biotechnology Co., Ltd.) on an ABI 7500 PCR Sequence Detector (Applied Biosystems; Thermo Fisher Scientific Inc., Waltham, MA, USA). Data were analyzed using the SDS software (version 2.0.6; Applied Biosystems; Thermo Fisher Scientific, Inc.). The fold-change in gene expression was determined by the $2^{-\Delta\Delta Cq}$ (23) method with GAPDH (housekeeping gene) as an internal control. The size of the amplified products were indicated to be 149, 149 and 102 bp for Bax, Bcl-2 and caspase 3, respectively.

Determination of hepatic ROS. Levels of ROS in liver slices were determined by the oxidation of dihydroethidium (DHE). Frozen sections of liver (5- μ m thick) were incubated for 30 min at 37°C in 1 ml of 4-(2-hydroxyethyl)-1-piperazineethanesulphonic acid buffer containing 50 μ M of DHE. Frozen sections were subsequently washed three times in phosphate-buffered saline. DHE was oxidized by ROS to produce fluorescent ethidium, which is able to bind to nucleic acids, thus further staining the nucleus a bright fluorescent red. The formation of ethidium was monitored by fluorescence inversion microscopy (magnification, x200) with excitation and emission wavelengths of 480 and 590 nm, respectively.

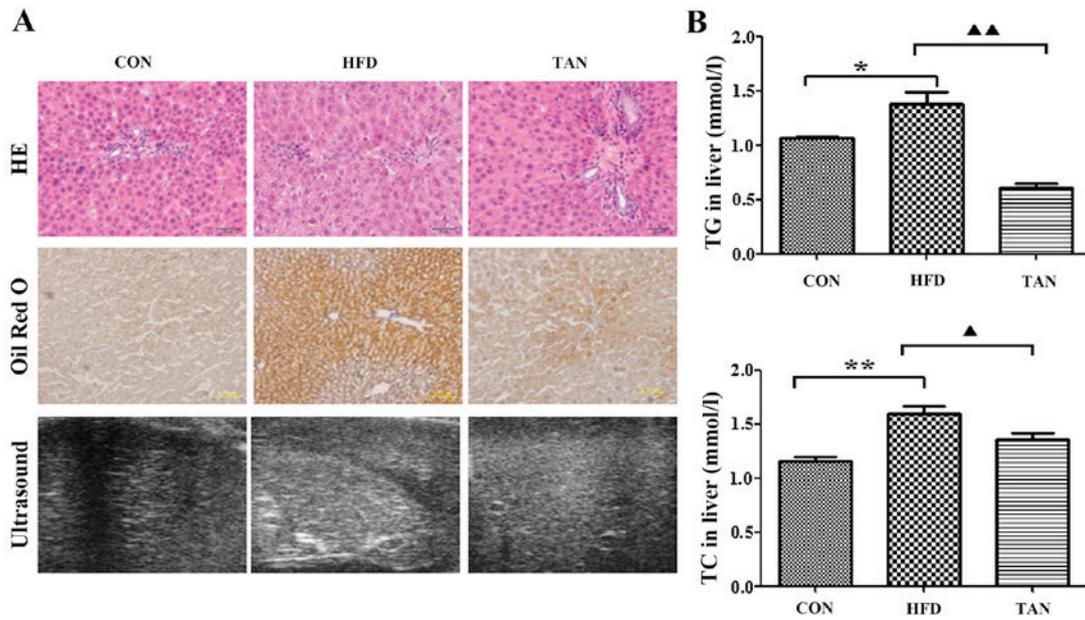


Figure 2. Differences in morphology and lipid deposition in rats from the CON, HFD and TAN groups. (A) Morphological observation of rat hepatocytes in CON, HFD and TAN groups by HE staining, Oil Red O staining and high-resolution ultrasound. (B) TG and TC were measured by the COD-CE-PAP and GPO-PAP methods, respectively, in the livers of CON, HFD and TAN rats. Data are presented as the mean + standard deviation (n=3). *P<0.05; **P<0.01; ▲P<0.05; ▲▲P<0.01. CON, control group; HFD, high-fat diet group; TAN, tanshinone II A treatment group; TG, thyroglobulin; TC, total cholesterol; HE, hematoxylin and eosin.

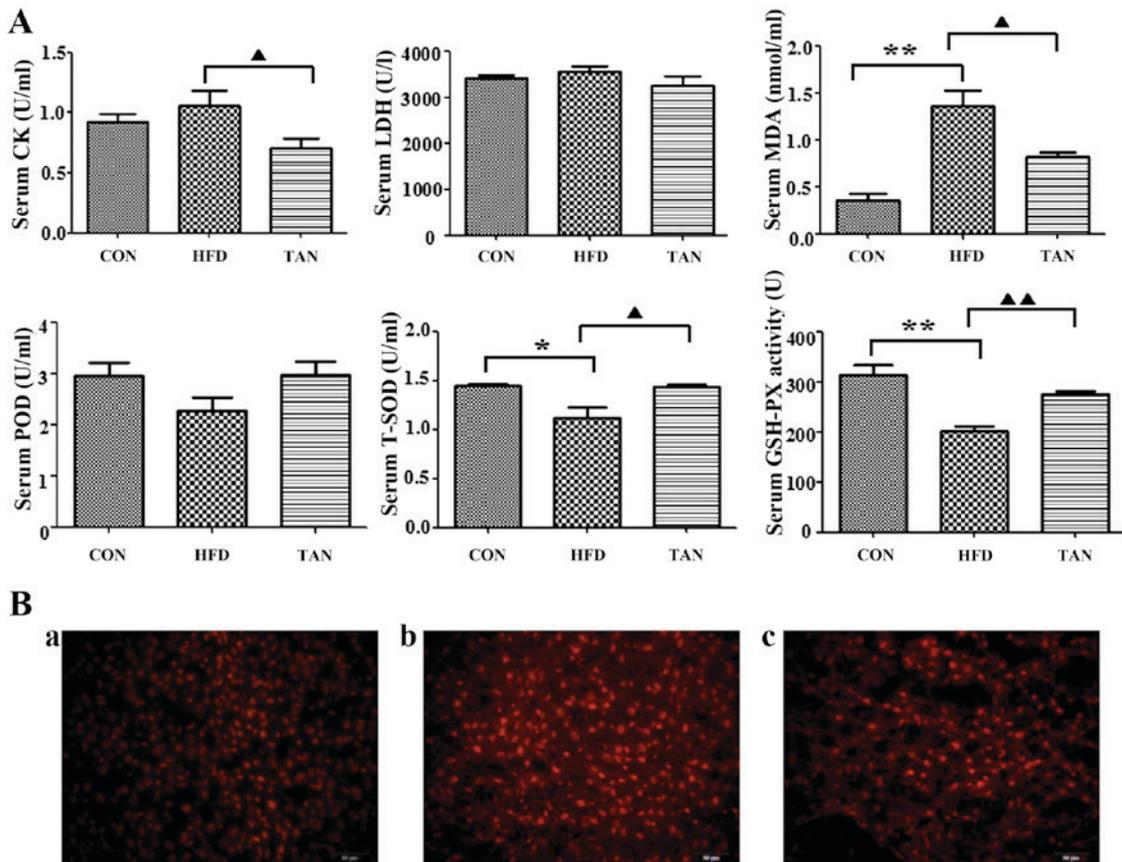


Figure 3. Effects of tanshinone IIA on high-fat diet-induced oxidative stress in the liver of rats. (A) Quantification of CK, LDH, MDA, POD, T-SOD and GSH-PX in the serum of rats from the CON, HFD and TAN groups. Rats were treated and the contents were subsequently assayed. Data are presented as the mean + standard deviation (n=3). *P<0.05; **P<0.01; ▲P<0.05; ▲▲P<0.01. (B) Detection of ROS in the liver of rats from the (a) CON group; (b) HFD group; and (c) TAN group. Rats were treated and dihydroethidium fluorescence was detected. Red color fluorescence represents ROS. CK, creatine kinase; LDH, lactate dehydrogenase; MDA, malondialdehyde; POD, peroxidase; T-SOD, total superoxide dismutase; GSH-PX, glutathione peroxidase; CON, control group; HFD, high-fat diet group; TAN, tanshinone II A treatment group; ROS, reactive oxygen species.

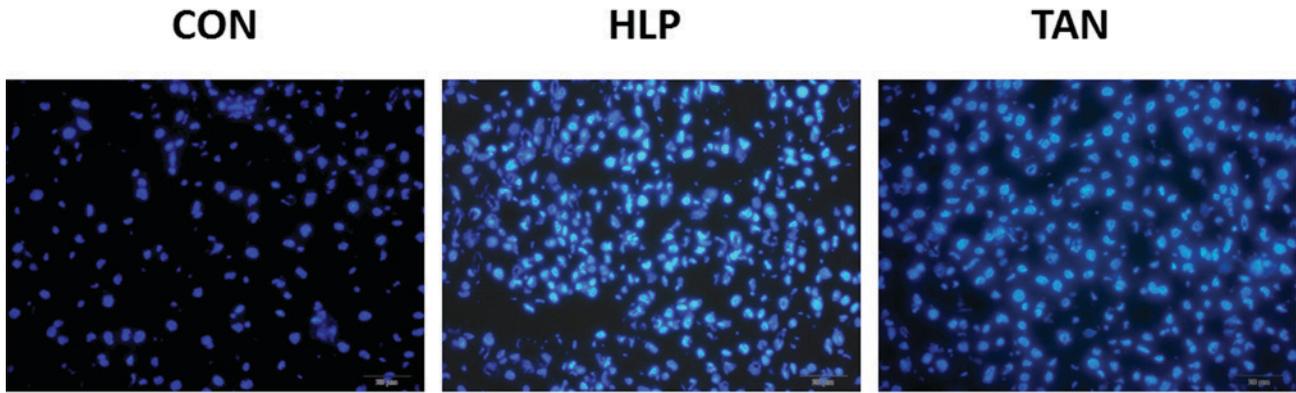


Figure 4. Hoechst assay in the liver of rats in the CON group (n=3), HFD group (n=3) and TAN group (n=3). Blue colour indicates apoptotic cells. CON, control group; HFD, high-fat diet group; TAN, tanshinone II A treatment group.

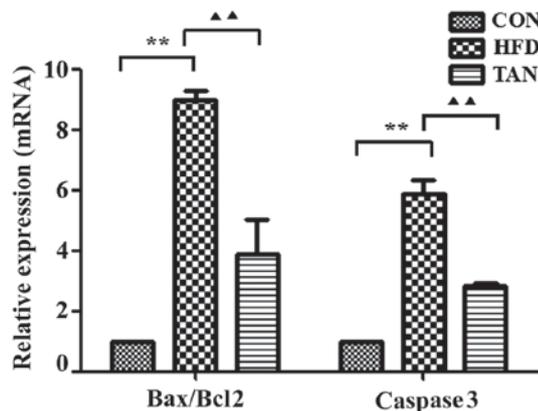


Figure 5. Apoptosis-related gene expression in the liver of rats. Expression levels of apoptosis-related genes, Bax, Bcl-2 and caspase 3, in rat livers were analyzed by reverse transcription-quantitative polymerase chain reaction in CON, HFD and TAN groups. Data are presented as mean + standard deviation (n=3). **P<0.01; ▲▲P<0.001. CON, control group; HFD, high-fat diet group; TAN, tanshinone II A treatment group; Bcl-2, B cell lymphoma-2; Bax, Bcl-2 associated protein X.

Protein extraction and western blotting. Total cellular protein was extracted from liver tissues using RIPA lysis buffer (Beyotime Institute of Biotechnology). Protein concentration was measured using a BCA protein assay kit (Beijing Dingguo Changsheng Biotechnology Co., Ltd., Beijing, China). To assess the protein expression levels of Bax, Bcl-2, and caspase 3 proteins, 80 μ g of protein samples were separated by 10% SDS-PAGE and transferred onto polyvinylidene difluoride membranes (Merck Millipore, Darmstadt, Germany). The membranes were incubated at 4°C overnight with primary antibodies against caspase 3 (1:800, 9662, Cell Signaling Technology, Inc., Danvers, MA, USA), Bax, Bcl-2 and GAPDH (1:500; sc-70407, sc-23960 and sc-32233; all Santa Cruz Biotechnology, Inc., Dallas, TX, USA) and subsequently with a secondary horseradish peroxidase-conjugated goat anti-rabbit and anti-mouse antibody (1:4,000, both Santa Cruz Biotechnology, Inc.) at room temperature for 1 h. Protein bands were visualized using an enhanced chemiluminescence detection kit (Thermo Fisher Scientific, Inc.) and exposed to X-ray film. Protein expression was normalized to the levels of GAPDH.

Statistical analysis. Data are presented as the mean + standard deviation. Differences among the groups were evaluated with the analysis of variance test for multiple groups, using GraphPad Prism 5.0 software (GraphPad Software, Inc., La Jolla, CA, USA). P<0.05 was considered to indicate a statistically significant difference.

Results

Serum lipid levels and lipid deposition in the liver. Compared with the CON group, the rats in the HFD group exhibited significantly increased levels of serum TG (P<0.05), TC (P<0.01) and LDL-C (P<0.01), and significantly decreased levels of serum HDL-C (P<0.01; Fig. 1). However, the TAN group did not indicate any significant difference in lipid profiles when compared with the HFD group (Fig. 1). Furthermore, hematoxylin and eosin-stained liver sections indicated severe hepatic steatosis in the HFD group (Fig. 2A). Compared with the CON group, the hepatocytes from the HFD group were swollen and round with a smaller cytoplasm and contained large lipid droplets. Furthermore, lipid vacuoles of different sizes and inflammatory cells were observed in hematoxylin and eosin-stained sections from the HFD group. In several hepatocytes, the nucleus appeared to be pushed to the side towards the cell membrane. The TAN group exhibited a reduction in hepatic steatosis, cell swelling and hepatocyte size in a number of lipid vacuoles when compared with HFD group (Fig. 2A). Oil Red O staining revealed that the hepatocytes in the HFD group contained a markedly increased number of lipid droplets when compared with the CON group. However, the number of lipid droplets was markedly decreased in the hepatocytes of the TAN group when compared with the HFD group. Compared with the homogeneous liver parenchyma of the HFD group, ultrasound examination of the liver of HFD rats indicated increased echogenicity of the liver parenchyma, characteristic of liver steatosis and heterogeneous parenchyma (24). The echo level of the liver parenchyma was revealed to be decreased in the TAN group when compared with the HFD group (Fig. 2A). Additionally, when compared with the CON group, the livers of rats in the HFD group exhibited significantly increased levels of TG (P<0.05) and TC (P<0.01); however, significantly decreased levels of TG

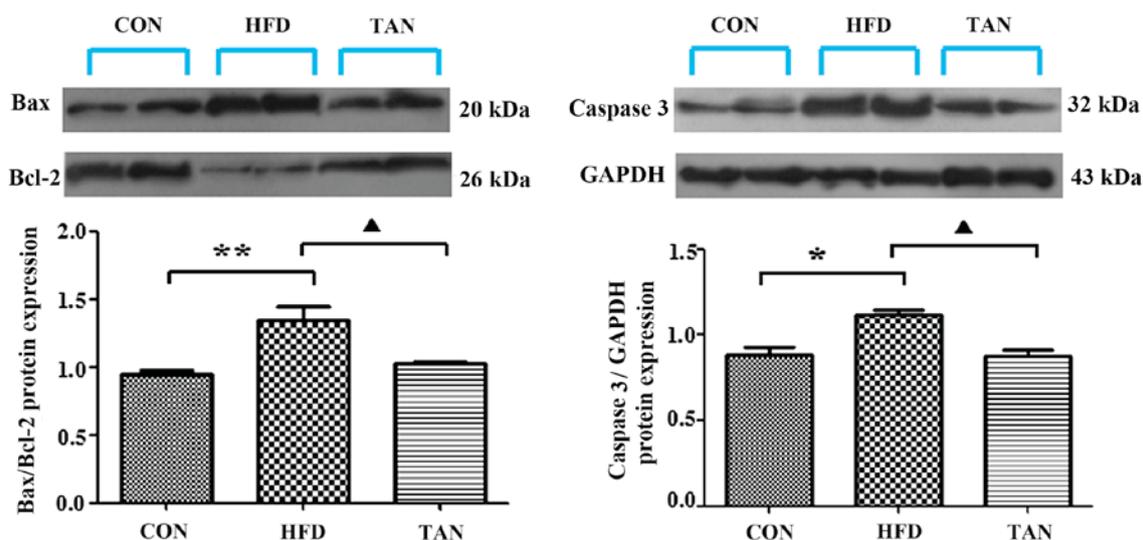


Figure 6. Apoptosis-related protein expression in the liver of rats. Expression levels of apoptosis-related proteins, Bax, Bcl-2 and caspase 3, in rat livers were analyzed by western blotting in the CON, HFD and TAN groups. Data are presented as the mean + standard deviation (n=3). *P<0.05; **P<0.01; ▲P<0.05. CON, control group; HFD, high-fat diet group; TAN, tanshinone II A treatment group; Bcl-2, B cell lymphoma-2; Bax, Bcl-2 associated protein X.

(P<0.01) and TC (P<0.05) were indicated in the TAN group when compared with the HFD group (Fig. 2B).

Effect of TSIIA on antioxidant enzymes and the activity of CK and LDH. Compared with the CON group, MDA content in the HFD group was significantly elevated (P<0.01; Fig. 3A), whereas the TAN group exhibited significantly decreased MDA content when compared with the HFD group (P<0.05). A significant decrease in T-SOD (P<0.05) and GSH-PX (P<0.01) was observed in the HFD group when compared with the CON group; however, significantly increased levels of T-SOD (P<0.05) and GSH-PX (P<0.01) were exhibited in the TAN group compared with the HFD group, indicating treatment with TSIIA reversed this effect. The level of POD activity was not significantly altered in all three groups. A comparison of the serum-marker enzymes LDH and CK, revealed a significant increase in CK activity (P<0.01) in the HFD group when compared with the TAN group; however, LDH activity was unchanged across all three groups.

Inhibitory effect of TSIIA on HFD-induced oxidative stress in the liver. Effects of TSIIA on HFD-induced oxidative stress in the liver of rats were indicated by fluorescence microscopy. Compared with the CON group, the HFD group exhibited markedly increased dihydroethidium fluorescence, which indicated increased ROS content in the liver. However, the level of fluorescence from livers of TAN group rats was indicated to be decreased when compared with the HFD group (Fig. 3B).

Effect of TSIIA on liver cell apoptosis. To determine whether HFD induces apoptosis in liver cells and to explore whether apoptosis may be ameliorated by TSIIA treatment, rates of liver cell apoptosis were quantified via Hoechst assay analysis. The Hoechst assay revealed that the HFD group displayed a markedly increased rate of apoptosis when compared with the CON group. Furthermore, the TAN group exhibited a markedly decreased rate of apoptosis when compared with the HFD

group, which indicated that TSIIA treatment was able to markedly reduce the rate of apoptosis, almost to the level observed in the CON group (Fig. 4).

Apoptosis-related gene and protein expression levels in the liver of rats. The ratio between the pro-apoptotic protein, Bax, and the anti-apoptotic protein, Bcl-2, in addition to the expression levels of caspase 3, are important apoptotic indicators (25,26). The results of RT-qPCR and western blotting showed that Bax/Bcl-2 mRNA and protein expression levels in the liver were significantly increased in the HFD group when compared with the CON group; however, these mRNA and protein expression levels were significantly decreased in the TAN group when compared with the HFD group (P<0.01; Figs. 5 and 6). Furthermore, caspase 3 mRNA and protein expression levels were significantly increased in the HFD group when compared with the CON group and these expression levels were significantly decreased in the TAN group when compared with the HFD group (P<0.05; Figs. 5 and 6).

Discussion

TSIIA is a major component of *Salvia miltiorrhiza* Bge (14). Previous studies have indicated that TSIIA possesses a range of biological properties (15,16,27), predominantly based on its anti-oxidative effects (18). In the present study, the effect of TSIIA on liver steatosis was assessed *in vivo*. TSIIA was identified to reduce lipid deposition in the liver and ameliorate liver steatosis; however, this effect was not based on alterations of serum lipid levels, despite 3 months of treatment. This suggests that actions other than the lipid-lowering effect of TSIIA are responsible for the reduction in liver steatosis exhibited in the present study.

It has been previously reported that oxidative stress is associated with the pathogenesis of NAFLD (28,29). Similarly, the present results indicated that the ROS levels, which are typically generated in a HFD (30), increased in rats in the HFD

group. Correlating with increased ROS, the present findings demonstrated increased levels of lipid peroxide products, in the form of MDA, and decreased T-SOD and GSH-PX were exhibited in rats of the HFD group. As a result, the present findings indicated that the hepatocytes of fatty liver rats were damaged, through increased liver steatosis and apoptosis.

Apoptosis is an active process of cell suicide. Under physiological conditions, it can clear senescent and injured cells to maintain the normal liver volume. However, dysregulation of apoptosis may result in liver diseases (31,32). Feldstein *et al* (33,34) considered that hepatocyte apoptosis was the distinguishing feature of NAFLD and increased with the development of fatty liver. Furthermore, Murakawa *et al* (35) and other previous studies (36,37) demonstrated that antioxidants had anti-apoptotic effects and implied that TSIIA may inhibit apoptosis due to its antioxidant activity. The present study identified that the rate of apoptosis was significantly increased in HFD group rats when compared to the CON group, by Hoechst staining. Moreover, the present findings further supported these effects at the molecular level by investigating Bax and Bcl-2 mRNA and protein expression. Bax and Bcl-2 are apoptosis-related genes; Bax is a pro-apoptotic factor, whereas Bcl-2 has anti-apoptotic activity (38). Xie *et al* (39) and McClintock *et al* (40) suggested that the survivability of cells post-apoptosis stimulation is dependent on the ratio of Bax/Bcl-2. Caspase 3, a cysteine protease, is able to promote apoptosis as a key protease (41). The present findings indicated an increase of the Bax/Bcl-2 ratio and caspase 3 mRNA and protein expression levels in the HFD group. However, these negative effects of a HFD on rat livers were reversed with TSIIA treatment, as indicated by the results from the TAN group. The present results indicate that enhanced anti-oxidative and anti-apoptotic capability may partially explain the protective effect of TSIIA on NAFLD.

Although TSIIA has no effect on serum lipid profiles, we found that attenuation of oxidative stress by TSIIA contributes to the amelioration of hepatic steatosis. In addition, the enhanced antioxidant capacity may account for the inhibition of apoptosis.

In summary, the present study showed that TSIIA improved the antioxidant capacity of the liver, leading to decreased apoptosis and improved hepatic steatosis in a rat model of fatty liver. In future studies, we plan to focus on the anti-apoptotic mechanisms of antioxidants, which may provide a potential target in the prevention and treatment of NAFLD.

Acknowledgements

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