

Novel insights into the effect of Xiaoyao san on corticosterone-induced hepatic steatosis: inhibition of glucocorticoid receptor/perilipin-2 signaling pathway

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Abstract

Objective: Xiaoyao san (XYS) is a classic traditional Chinese medicinal formula. It has been clinically administered to regulate liver function. However, its mechanisms in glucocorticoid-induced hepatic steatosis are unknown. This study aimed to investigate whether YYS protects against corticosterone (CORT)-induced hepatic steatosis, and to explore its mechanism.

Methods: High-fat diet mice induced with hepatic steatosis by 2 mg/kg CORT were administered 2.56 g/kg or 5.12 g/kg YYS daily for 7 weeks. The effects of YYS on hepatic steatosis in mice were evaluated by H&E and Oil Red O staining and by measuring their plasma lipids (triglyceride, total cholesterol, and free fatty acids). The mechanism of YYS against hepatic steatosis was investigated by network pharmacology, immunohistochemistry, western blotting, and gain-of-function/loss-of-function experiments.

Results: YYS alleviated CORT-induced steatosis, decreased plasma lipids, and inhibited glucocorticoid receptor (GR) activation in the liver. Network pharmacology data indicated that YYS may have mitigated hepatic steatosis *via* GR which mediated adipose differentiation-related protein (ADFP). Gain-of-function/loss-of-function experiments *in vitro* confirmed that GR positively regulated ADFP expression.

Conclusions: YYS ameliorated CORT-induced hepatic steatosis by downregulating the GR/ADFP axis and inhibiting lipid metabolism. Our studies implicate that YYS is promising as a therapy for CORT-induced hepatic steatosis, and lay the foundation for designing novel prophylactic and therapeutic strategies on CORT-induced hepatic steatosis.

Keywords: Adipose differentiation-related protein, Glucocorticoid receptor, Hepatic steatosis, Network pharmacology, Xiaoyao san

Introduction

Non-alcoholic fatty liver disease (NAFLD) is the most common metabolic liver disease in the world^[1], and nonalcoholic hepatic steatosis (NASH) is a serious consequence of NAFLD^[2]. It is reported that the prevalence

of NASH is estimated to be approximately 3% to 6% in the US population, and more than 20% of patients with NASH progress to cirrhosis, end-stage liver disease, and even death with time^[3]. Hepatic steatosis is also detected in 20% of all hypercortisolism patients^[4-5] who have presented with excessive glucocorticoid secretion or have been subjected to glucocorticoid therapy^[6-7]. Hypercortisolism-related hepatic steatosis is attributed mainly to hyperactivation of the glucocorticoid receptor (GR) *via* the hypothalamic-pituitary-adrenal axis^[8-9] and is regulated by a GR-dependent pathway^[9-10]. GR activation inhibits fatty acid β -oxidation through the peroxisome proliferator-activated receptor pathway^[11]. Heat shock cognate protein 70 (HSP 70) is a GR chaperonin protein that binds perilipin-2 (adipose differentiation related protein, ADFP) which stabilizes lipid droplets^[12]. The foregoing evidence indicates that interference with GR activation and its downstream pathway could ameliorate hypercortisolism-related hepatic steatosis. This condition may potentially develop into steatohepatitis or cirrhosis^[13-14] when therapy is delayed. Sterol regulatory element-binding proteins-1 inhibitor, acetyl-coenzyme A hydroxylase inhibitor, stearoyl-coenzyme A desaturase 1 inhibitor, peroxisomal proliferation-activated receptor activator, and so on have been widely applied in clinics for prevention and therapy hepatic steatosis^[15]. However, the GR antagonists are still few applied for hepatic steatosis.

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Even though mifepristone (RU-486), a well-known GR antagonist, can reverse the negative effects of GR including hepatic steatosis, it is still limited in clinic application because of hypovolemia, hypotension, and infertility^[16]. Hence, a new safe and efficacious therapy against glucocorticoid-induced hepatic steatosis is urgently required. Therefore, in this study, we devote ourselves to discovering an effective and safe therapy against glucocorticoid-induced hepatic steatosis.

Xiaoyao san (XYS) is a classical traditional Chinese medicine (TCM) prescription. It is recorded in *Prescriptions of the Bureau of Taiping People's Welfare (Tai Ping Hui Min He Ji Ju Fang)* from the Song dynasty and has been administered to alleviate depression and liver diseases for >800 years. It has had obvious efficacy and induced few side effects^[17]. YYS consists of *Bupleuri Radix*, *Angelicae Sinensis Radix*, *Paeoniae Radix Alba*, *Atractylodis Macrocephalae Rhizoma*, *Poria*, *Glycyrrhizae Radix et Rhizoma*, *Menthae Haplocalycis Herba*, and *Zingiberis Rhizoma Recens* in a 5:5:5:5:5:4:1:5 weight ratio according to the Xiaoyao wan (condensed pill) entry in the "Chinese Pharmacopoeia 2020 Edition". The active ingredients in YYS were confirmed to have anti-neuritis^[18], anti-hepatitis^[19–20], hepatoprotective^[21–22], and antihepatotoxic^[23] effects. However, the modes of action of YYS against glucocorticoid-induced hepatic steatosis are unknown. In this study, we aimed to investigate whether YYS protects against corticosterone (CORT)-induced hepatic steatosis, and to explore its mechanism. Several reports indicated that YYS restores serum CORT levels, increases energy metabolism, and reduces lipid accumulation^[24–25]. Glycyrrhizin in YYS lowers HSP 90 levels and attenuates GR ligand affinity^[26]. Moreover, aikosaponin D in YYS regulates GR translocation^[27]. GR potentially regulates ADFP by binding HSP 70^[12]. Therefore, we hypothesized that YYS protection against glucocorticoid-induced hepatic steatosis is mediated by GR and ADFP.

Materials and methods

Materials

YYS was prepared by Jiuzhitang Co. Ltd., Changsha, China, in accordance with the formulation process described in the "Chinese Pharmacopoeia 2020 Edition". A 0.42 extraction ratio of YYS was used in this study. YYS extractions were analyzed by liquid chromatography-mass spectrometry (LC-MS, TM Q ExactiveTM Orbitrap LC-MS/MS system, Thermo Scientific, Waltham, MA, USA) and quantified by ultra-high performance liquid chromatography (UHPLC) at 280 nm^[28] (Supplemental Digital Content: YYS extraction was analyzed by UPLC, <http://links.lww.com/AHM/A0>). In brief, the mobile phases comprised eluent A (0.1% formic acid) and eluent B (methanol). The gradient flow was as follows: 0 to 5 min, 10% to 30% B; 5 to 12 min, 30% to 35% B; 12 to 20 min, 35% to 90% B. The analysis was performed at a flow rate of 0.3 mL/min with ACQUITY UPLC HSS T₃ column 1.8 μm, 2.1 mm × 100 mm (Waters, Milford, MA, USA). UHPLC separations were conducted by Dionex Ultimate UHPLC 3000 system (Thermo Scientific, Waltham, MA, USA). The injection volume was 4 μL. The spectrum only identified and recorded the sample components within the first 20 min. CORT and RU-486 were purchased from Sigma-Aldrich Corp. (St. Louis, MO, USA).

Animal experiments

Male C57BL/6J mice aged 3 weeks were purchased from Guangdong Provincial Medical Laboratory Animal Center, Guangdong, China and managed in accordance to protocols approved by the Institutional Animal Care and Use Committee of Jinan University, Guangzhou, China (Approval number: 20130904001). All mice were housed at (22 ± 2)°C under a 12/12 h light/dark cycle and randomly assigned to six groups (*n* = 6). Normal diet (D12450B; 3.84 kcal/g; 10% kcal from fat consisting of 25 g/kg soybean oil and 20 g/kg lard) and high-fat diet (HFD) (D12492; 5.24 kcal/g; 60% kcal from fat consisting of 25 g/kg soybean oil and 245 g/kg lard) were purchased from Research Diet, New Brunswick, NJ, USA. All mice except those administered the normal diet (ND) received a HFD for 14 weeks to induce hepatic steatosis^[29–30]. From the 8th week, the HFD-fed mice were subcutaneously injected with 2 mg/kg CORT (Sigma-Aldrich Corp., St. Louis, MO, USA) once daily for 7 weeks to promote glucocorticoid-induced hepatic steatosis^[31–32]. The daily YYS dosage was determined based on an average adult human body weight of 70 kg, a conversion factor of 9.1 between humans and mice, and a 0.42 extraction rate of YYS. To reflect the protective mechanism of YYS against CORT-induced hepatic steatosis, the low and high experimental YYS dosages (2.56 g/kg and 5.12 g/kg, respectively) were set to fourfold and eightfold the clinical dosage reported having satisfactory efficacy^[17,33–34]. YYS and the GR antagonist RU-486 (12.5 mg/kg; positive control; Sigma-Aldrich Corp., St. Louis, MO, USA) were intragastrically administered and subcutaneously injected, respectively, to the CORT-treated mice once daily for 7 weeks.

The mice were sacrificed under ether anesthesia at the end of the experiment. Plasma was isolated from blood samples treated with heparin sodium. Plasma triglyceride (TG), total cholesterol (TC), and free fatty acid (FFA) levels were determined with commercial kits (Nanjing Jiancheng Bioengineering Co. Ltd., Nanjing, China). The livers were promptly excised for the subsequent experiments.

Histomorphology, immunohistochemistry (IHC), and immunofluorescence

For histomorphology, the liver tissues were fixed in 4% (v/v) paraformaldehyde, embedded in paraffin, cut into 5 μm slices by microtome (RM2235, Leica, Wetzlar, Germany), and stained with H&E and Oil Red O staining according to the manufacturer's instructions. Tissue images (200×) were viewed and recorded with the M8 microscope and scanner (Precipoint M8, Freising, Germany).

For IHC, the sections were incubated with primary antibodies against ADFP (NB110-40877; Novus Biologicals, Littleton, CO, USA) and GR (#12041; Cell Signaling Technology, Danvers, MA, USA). A 3,3'-diaminobenzidine detection kit (Gene Technology, Shanghai, China) was used according to the manufacturer's instructions. Tissue images (400×) were viewed and recorded with the M8 microscope and scanner (Precipoint M8, Freising, Germany).

For immunofluorescence, the liver sections were incubated with primary antibody against ADFP at 4°C overnight, incubated with secondary antibody conjugated to Alexa Fluor 488 at 37°C for 1 h, and stained with 4',6-diamidino-2-phenylindole at 25°C for 10 min.

Immuno-fluorescence images (400×) were viewed and recorded with a Zeiss LSM510 Meta Duo Scan laser scanning confocal microscope (Carl Zeiss AG, Oberkochen, Germany).

Lipid droplet extraction

Lipid droplets were extracted according to the methods previously reported^[35–36]. In brief, liver tissues were gently ground with phosphate-buffered saline in a Dunes homogenizer (Wheaton, Millville, NJ, USA) and the supernatants were collected by centrifugation (12,000 × g, 4°C, 10 min). Then 3 mL supernatant was mixed with 3 mL of 1.08 M sucrose and the mixture was gently layered onto a gradient separation solution consisting of equal volumes of 270 mM sucrose, 135 mM sucrose, and Tris-EDTA-EGTA solution. The gradient solution samples were separated by ultra-high-speed refrigerated centrifugation (150,000 × g, 4°C, 60 min). The milky white upper fraction contained the lipid droplets and was collected.

Reverse transcription quantitative polymerase chain reaction (RT-qPCR) analysis

Total RNA was extracted from the liver samples with TRIzol reagent (Thermo Fisher Scientific, Waltham, MA, USA) and reverse-transcribed to cDNA with a reverse transcription kit (Transgen Biotech, Guangzhou, China). The cDNA samples were combined with specific primers (Table 1) and SYBR qPCR Master Mix (Transgen Biotech, Guangzhou, China). The mixture was amplified on a CFX Connect™ RT-qPCR system. The relative mRNA expression of each gene was normalized with 18s by the $2^{-\Delta\Delta C_t}$ method^[37].

Western blotting

Total protein from the liver tissues and cells was extracted by lysis buffer and quantified with a Pierce® bicinchoninic acid protein assay kit (Thermo Fisher Scientific, Franklin, MA, USA). Ten-microgram protein samples were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis and transferred to polyvinylidene fluoride membranes (EMD Millipore, Billerica, MA, USA). Target proteins were measured by incubating them with primary antibodies including anti-ADFP, anti-GR, and anti-glyceraldehyde-3-phosphate dehydrogenase (GAPDH; Transgen Biotech, Guangzhou, China). The target proteins were then incubated with secondary antibodies and the immunoblots were visualized with Pierce® enhanced chemiluminescence western blotting substrate (Thermo Fisher Scientific, Franklin, MA, USA). GAPDH expression was the quantitative reference. The expression of proteins was quantified by ImageJ software (version 1.46, NIH, Bethesda, MD, USA).

Table 1
Gene primer sequences

Gene	Forward (5'–3')	Reverse (5'–3')
18s	ACGGCTACCACATCC	CAGACTTGCCCTCCA
Plin1	TACCTAGCTGCTTTCTCGGTGTTAC	GGGCTTCTTTGGTGCTGTTGT
Plin2	GGAGTGGAAAGAGAAGCATCG	CAACACAGTGGGACTCATGG
Plin3	TGCTGGGTGATGTCCCTGAAC	GCCTGATGGAATCCGTGAAAG
Plin4	GTGCTGGCTCATTG	TGGCTTGTCTTACC
Plin5	ACGAGGTGGTGGGTGA	GAAGGGAACAGAAGCC

Prediction of potential signaling pathways of YYS

Six constituent herbs in YYS were submitted to BATMAN-TCM (<http://bionet.ncpsb.org/batman-tcm>) to discover their therapeutic mechanisms. After the ingredients in the herbs were identified, the candidate targets were calculated by the similarity-based target prediction method^[38]. Putative targets were predicted because the score cutoff was >30. YYS regulated GR and ADFP expression. Hence, a subsequent functional analysis calculated the intersections of GR, ADFP, the YYS compounds, and the putative YYS targets in DAVID 6.7 (<https://david.ncifcrf.gov>) and the STRING database (<https://string-db.org/>) and indicated the KEGG pathways. The herb-compound-target-pathway network was constructed with Cytoscape 3.7.2.

Plasmid and siRNA transfection

HepG2 cells (American Type Culture Collection, Rockville, MD, USA) were incubated in Roswell Park Memorial Institute 1640 medium with fetal bovine serum. HepG2 cells were transfected with NR3C1 plasmid to overexpress cellular GR. The overexpression-NR3C1 plasmid and the vector (NC) were purchased from Beijing TransGen Biotech Co. Ltd. (Beijing, China). Si-NR3C1 primers were used to knock down NR3C1 (Guangzhou Ribobio Co. Ltd., Guangzhou, China) by targeting the NR3C1 sequence CAACGGTGGCAATGTGAAA. Negative control siRNA primers (si-NC) were purchased from Guangzhou Ribobio Co. Ltd. and used as normal controls. HepG2 cells were transfected with plasmids, siRNAs, and 0.5% Lipofectamine™ LTX (Invitrogen, Carlsbad, CA, USA).

Statistical Analysis

Data are mean ± standard deviation (SD) and were plotted with GraphPad Prism 5 (GraphPad Software, La Jolla, CA, USA). Statistically significant differences between treatment means were analyzed by one-way analysis of variance. $P < 0.05$ represented significant differences among groups.

Results

YYS ameliorated hepatic steatosis in CORT-induced fat liver disease (FLD) mice

The peaks indicated in Supplemental Digital Content (YYS extraction was analyzed by UPLC, <http://links.lww.com/AHM/A0>) were (1) paeoniflorin, (2) ferulic acid, and (3) liquiritin. H&E and Oil Red O staining in Figure 1A and 1B showed that both 2.56 g/kg and 5.12 g/kg YYS reduced the number of fat vacuoles in the livers of the CORT-induced FLD mice. YYS also lowered the levels of plasma lipids including TG, TC, and FFA in a dose-dependent manner in CORT-treated FLD mice (Figure 1C–1E, $P < 0.05$). The GR antagonist RU-486 reversed hepatic steatosis symptoms and lowered the plasma lipid levels in CORT-induced FLD mice (Figure 1, $P < 0.01$).

YYS inhibited GR activation in CORT-induced FLD mice

The effects of YYS on hepatic GR expression in CORT-treated FLD mice were shown in Figure 2. The IHC results in Figure 2A show that YYS decreased nuclear GR translocation in CORT-induced FLD mice in a manner similar to that of the GR antagonist RU-486.

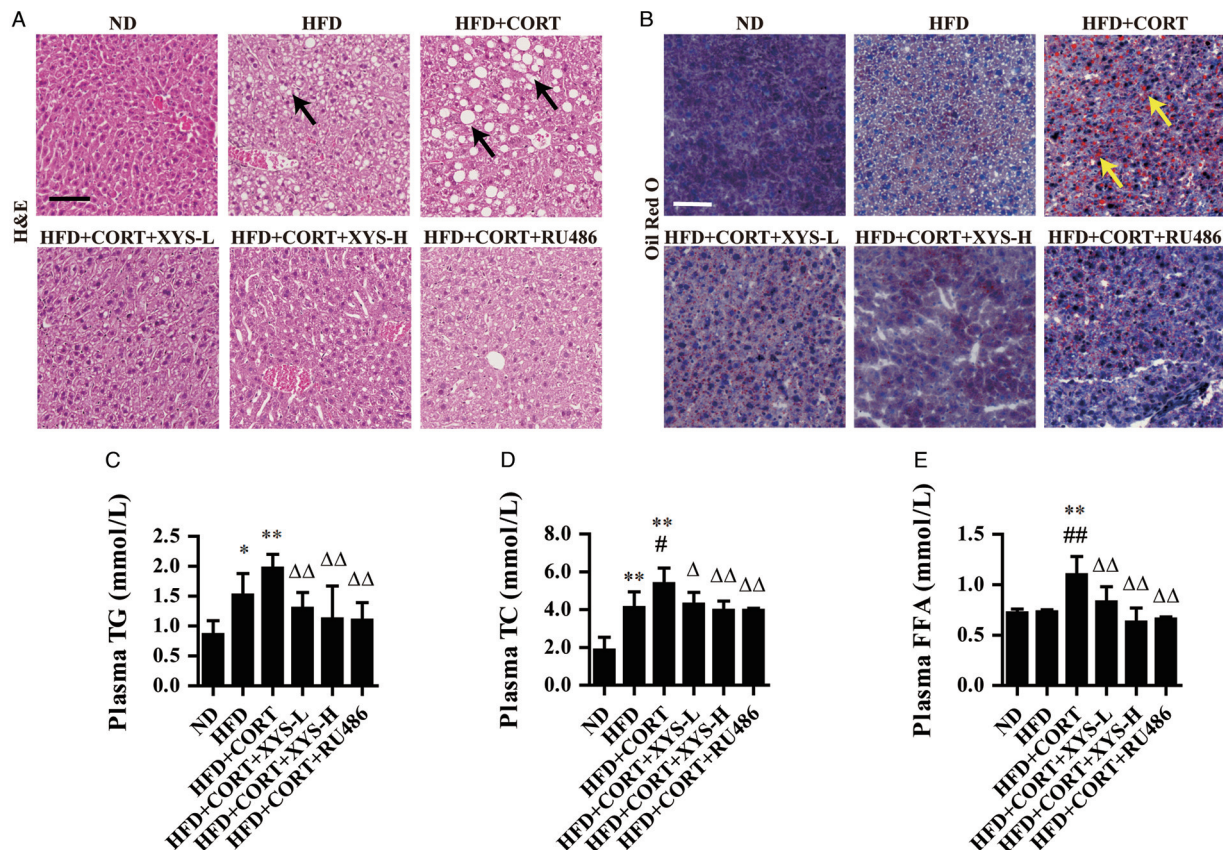


Figure 1. XYS ameliorated hepatic steatosis in CORT-induced FLD mice. HFD mice were used to assess the impact of XYS (2.56 g/kg and 5.12 g/kg daily for 7 weeks) on CORT (2 mg/kg)-induced hepatic steatosis. (A) H&E staining and (B) Oil Red O staining were used to observe morphological changes in liver sections ($n=3$; $200\times$, bar = $50\mu\text{m}$). Dark arrows indicate fat vacuoles and yellow arrows indicate lipid droplets. (C–E) TG ($n=4$), TC ($n=4$), and FFA ($n=5$) levels were measured. Data are mean \pm SD. Significant differences between means were determined by one-way ANOVA. * $P < 0.05$ and ** $P < 0.01$ vs. ND group; # $P < 0.05$ and ## $P < 0.01$ vs. HFD group; $\Delta P < 0.05$ and $\Delta\Delta P < 0.01$ vs. CORT group. ANOVA: analysis of variance; CORT: corticosterone; FFA: free fatty acid; FLD: fatty liver disease; HFD: high-fat diet; ND: normal diet; SD: standard deviation; TC: total cholesterol; TG: triglyceride; XYS-L: Xiaoyao san low dosage; XYS-H: Xiaoyao san high dosage; RU486: mifepristone.

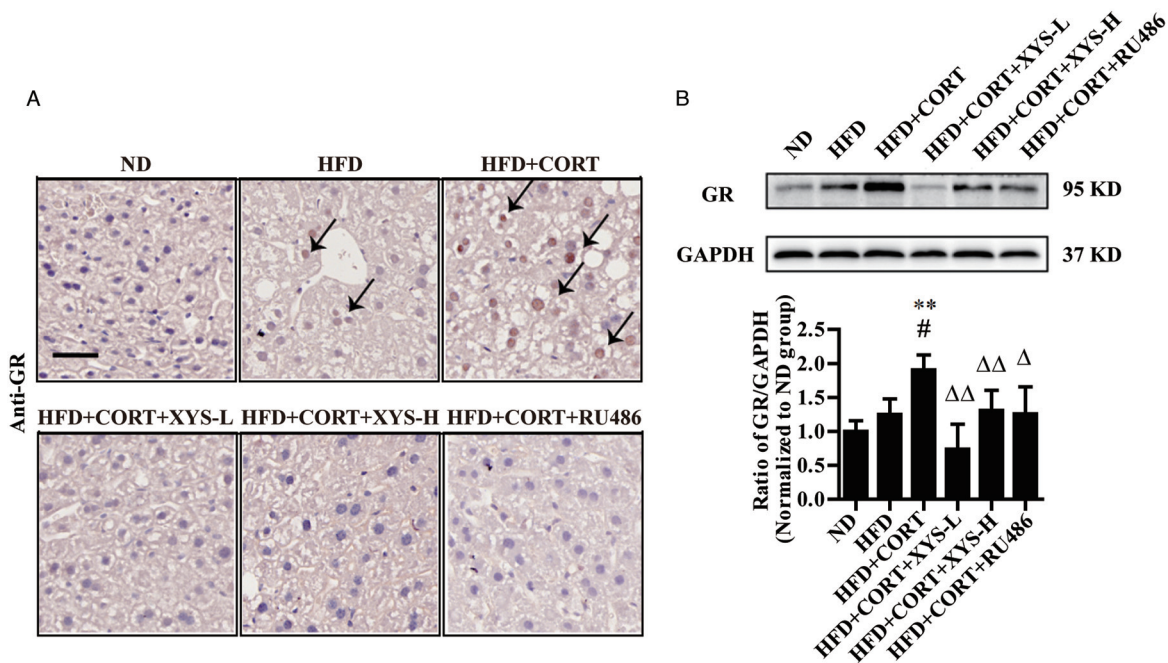


Figure 2. XYS inhibited GR activation in CORT-induced FLD mice. (A) Nuclear GR translocation in livers was detected by IHC. Positive brown staining is marked with dark arrows ($n=3$; $400\times$, bar = $25\mu\text{m}$). (B) GR protein expression in the total lysate was determined by western blotting and quantified with ImageJ (version 1.46, NIH, Bethesda, MD, USA) ($n=3$). Data are mean \pm SD. Significant differences between means were determined by one-way ANOVA. ** $P < 0.01$ vs. ND group; # $P < 0.05$ vs. HFD group; $\Delta P < 0.05$ and $\Delta\Delta P < 0.01$ vs. CORT group. ANOVA: analysis of variance; CORT: corticosterone; GR: glucocorticoid receptor; HFD: high-fat diet; FLD: fatty liver disease; IHC: immunohistochemistry; ND: normal diet; SD: standard deviation; XYS-L: Xiaoyao san low dosage; XYS-H: Xiaoyao san high dosage; RU486: mifepristone.

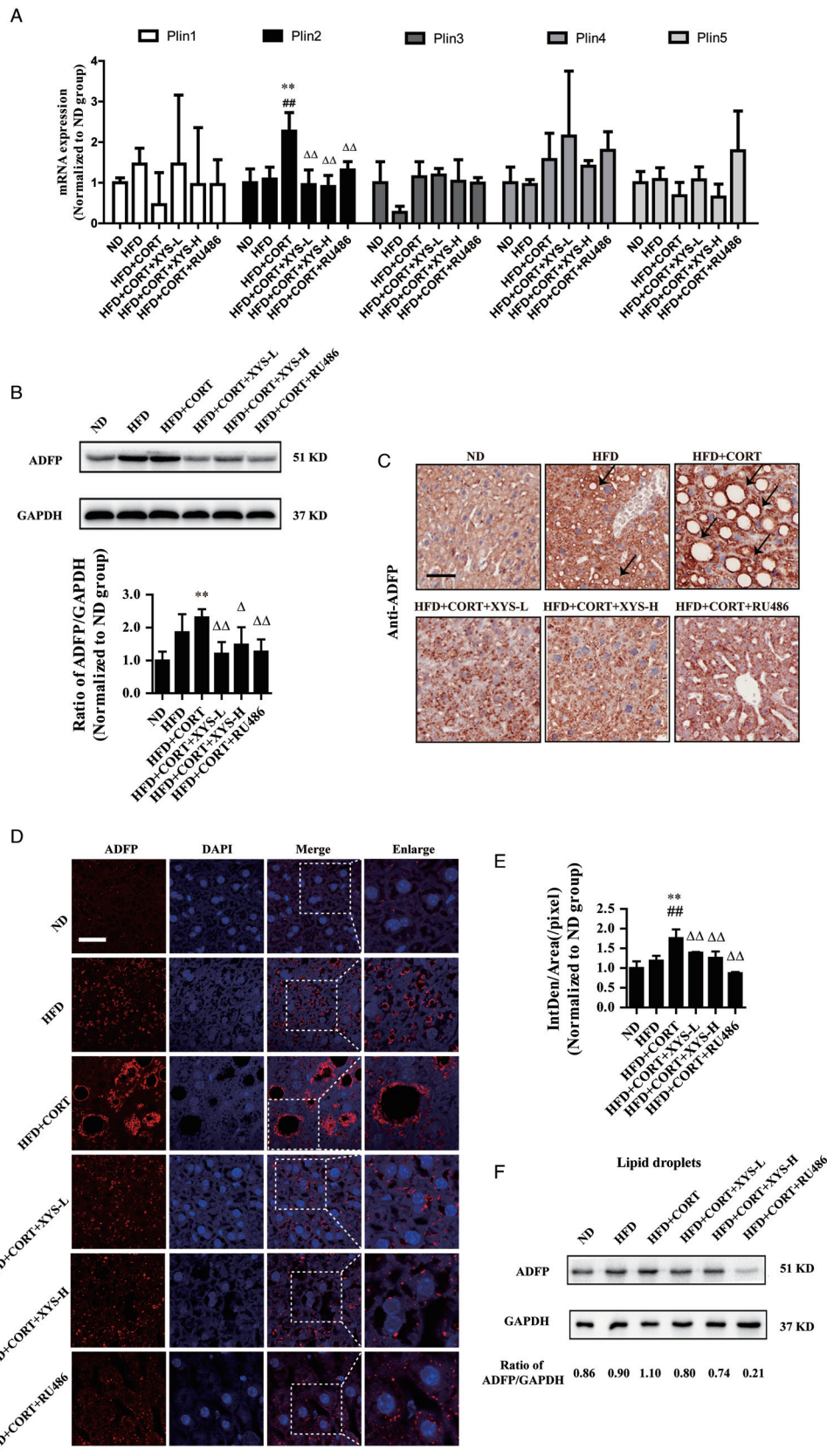
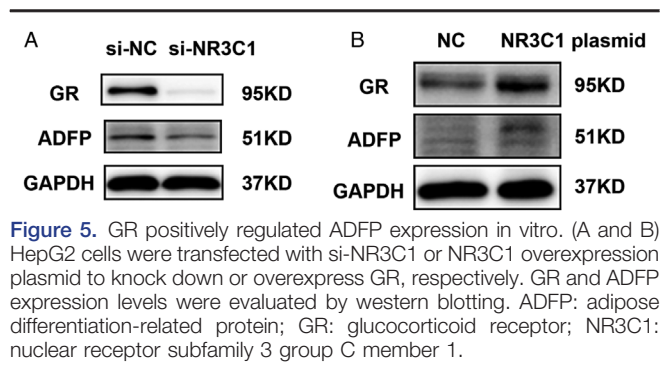


Figure 4. XYs reduced ADFP recruitment and expression around lipid droplets in livers of CORT-induced FLD mice. (A) mRNA and (B) protein expression levels of ADFP were detected by RT-qPCR and western blotting, respectively, and quantified with ImageJ (version 1.46, NIH, Bethesda, MD, USA) ($n=3$). (C) IHC data indicated that ADFP was stained brown. Here, ADFP is labeled with dark arrows ($n=3$; $400\times$, bar = $25\mu\text{m}$). (D) ADFP around lipid droplets was investigated by immunofluorescence and representative images were viewed and recorded by confocal microscopy ($400\times$, bar = $10\mu\text{m}$). (E) Positive fluorescence densities (IntDen/Area/pixel) were quantified with ImageJ (version 1.46, NIH, Bethesda, MD, USA) ($n=3$). (F) After the lipid droplet membranes were extracted, the ADFP expression in the lipid droplet lysate was determined by western blotting. Data are mean \pm SD. Significant differences between means were determined by one-way ANOVA. $**P < 0.01$ vs. ND group; $\#\#P < 0.01$ vs. HFD group; $\Delta P < 0.05$ and $\Delta\Delta P < 0.01$ vs. CORT group. ADFP: adipose differentiation-related protein; CORT: corticosterone; FLD: fatty liver disease; IHC: immunohistochemistry; ND: normal diet; SD: standard deviation; XYs-L: Xiaoyao san low dosage; XYs-H: Xiaoyao san high dosage; RU486: mifepristone; Plin1: perilipin 1; Plin2: perilipin 2; Plin3: perilipin 3; Plin4: perilipin 4; Plin5: perilipin 5; HFD: high-fat diet.



cells showed that GR overexpression upregulated the ADFP protein (Figure 5B).

Discussion

XYS has been clinically administered for Liver-*Qi* Stagnation and Spleen Deficiency Syndrome, and depression^[39]. Its putative mode of action is serum glucocorticoid level reduction^[40]. TCM theory states that YYS has efficacy on the symptoms of energy metabolism disorders such as hypercortisolism-related hepatic steatosis. However, the associated mechanisms of YYS against glucocorticoid-induced FLD are unclear.

Current therapeutic approaches for the management of hepatic steatosis include preventing lipid droplet formation and enlargement^[39], improving lipid oxidation^[41], and decreasing dietary fat intake^[42]. However, reducing fat

consumption was relatively less efficacious at ameliorating hepatic steatosis in obese patients with hypercortisolism^[4-5]. The present study revealed that YYS protected against the development of glucocorticoid-induced hepatic steatosis in CORT-induced FLD mice by mitigating lipid droplets. The results of this study indicated that YYS activates fat turnover and metabolism. We found that YYS inhibited GR activation in CORT-induced hepatic steatosis by decreasing hepatic GR expression and nuclear translocation. Network pharmacology data disclosed that several active components in XFS targeted GR-dependent endocrine resistance and PPAR signaling pathways. The liver-specific lipid droplet membrane protein ADFP is associated with lipid droplet formation and growth in FLD^[43]. Lipolysis was preceded by chaperone-mediated ADFP autophagy^[12,44]. This mechanism shrank lipid droplets. Network pharmacology data revealed that ADFP is involved in YYS-mediated, GR-dependent pathways. Our data further demonstrated that YYS downregulates hepatic ADFP and decreases ADFP recruitment around lipid droplets. We also used loss-of-function and gain-of-function experiments to investigate the relationship between GR and ADFP protein and discovered that GR positively regulated ADFP protein expression. Hence, we identified a new GR/ADFP signaling pathway contributing to the mechanism by which YYS ameliorates hepatic steatosis.

The present study elucidated the effects of YYS on glucocorticoid-induced hepatic steatosis. YYS inhibits ADFP expression by decreasing hepatic GR activation and expression (Figure 6). GR/ADFP signaling pathway

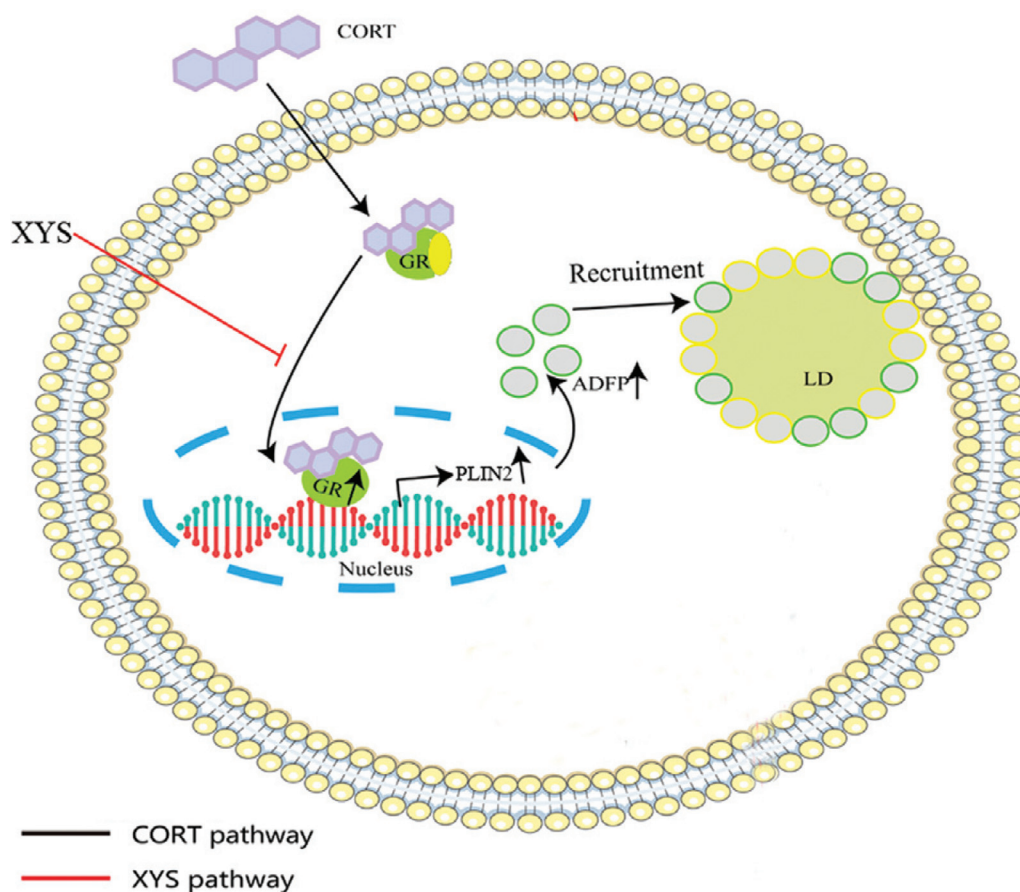


Figure 6. Schematic diagram. YYS ameliorated glucocorticoid-induced hepatic steatosis by downregulating the GR/ADFP signaling pathway. ADFP: adipose differentiation-related protein; GR: glucocorticoid receptor; YYS: Xiaoyao san.

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