Research Article

Antistress Effects of San-Huang-Xie-Xin Decoction on Restraint-Stressed Mice Revealed by $^1$H NMR-Based Metabolomics and Biochemistry Analysis

Wei Peng 1, Huan Du 1, Guangli Liu 1, Qing Zhang 1, Tingting Kuang 1, Zhang Wang 2, and Gang Fan 2

1School of Pharmacy, Chengdu University of Traditional Chinese Medicine, Chengdu 611137, China
2School of Ethnic Medicine, Chengdu University of Traditional Chinese Medicine, Chengdu 611137, China

Correspondence should be addressed to Zhang Wang; wzqcd@163.com and Gang Fan; fangang1111@163.com

Received 6 November 2018; Revised 18 January 2019; Accepted 10 March 2019; Published 16 April 2019

1. Introduction

Previous reports have indicated that stress could negatively affect physical and mental performances of a human being and result in various stress-related disorders, such as anxiety and insomnia [1, 2]. Importantly, it is reported that depression and anxiety, two substantive public health problems, which often lead to heavy social burdens, have the high incidences under stress exposure [3, 4]. However, the current available pharmacotherapies to treat these stress-related disorders are limited due to low efficacy and oppressive side effects [5]. In addition, increasing scientific evidences have suggested that traditional Chinese medicines (TCMs), which have been utilized to treat diseases for thousands of years in China, possess promising curative effects for treating some difficult miscellaneous diseases, including some neuropsychiatric disorders such as depression, anxiety, and insomnia [6, 7].

San-Huang-Xie-Xin decoction (SHXXD), composed of Coptidis Rhizoma, Rhei Radix et Rhizoma, and Scutellariae Radix, is a representative antipyretic and detoxifying prescription in traditional Chinese medicine. In this study, we investigated the antistress effects and underlying mechanisms of San-Huang-Xie-Xin decoction (SHXXD) on restraint-stressed mice by $^1$H NMR-based metabolomics combined with biochemistry assay. A total of 48 male mice (5 weeks old, 18–22 g) were divided randomly into 6 groups (n = 8), including the normal group, restraint-stressed group, vitamin C group (positive drug, 17 mg/kg), and 3-dosage groups of SHXXD (200, 400, and 800 mg/kg). The stress model was induced by restraining mice in a polypropylene centrifuge tube for 6 h every day. The rotarod test was performed, and several biochemical indicators were measured. Moreover, other 24 animals were divided into 3 groups (n = 8) including the normal group, restraint-stressed group, and SHXXD group (800 mg/kg) for $^1$H NMR-based metabolomics analysis. Our results showed that SHXXD significantly increased the rotarod time, thymus index, spleen index, and the levels of glutathione peroxidase (GSH-Px), superoxide dismutase (SOD), and interleukin-2 (IL-2), but decreased the levels of malondialdehyde (MDA), IL-1β, tumor necrosis factor-α (TNF-α), corticosterone (CORT), and adrenocorticotropic hormone (ACTH) in restraint-stressed mice. Moreover, the contents of eight endogenous metabolites that were changed by restraint stress were significantly reversed by SHXXD. The results of both metabolomics and biochemical analysis indicated that SHXXD (800 mg/kg, p.o.) could improve the biochemical changes and metabolic disorders in restraint-stressed mice by antioxidation and anti-inflammation, enhancing the body’s immune function and restoring several disturbed metabolic pathways (i.e., lipid metabolism, glycolysis and gluconeogenesis, inflammatory injury, and energy metabolism). Taken together, these results indicated that SHXXD has a potential antistress effect in restraint-stressed mice and could be considered as a candidate drug for stress-related disorders.
Radix, is a representative antipyretic and detoxifying TCM prescription and first recorded in Synopsis of Golden Chamber (a known TCM monograph written by Zhang Zhongjing) [8]. Current phytochemical researches have revealed that the main chemical constituents of SHXXD are alkaloids, anthraquinones, and flavonoids, including berberine, coptisine, palmatine, baicalin, wogonoside, jatrorrhizine, baicalein, aloe-emodin, rhein, and emodin [9, 10]. In addition, previous investigations reported that SHXXD has various pharmacological effects, such as anti-inflammatory, antioxidant, antimicrobial, antiatherosclerosis, and antitumor activities [11–13]. Besides, there has been a clinical report of mental regulative effects of SHXXD [14]. However, to the best of our knowledge, there is no systemic research regarding the effects of SHXXD against stress-related disorders.

Previous evidences revealed that stress possesses detrimental influences on the functions of several cells due to free radical damage, leading to various oxidative stress-related neuropsychiatric diseases [1]. Moreover, previous studies reported that SHXXD could inhibit oxidative stress and inflammation injury of global cerebral ischemia-reperfusion rats [14, 15]. Consequently, we come to the hypothesis that SHXXD might have beneficial effects against restraint stress-induced disorders, and the possible mechanisms might be correlated to its antioxidant and anti-inflammatory properties. This study was designed to evaluate the antistress effects and underlying mechanisms of SHXXD on restraint-stressed mice by using the 1H NMR-based metabolomics and biochemistry assay methods, which will not only facilitate the understanding of the pathological changes of restraint stress but also provide valuable information for revealing the antistress effects of SHXXD.

2. Materials and Methods

2.1. Animals and Ethics Statement. A total of 72 male ICR (Institute for Cancer Research) mice (5 weeks old, 18-22 g) were purchased from the Chengdu Dashuo Biological Technology Co. Ltd. (Chengdu, China). All animals were housed in standard mouse cages (29 cm × 17.8 cm × 16 cm) at 21 ± 1°C under a 12 h light/dark cycle and had free access to standard pellet diet (Purina Chow) and tap water, and each cage contains 8 mice. Among these animals, 48 mice were used to evaluate the antistress effects of SHXXD, and other 24 mice were used for 1H NMR-based metabolomics analysis. All animal protocols were strictly followed in accordance with the National Institutes of Health Guide concerning the Care and Use of Laboratory Animals, and all the experimental protocols were carried out with the approval of the Animal Experimentation Ethics Committee of the Chengdu University of Traditional Chinese Medicine (No. 2016-12).

2.2. Chemicals and Reagents. ELISA kits of malondialdehyde (MDA), superoxide dismutase (SOD), glutathione peroxidase (GSH-Px), interleukin- (IL-) 1β, IL-2, tumor necrosis factor- (TNF-) α, corticosterone (CORT), and adrenocorticotropic hormone (ACTH) were purchased from the Beijing Yonghui Biological Technology Co. Ltd. (Beijing, China). Deuterium oxide (D2O) was purchased from the Cambridge Isotope Laboratories Inc. (Cambridge, UK). Acetonitrile purchased from Fisher Chemicals (Pittsburgh, PA, USA) was of chromatographic grade. Deionized water was obtained from a hi-tech water purification system (Henan, China). The standard compounds of aloe-emodin, rhein, jatrorrhizine, palmatine, berberine, baicalin, wogonoside, baicalein, and wogonin were purchased from Chengdu Push Biotechnology Co. Ltd. (Chengdu, China). All other chemicals used in this study were of analytical reagent grade.

2.3. Materials and the Preparation of SHXXD. All the herbs were purchased from Sichuan Neautus Traditional Chinese Medicine Co. Ltd. (Chengdu, China) and authenticated by their appearance traits and the thin-layer chromatography method documented in the Chinese Pharmacopoeia. The voucher specimens (Nos. 16031701, 16031702, and 16031703 for the Rhei Radix et Rhiza, Copidis Rhizoma, and Scutellariae Radix, respectively) were deposited in the College of Ethnic Medicine, Chengdu University of Traditional Chinese Medicine, Chengdu, China. The Synopsis of Golden Chamber, a traditional TCM monograph, recorded that SHXXD consists of Rhei Radix et Rhiza 6 g, Copidis Rhizoma 3 g, and Scutellariae Radix 3 g. Therefore, Rhei Radix et Rhiza, Copidis Rhizoma, and Scutellariae Radix at the weight ratio of 2 : 1 : 1, reaching a total weight of 500 g, were extracted twice with pure water (1 : 10, w/v) by decocting (each extraction period lasted 1 h). Subsequently, the decoction was filtered and concentrated under 50°C in vacuum using an RE-2000B rotary vacuum evaporators (Shanghai Yarong Biochemical Instrument Co., Shanghai, China), and the dried residues were treated as the SHXXD extracts which were stored at -20°C before use.

2.4. High-Performance Liquid Chromatography (HPLC) Analysis of SHXXD. A conventional HPLC method was applied to qualitatively and quantitatively analyze the main chemical constituents of SHXXD extract. Chromatographic analysis was performed on an Agilent 1260 Series LC system (Agilent Technologies, Germany), consisted of an online degasser, a quaternary pump, a diode array detector, an autosampler, and a column thermostat. The samples were separated on an InertSustain C18 column (4.6 mm × 250 mm, 5 μm) at a column temperature of 30°C. The mobile phase was acetonitrile (A) and 0.2% phosphoric acid (B) with a gradient elution program of 5–15% A at 0–10 min, 15–20% A at 10–15 min, 20–23% A at 15–40 min, 23–24% A at 40–55 min, 24–35% A at 55–65 min, and 35–45% A at 65–75 min. The flow rate was 1.0 mL/min, and the injection volume was 10 μL. The detection wavelength was set at 254 nm. The concentrations of aloe-emodin, rhein, jatrorrhizine, palmatine, berberine, baicalin, wogonoside, baicalein, and wogonin in SHXXD extract were determined by using the external standard method.

2.5. Animal Grouping and Experimental Protocols. The experimental doses of the testing samples were selected based on commonly used clinical doses of SHXXD for human. The commonly used clinical dosage of SHXXD is 20 g/person/day.
Visceral weight (crude herb equivalent), and this dose converted to mice should be about 2.6 g/kg/day (crude herb equivalent). In addition, the yield of water extract of SHXXD is about 16%, and so the dose for mice regarding water extract of SHXXD is about 400 mg/kg. Consequently, the 400 mg/kg was treated as the middle dose. In this study, we selected the 200, 400, and 800 mg/kg/day as the low, middle, and high dose for mouse experiments, respectively.

After adaption of the environment for 7 days, a total of 48 mice were divided randomly into 6 groups (each group was consisted of 8 mice), including the normal group (normal): normal mice treated with saline; the restraint-stressed group (control): restraint-stressed mice treated with saline; the vitamin C group (VC): restraint-stressed mice pretreated with VC (17 mg/kg/day, p.o.); the SHXXD-L group: restraint-stressed mice pretreated with low-dose of SHXXD (200 mg/kg/day, p.o.); the SHXXD-M group: restraint-stressed mice pretreated with middle dose of SHXXD (400 mg/kg/day, p.o.); and the SHXXD-H group: restraint-stressed mice pretreated with high-dose of SHXXD (800 mg/kg/day, p.o.). All the mice were administered for 14 consecutive days. The restraint-stressed mice were prepared according to the previous reported methods [16]. Briefly, one hour after the administration of VC or SHXXD, all mice except the normal mice were physically restrained in the 50 mL polypropylene centrifuge tube for 6 h once daily lasting for continuous 14 days. Then, the mice received a rotarod test in the 15th day, and blood samples were collected immediately using an orbital blood sampling after the rotarod tests for further ELISA assays, and the mice were subsequently sacrificed by decapitation at the end of the experiment. The brain, thymus, and spleen were dissected from each mouse to determine their visceral indices according to the following formula (1). In addition, brain tissues were also collected for further ELISA assays. The experimental procedure is shown in Figure 1.

\[
\text{%Visceral indices} = \frac{\text{Visceral weight}}{\text{Brain weight}} \times 100. \quad (1)
\]

2.6. Rotarod Test. The rotarod test was carried out according to the previous reported method with minor modifications [17]. After the final 6-hour restraint in the 14th day, the mice were suffered an adaptive training on a rotarod test at a speed of 20 rpm for 2 min by using a ZB-200 rotarod apparatus (Techman Soft Co., Chengdu, China) before the formal rotarod test. On the next day, the standing time on rolling rotarod (Techman Soft Co., Chengdu, China) before the formal tests were su

2.7. Biochemistry Tests. Serum samples were prepared by leaving whole blood to stand at room temperature for coagulation and then centrifuged for 10 min at 4000 rpm. Brain homogenate 10% (w/v) was prepared in phosphate-buffered saline (PBS) and centrifuged at 4000 rpm for 10 min under 0°C. All the testing samples were stored at -80°C until ELISA analysis. Furthermore, the contents of CORT, ACTH, IL-1β, IL-2, and TNF-α in serum and contents of MDA, SOD, and GSH-Px in the brain were determined using commercial ELISA kits according to the manufacturer’s protocols and instructions.

2.8. Statistical Analysis. Data are represented as the mean ± S.D. One-way analysis of variance (ANOVA) followed by Dunnett’s test was done with GraphPad Prism software version 5.0 (GraphPad Software Inc., San Diego, CA, USA) to detect intergroup differences. A probability (p) value of less than 0.05 was considered statistically significant.

2.9. 1H NMR-Based Metabolomics Analysis

2.9.1. Sample Preparation. In view of the good results of the high-dose group of SHXXD in biochemical tests, its serum samples were further used for metabolomics analysis. After adaption of the environment for 7 days, a total of 24 mice were divided randomly into 3 groups (each group was consisted of 8 mice), including the normal group (normal): normal mice treated with saline; the restraint-stressed group (control): restraint-stressed mice treated with saline; and the SHXXD group: restraint-stressed mice pretreated with high-dose of SHXXD (800 mg/kg/day, p.o.). All the mice were administered for 14 consecutive days. Blood samples were collected using an orbital blood sampling after the final 6-hour restraint in the 14th day.

1H NMR sample preparation was performed according to the method described in previous investigations with some modifications [18]. Briefly, 200 μL serum samples were added to a 1.5 mL centrifuge tube, and then 100 μL D2O and 300 μL NaH2PO4/K2HPO4 buffer (45 mM, pH 7.4) were added. Furthermore, the mixture was centrifuged for 10 min at 12,000 rpm, and subsequently, 550 μL supernatant was transferred into a standard 5 mm NMR tube. Finally, the NMR tube was sealed and further applied for 1H NMR measurement.

2.9.2. 1H NMR Spectroscopy Analysis. All 1H NMR spectra were recorded on an Agilent DD2 600 MHz NMR spectrometer (Agilent Technologies, Santa Clara, USA) operating at a proton frequency of 600.13 MHz. 1H NMR experiments were performed with the following parameters: relaxation delay, 2.0 s; acquisition time, 1.36 s; number of scans, 64; pulse program, cpmpgr1d (Bruker notation); and spectral width, 12,019.2 Hz.

2.9.3. Data Processing and Statistical Analysis. All the sample spectra were phased and manually baseline corrected using MestReNova software (version 9.0.1, Mestrelabs Research SL). The sample spectra were referenced to a lactate signal at 4.11 ppm. In addition, they were segmented into consecutively nonoverlapping regions of 0.002 ppm chemical shift bins between 0.50 and 9.50 ppm. The region from 4.68 to 5.00 ppm, which contained the residual water signal, was excluded. The spectra were normalized to the total area of metabolites’ spectral integrals to compensate for differences in sample concentration. The 1H NMR spectra were automatically reduced to ASCII files using MestReNova software. Output data were imported into SIMCA software (version 11.5, Umetrics, Umeå, Sweden) for multivariate statistical
analysis. Metabolites that were detected were assigned by referring to relevant literatures [19–24] and querying several databases (Madison, http://mmcd.nlm.nih.gov/; KEGG, http://www.genome.jp/kegg/; and HMDB, http://www.hmdb.ca/).

The Pareto-scaled NMR data were analyzed with nonsupervised principal component analysis (PCA) and supervised partial least squares discriminant analysis (PLS-DA) using SIMCA software (version 11.5, Umetrics, Umeå, Sweden). The parameters of $R^2$Y indicated the validity of the models against overfitting, while the parameters of $Q^2$ indicated the predictive ability. Moreover, the mean integral areas of bins belonging to the potential biomarkers that were obtained from PLS-DA were calculated and normalized to the total spectral intensity. To determine if changes in bin intensities were statistically significant fold changes, Student’s $t$-test was performed using MetaboAnalyst 4.0 (http://www.metaboanalyst.ca/). Altered features were considered significant when pertaining $p$ values were less than 0.05.

3. Results

3.1. The Chemical Constituents of SHXXD Determined by HPLC. The HPLC chromatogram of SHXXD extract is shown in Figure 2. A total of 9 constituents including aloe-emodin, rhein, jatrorrhizine, palmitate, berberine, baicalin, wogonoside, baicalein, and wogonin were identified by comparing their retention time and ultraviolet spectra with those of the standard compounds. Among them, aloe-emodin and rhein are mainly from Rhei Radix et Rhizoma; jatrorrhizine, palmitate, and berberine are from Coptidis Rhizoma; and baicalin, wogonoside, baicalein, and wogonin are from Scutellariae Radix. Moreover, the contents of the nine compounds were calculated according to the external standard method. It was found that baicalin showed the highest level in SHXXD extract and its content was 49.63 mg/g, followed by wogonoside (11.99 mg/g). The other 7 compounds (i.e., rhein, aloe-emodin, baicalein, wogonin, jatrorrhizine, palmitate, and berberine) showed low concentrations in SHXXD extract, and their contents are 1.80, 0.89, 1.29, 2.96, 1.94, 3.70, and 5.06 mg/g, respectively.

3.2. Effects of SHXXD on the Rotarod Time of Restraint-Stressed Mice. As shown in Figure 3(a), after the 14-day restrain stress, the rotarod time of restraint-stressed mice (control group) was significantly decreased compared with the normal mice ($p < 0.01$). However, our results revealed that treatment with vitamin C ($p < 0.01$) and SHXXD at low ($p < 0.05$) and high ($p < 0.01$) doses could significantly increase the rotarod time, compared with the control group.

3.3. Effects of SHXXD on Thymus and Spleen Indexes of Restraint-Stressed Mice. Furthermore, the organ indexes of the thymus and spleen were determined in our present study, and the results shown in Figures 3(b) and 3(c) indicated that the thymus and spleen indexes were significantly lower in restraint-stressed mice than those in the normal mice ($p < 0.01$). After treatment with vitamin C, the spleen index was significantly increased ($p < 0.05$) compared with the control group. Similarly, SHXXD treatment at middle and high doses also reversed the decrease in the spleen index ($p < 0.05$), compared with the control group. In addition, administration of SHXXD at middle and high doses could also increase significantly the thymus index ($p < 0.05$), compared with the control group.

3.4. Effects of SHXXD on Serum CORT and ACTH of Restraint-Stressed Mice. As can be seen from Figure 4, the levels of CORT (Figure 4(a)) and ACTH (Figure 4(b)) in the serum of the restraint-stressed mice were significantly higher than those in the normal mice ($p < 0.01$). However, our results revealed that vitamin C could significantly decrease the serum contents of CORT ($p < 0.05$) and ACTH ($p < 0.01$), compared with the control group. Similarly, treatment with SHXXD at low, middle, and high doses also significantly reduced the serum levels of CORT ($p < 0.05$) and ACTH ($p < 0.05$), compared with the control group.

3.5. Effects of SHXXD on Serum Cytokines of Restraint-Stressed Mice. As shown in Figures 5(a)–5(c), the levels of TNF-$\alpha$ ($p < 0.01$) and IL-1$\beta$ ($p < 0.001$) were significantly higher, whereas the IL-2 level was significantly lower ($p < 0.01$) in the serum of the restraint-stressed mice than that in the normal mice. However, vitamin C could significantly decrease the levels of TNF-$\alpha$ ($p < 0.05$) and IL-1$\beta$ ($p < 0.01$) compared with the control group (Figures 5(a) and 5(b)). Interestingly, similar to the vitamin C, SHXXD at low and high doses could significantly decrease the serum levels of TNF-$\alpha$ ($p < 0.05$) and IL-1$\beta$ ($p < 0.01$), and the

![Figure 1: Experimental procedure of the antistress activity evaluation of SHXXD.](image-url)
middle dose of SHXXD could also reduce the TNF-α level (\(p < 0.05\)), compared with the control group (Figures 5(a) and 5(b)). Besides, our results also suggested that SHXXD at middle and high doses could increase significantly the IL-2 level (\(p < 0.05\)), compared with the control group (Figure 5(c)).
3.6. Effects of SHXXD on GSH-Px, MDA, and SOD of Restraint-Stressed Mice. The effects of SHXXD on the levels of GSH-Px, MDA, and SOD in brain tissues of restraint-stressed mice are shown in Figures 5(d)–5(f). After the 14-day restraint stress, the levels of GSH-Px (p < 0.01) and SOD (p < 0.01) in brain tissues of the restraint-stressed group were significantly reduced, whereas the MDA (p < 0.01) was significantly increased, compared with the normal mice. However, our results revealed that vitamin C treatment could significantly enhance the levels of GSH-Px (p < 0.05) (Figure 5(d)) and SOD (p < 0.05) (Figure 5(e)) in brain tissues compared with the control group, but reduced the level of MDA (p < 0.01) (Figure 5(f)). Similar to the vitamin C, SHXXD at the high dose could significantly increase the levels of GSH-Px (p < 0.01) (Figure 5(d)) and SOD (p < 0.01) (Figure 5(e)) in brain tissues, compared with the control group. Furthermore, SHXXD at the low, middle, and high doses could significantly reduce the MDA level in brain tissues (p < 0.05) (Figure 5(f)), compared with the control group.

3.7. Effects of SHXXD on Serum Metabolites of Restraint-Stressed Mice by 1H NMR Metabolomics Analysis. In this study, the 1H NMR-based metabolomics method was carried out to study the effects of SHXXD on the serum metabolites of restraint-stressed mice. The representative 1H NMR spectra of the serum samples from the normal, restraint-stressed, and SHXXD groups are displayed in Figure 6 with major metabolites labelled. By referencing some literatures [19–24] and comparison with several databases, a total of 29 metabolites were identified including tyrosine, formate, histidine, phenylalanine, α-glucose, β-glucose, threonine, lactate, inositol, 1,2-propanediol, glycerol, phosphocholine, choline, malonic acid, lysine, N,N-dimethylglycine, creatinine, citrate, glutamine, 3-hydroxybutyrate, N-acetyl glycoprotein, methionine, acetate, acetate, alanine, valine, isoleucine, low-density lipoprotein (LDL) and very low-density lipoprotein (VLDL), and lipids (saturated as well as unsaturated).

In order to find patterns in the 1H NMR data and differential metabolites, multivariate data analyses were performed. Firstly, PCA, a classical unsupervised approach, was performed to detect patterns in the serum metabolites. The normal, restraint-stressed, and SHXXD groups cannot be clearly separated in the PCA score plot (data not shown). Subsequently, PLS-DA, a supervised pattern recognition method, was further conducted to maximize the separation between groups. The score plot of PLS-DA (Figure 7(a)) showed that this model could reliably differentiate the restraint-stressed group from the normal group (with $R^2 = 0.850$, $R^2_Y = 0.935$, and $Q^2 = 0.803$), suggesting significant metabolic changes induced by restraint stress. Similarly, a clear separation between the control group and the SHXXD group (Figure 7(c)) was obtained by the PLS-DA analysis (with $R^2 = 0.839$, $R^2_Y = 0.898$, and $Q^2 = 0.757$), which indicated that their metabolic profiles were different. Moreover, the corresponding loading plot of PLS-DA revealed the metabolites having significant contributions to the intergroup differences. The loading plot (Figure 7(b)) showed that the separation between the restraint-stressed group and the normal group mainly arose from the signals of lactate, alanine, N-acetyl glycoproteins, glycerol, choline, acetate, LDL/VLDL, lipids, phosphocholine, and valine. In addition, eight metabolites (i.e., lactate, alanine, N-acetyl glycoproteins, glycerol, choline, lipids, acetate, and valine) were found to play an important role in separating the SHXXD group from the restraint-stressed group (Figure 7(d)). These results indicated that these metabolites were significantly affected by the restraint-stressed procedure and drug administration and could serve as potential biomarkers. Table 1 shows these 10 discriminatory metabolites.

Moreover, in order to validate the PLS-DA results, mean integral areas of these 10 metabolites in normalized spectral bins were calculated and compared in the three groups by the use of Student's t-test as reported in Figure 8 and Table 1. The results indicated that lactate, LDL/VLDL, alanine, lipids, N-acetyl glycoproteins, glycerol, choline, and phosphocholine levels were significantly increased, whereas acetate and valine were significantly decreased by the restraint stress procedure. These findings were in agreement with several previous reports [22–24]. After drug

**Figure 4:** Effects of SHXXD on serum CORT (a) and ACTH (b) levels. The data are presented as the mean ± S.D. (n = 8); *p < 0.05 and **p < 0.01 vs. the control group.

**Figure 6:** Restraint-Stressed Mice. 3.7. E

dose could significantly reduce the MDA level in brain tissues compared with the control group. Furthermore, SHXXD at the low, middle, and high doses significantly increased, compared with the normal mice. However, our results revealed that vitamin C treatment could significantly enhance the levels of GSH-Px, MDA, and SOD in brain tissues of restraint-stressed mice. The representative 1H NMR spectra were identified including tyrosine, formate, histidine, phenylalanine, α-glucose, β-glucose, threonine, lactate, inositol, 1,2-propanediol, glycerol, phosphocholine, choline, malonic acid, lysine, N,N-dimethylglycine, creatinine, citrate, glutamine, 3-hydroxybutyrate, N-acetyl glycoprotein, methionine, acetate, acetate, alanine, valine, isoleucine, low-density lipoprotein (LDL) and very low-density lipoprotein (VLDL), and lipids (saturated as well as unsaturated). In order to find patterns in the 1H NMR data and differential metabolites, multivariate data analyses were performed. Firstly, PCA, a classical unsupervised approach, was performed to detect patterns in the serum metabolites. The normal, restraint-stressed, and SHXXD groups cannot be clearly separated in the PCA score plot (data not shown). Subsequently, PLS-DA, a supervised pattern recognition method, was further conducted to maximize the separation between groups. The score plot of PLS-DA (Figure 7(a)) showed that this model could reliably differentiate the restraint-stressed group from the normal group (with $R^2 = 0.850$, $R^2_Y = 0.935$, and $Q^2 = 0.803$), suggesting significant metabolic changes induced by restraint stress. Similarly, a clear separation between the control group and the SHXXD group (Figure 7(c)) was obtained by the PLS-DA analysis (with $R^2 = 0.839$, $R^2_Y = 0.898$, and $Q^2 = 0.757$), which indicated that their metabolic profiles were different. Moreover, the corresponding loading plot of PLS-DA revealed the metabolites having significant contributions to the intergroup differences. The loading plot (Figure 7(b)) showed that the separation between the restraint-stressed group and the normal group mainly arose from the signals of lactate, alanine, N-acetyl glycoproteins, glycerol, choline, acetate, LDL/VLDL, lipids, phosphocholine, and valine. In addition, eight metabolites (i.e., lactate, alanine, N-acetyl glycoproteins, glycerol, choline, lipids, acetate, and valine) were found to play an important role in separating the SHXXD group from the restraint-stressed group (Figure 7(d)). These results indicated that these metabolites were significantly affected by the restraint-stressed procedure and drug administration and could serve as potential biomarkers. Table 1 shows these 10 discriminatory metabolites.

Moreover, in order to validate the PLS-DA results, mean integral areas of these 10 metabolites in normalized spectral bins were calculated and compared in the three groups by the use of Student’s t-test as reported in Figure 8 and Table 1. The results indicated that lactate, LDL/VLDL, alanine, lipids, N-acetyl glycoproteins, glycerol, choline, and phosphocholine levels were significantly increased, whereas acetate and valine were significantly decreased by the restraint stress procedure. These findings were in agreement with several previous reports [22–24]. After drug...
administration, the levels of these metabolites that were changed by the restraint stress were significantly reversed by SHXXD except for LDL/VLDL and phosphorylcholine. These results suggested that SHXXD has a therapeutic effect in adjusting the changed levels of metabolites in restraint-stressed mice.

Figure 5: Effects of SHXXD on serum cytokines (IL-1β (a), TNF-α (b), and IL-2 (c)) and GSH-Px (d), MDA (e), and SOD (f) in the brain of restraint-stressed mice. Each point represents an individual sample. Error bars indicate the mean ± S.D. (n = 8). *p < 0.05, **p < 0.01, and ***p < 0.001 vs. the control group.
4. Discussion

Increasing researches have revealed that stress-related mental disorders show the rising incidence due to the intensifying competitions and rapid developing society [25, 26]. Furthermore, herbal medicines are reported be a precious resource for discovering novel drugs for treating various diseases [27, 28]. To the best of our knowledge, this study is the first systemic report on ameliorative effects of SHXXD, a representative antipyretic and detoxifying herbal medicinal prescription in China, against restraint stress-induced symptoms in mice.

Previous investigations have demonstrated that restraint is an ideal and preferred way for preparing stressed animals, and the restraint-stressed mouse model has been extensively used to investigate the pathology of stress-related disorders and to screen the efficacy of candidate antistress drugs [1, 29]. To the best of our knowledge, this study is the first systemic report on ameliorative effects of SHXXD, a representative antipyretic and detoxifying herbal medicinal prescription in China, against restraint stress-induced symptoms in mice.

Activation of the hypothalamic-pituitary-adrenal (HPA) axis plays an important role in physiological response of animals to stress and commonly results in immune organs atrophy in stress, such as the thymus and spleen [1, 6]. Furthermore, HPA axis activation could also increase the serum levels of CORT and ACTH. Previous reports have demonstrated that restraint stress could activate the HPA axis, and downregulating the abnormal activation of HPA axis might be beneficial for controlling stress-related disorders [34]. Restraint stress could induce chronic fatigue by overreleasing various hormones such as CORT, ACTH, and catecholamines [35, 36]. The rotarod test is a commonly used way for evaluating the antifatigue effects of candidate drugs [37]. In the present study, we found that SHXXD could decrease the serum levels of CORT and ACTH, but increase the organ indexes of the thymus and spleen. Besides, the results indicated that SHXXD treatment could prolong the rotarod time of the restraint-stressed mice. All these results mentioned above suggested that SHXXD could ameliorate the stress-induced fatigue and reduce the abnormal activation of the HPA axis in stress.
Figure 7: Score plots and corresponding loading plots generated from pairwise PLS-DA analysis of $^1$H NMR data: (a) score plot of the normal and restraint-stressed groups; (b) loading plot of the normal and restraint-stressed (control) groups; (c) score plot of the restraint-stressed and SHXXD groups; (d) loading plot of the restraint-stressed and SHXXD groups.

Table 1: Potential biomarkers in mouse serum and their variations among groups.

| No. | Metabolite               | Chemical shift (ppm)$^a$ | Assignment                  | Control vs. normal$^b$ | SHXXD vs. control$^b$ |
|-----|--------------------------|--------------------------|-----------------------------|------------------------|------------------------|
| 1   | LDL/VLDL                 | 0.87 (br), 1.27 (br)     | $-\text{CH}_3$, $-\text{CH}_2$ | $\uparrow^*$           | —                      |
| 2   | Lactate                  | 1.32 (d), 4.11 (q)      | $\beta$–$\text{CH}_3$, $\alpha$–$\text{CH}$ | $\uparrow^{**}$        | $\downarrow^{**}$      |
| 3   | Alanine                  | 1.47 (d)                 | $\beta$–$\text{CH}_3$ | $\uparrow^*$           | $\downarrow^*$         |
| 4   | Lipids                   | 1.57 (br)                | $-\text{CH}_2$–$\text{CH}_2$–$\text{C}=\text{O}$ | $\uparrow^{***}$       | $\downarrow^{**}$      |
| 5   | N-acetyl glycoproteins   | 2.04 (s)                | $-\text{CH}_3$ | $\uparrow^{***}$       | $\downarrow^{***}$     |
| 6   | Glycerol                 | 3.56 (m), 3.69 (m)      | $-\text{CH}_2$, $-\text{CH}_2$ | $\uparrow^{***}$       | $\downarrow^{***}$     |
| 7   | Choline                  | 3.20 (s)                | $-\text{N}$–$(\text{CH}_3)_3$ | $\uparrow^*$           | $\downarrow^*$         |
| 8   | Phosphorylcholine        | 3.22 (s)                | $-\text{N}$–$(\text{CH}_3)_3$ | $\uparrow^*$           | —                      |
| 9   | Acetoacetate             | 2.23 (s)                | $-\text{CH}_3$ | $\uparrow^*$           | $\downarrow^*$         |
| 10  | Valine                   | 0.98 (d), 1.03 (d)      | $\gamma$–$\text{CH}_3$, $\gamma$–$\text{CH}_3$ | $\downarrow^*$         | $\uparrow^*$           |

$^a$: singlet; d: doublet; q: quartet; m: multiplet; br: broad resonance. $^b$: Metabolites with “$\uparrow$” means increased/decreased; “$\downarrow$” means no significant differences. $^*$, $^{**}$, and $^{***}$ indicate $p < 0.05$, $p < 0.01$, and $p < 0.001$, respectively.
Previous investigations also revealed that besides the immune organs, the serum level of IL-2, released by activated T cells and closely related to the body’s immunity, is also reduced by various stress conditions [6]. Interestingly, our results indicated that SHXXD could increase the IL-2 level in the serum of restraint-stressed mice as well as the thymus and spleen indices, suggesting that SHXXD could enhance the body’s immunity of restraint-stressed mice. It is reported that increased releases of proinflammatory cytokines such as TNF-α and IL-1β commonly appeared in individuals with stress-related disorders [38], and the present results revealed that SHXXD could suppress the overexpressed proinflammatory cytokines of TNF-α and IL-1β. Stress has harmful effects on cell functions via damaging antioxidant defense system, resulting in oxidative and finally leading to various oxidative stress-related diseases [39]. It is reported that GSH-Px and SOD are two major intracellular antioxidants for antioxidative stress [30]. Lipid peroxidation is recognized as a potential molecular mechanism of cell or tissue damages during oxidative stress, and reactive oxygen species in the brain induced by stress could result in lipid peroxidation [40]. MDA is one of the major end products of lipid peroxidation and is commonly used as an indicator for oxidative injury [41]. Our present findings revealed that SHXXD could increase the levels of GSH-Px and SOD, but decrease the MDA in brain tissues of restraint-stressed mice, indicating that SHXXD might have potential antioxidant activity, which is beneficial to maintain the normal antioxidant defense system.

Metabolomics is a useful strategy for investigating the holistic metabolic changes and potential metabolic pathways of TCMs in vivo [21, 23]. In this study, the effects of SHXXD on restraint-stressed mice were investigated for the first time using a 1H NMR-based metabolomics approach combined with statistical analysis. The results indicated that SHXXD treatment could change the endogenous metabolites in restraint-stressed mice, and the main biomarkers are lactate, alanine, N-acetyl glycoproteins, glycerol, choline, lipids, acetocacetate, and valine. Furthermore, by analyzing the biochemical processes of these biomarkers, we found that the antistress effects of SHXXD might be predominantly related to the glycolysis and gluconeogenesis, lipid metabolism, inflammatory injury, and energy metabolism.

Alanine and lactate play important roles in various biochemical processes. Lactate is the end product of glycolysis. The two metabolites are easily converted from pyruvate by alanine aminotransferase (ALT) and lactate dehydrogenase (LDH), respectively [19]. Some studies have found that restraint stress can cause a significant increase in serum ALT and LDH activities [42, 43]. In addition, chronic restraint stress for 30 days could impair glucose metabolism in rats [44]. Therefore, in the present study, the higher levels of alanine and lactate were observed in the serum of restraint-stressed mice than those in normal mice, which might mean a disorder of glycolysis and gluconeogenesis induced by restraint stress. The increased alanine level might be caused by the transformation of acetyl-CoA, while gluconeogenesis from amino acids was suppressed [20]. Moreover,
the increase of serum lactate was related to anaerobic cell respiration, which is a kind of energy metabolism process. After SHXXD intervention, the results showed a tendency of bringing the levels of alanine and lactate back to normal, suggesting that SHXXD could adjust the disorder of glycolysis and gluconeogenesis caused by restraint stress. Choline, an important membrane phospholipid constituent, was observed to be increased significantly in the restraint-stressed mice compared with the level in the normal mice, which would be the evidence that lipid metabolism and membrane fluidity were disrupted. Besides, considerably increased levels of glycerol and lipids were observed in the restraint-stressed mice, suggesting a dysfunction of lipid metabolism. After SHXXD intervention, the levels of glycerol, lipids, and choline were decreased in the restraint-stressed mice, indicating an inflammatory response resulting from the restraint stress procedure. Moreover, after SHXXD treatment, the level of N-acetyl glycoproteins was remarkably attenuated, indicating that SHXXD could alleviate the lipid metabolism disorder caused by restraint stress. N-acetyl glycoproteins are acute-phase reaction glycoproteins, which are related to the inflammatory response [21]. They increased when the body was subjected to external or internal challenges, such as infection, inflammation, or stress [21]. In the present study, the increased N-acetyl glycoproteins in the restraint-stressed mice provided evidence for an inflammatory response according to the results of biochemical tests. In this study, acetoacetate was observed to be decreased significantly in the restraint-stressed mice compared with the level in the normal group, suggesting that a disorder of energy metabolism may have been present in restraint-stressed mice. In addition, valine plays an important role in the TCA cycle after transformation into succinyl-CoA [24]. In the present study, the level of valine was significantly reduced in restraint-stressed mice measured by 1H NMR, indicating an impairment of the TCA cycle. Moreover, the increase in serum lipids indicated a reduction of energy production. After SHXXD treatment, acetoacetate, valine, and lipid levels all show a tendency to return to normal status. These changes suggested that SHXXD could regulate the energy metabolism of the restraint-stressed mice and maintain the energy balance of the body.

5. Conclusion

In conclusion, our present study firstly investigated the anti-stress effects of San-Huang-Xie-Xin decoction (SHXXD) on restraint-stressed mice in vivo. The results revealed that SHXXD could ameliorate the restraint stress-induced disorders, and the underlying mechanisms may be related to its antioxidant and anti-inflammatory, enhancing the body’s immune function and restoring several disturbed metabolic pathways (i.e., lipid metabolism, glycolysis and gluconeogenesis, inflammatory injury, and energy metabolism). Further investigations are needed to fully elucidate the antistress mechanism of SHXXD.

Data Availability

The data used to support the findings in this paper are available from the corresponding author upon request.

Conflicts of Interest

The authors declare that they have no conflicts of interest.

Authors’ Contributions

Wei Peng and Huan Du contributed equally to this work.

Acknowledgments

The authors gratefully acknowledge the financial support from the Cultivation Program of Outstanding Young Academic and Technological Leaders of Sichuan Province (No. 2014JQ0050) and the National Key Research and Development Program of China (No. 2017YFC1703900).

References

[1] T. Buynitsky and D. I. Mostofsky, “Restraint stress in biobehavioral research: recent developments,” Neuroscience and Biobehavioral Reviews, vol. 33, no. 7, pp. 1089–1098, 2009.

[2] A. M. Bao, G. Meynen, and D. F. Swaab, “The stress system in depression and neurodegeneration: focus on the human hypothalamus,” Brain Research Reviews, vol. 57, no. 2, pp. 531–553, 2008.

[3] J. M. Jung, S. J. Park, Y. W. Lee et al., “The effects of a standardized Acanthopanax koreanum extract on stress-induced behavioral alterations in mice,” Journal of Ethnopharmacology, vol. 148, no. 3, pp. 826–834, 2013.

[4] P. Han, T. Han, W. Peng, and X. R. Wang, “Antidepressant-like effects of essential oil and asarone, a major essential oil component from the rhizome of Acorus tatarinowii,” Pharmaceutical Biology, vol. 51, no. 5, pp. 589–594, 2013.

[5] W. W. Chen, R. R. He, Y. F. Li, S. B. Li, B. Tsoi, and H. Kurihara, “Pharmacological studies on the anxiolytic effect of standardized Schisandra lignans extract on restraint-stressed mice,” Phytomedicine, vol. 18, no. 13, pp. 1144–1147, 2011.

[6] S. Lee, D. H. Kim, J. W. Jung et al., “Schizandra chinensis and Scutellaria baicalensis counter stress behaviors in mice,” Phytotherapy Research, vol. 21, no. 12, pp. 1187–1192, 2007.

[7] C. Wang, C. Wu, Z. Yan, and X. Cheng, “Ameliorative effect of Xiaoyao-jieyu-san on post-stroke depression and its potential mechanisms,” Journal of Natural Medicines, vol. 73, no. 1, pp. 76–84, 2019.

[8] T. T. Miao, “The research progress of San-Huang-Xie-Xin decoction,” Hunan Journal of Traditional Chinese Medicine, vol. 32, no. 3, pp. 190–191, 2016.

[9] J. J. Liu, G. J. Zhang, H. Peng, Y. Mao, B. L. Xu, and R. Huang, “Analysis of active components from San-Huang-Xie-Xin decoction,” Chinese Journal of Experimental Traditional Medical Formulae, vol. 19, no. 18, pp. 103–108, 2013.

[10] H. C. Geng, Y. Xin, F. W. Ai, and Y. L. Ma, “Fingerprint of San-Huang-Xie-Xin decoction by HPLC,” Chinese Traditional and Herbal Drugs, vol. 39, no. 4, pp. 524–529, 2008.
[11] Z. J. Chen, W. Li, M. S. Lin et al., “Research status of San-Huang-Xie-Xin decoction and analyses of key technologies and core issues,” Chinese Traditional and Herbal Drugs, vol. 47, no. 22, pp. 4111–4115, 2016.

[12] X. L. Zhou, H. Zhang, X. L. Meng, and B. Y. Hu, “Effects of San-Huang-Xie-Xin decoction on inflammatory cytokines release from RAW 264.7 cells induced by LPS,” Sichuan Journal of Physiological Sciences, vol. 33, no. 4, pp. 145–147, 2011.

[13] Q. J. Liu, X. D. Tian, Z. Q. Wu, P. Sun, and Y. L. Li, “Effect of San-Huang-Xie-Xin decoction on the proliferation of gastric cancer cells,” Progress in Modern Biomedicine, vol. 17, no. 31, pp. 6028–6032, 2017.

[14] Y. K. Yao, “The experiences of Shiyan Wu for treating schizophrenia in clinical,” Zhejiang Journal of Traditional Chinese Medicine, vol. 28, no. 4, p. 177, 1993.

[15] Y. Zhang, W. Chen, and X. L. Meng, “The protective effect of Sanhuangxixin decoction against oxidative stress and inflammatory injury of global cerebral ischemia reperfusion rats,” Pharmacy and Clinics of Chinese Materia Medica, vol. 30, no. 4, pp. 1–5, 2014.

[16] H. I. Choi, H. W. Lee, T. M. Eom et al., “A traditional Korean multiple herbal formulae (Yuk-Mi-Jilhwang-Tang) attenuates acute stress-induced brain tissue oxidation,” Drug and Chemical Toxicology, vol. 40, no. 2, pp. 125–133, 2017.

[17] B. H. Lee, J. Kim, R. M. Lee et al., “Gintonin enhances performance of mice in rotarod test: involvement of lysophosphatic acid receptors and catecholamine release,” Neurosciences, vol. 612, pp. 256–260, 2016.

[18] T. M. Wu, J. S. Chen, W. J. Xue, and X. L. Gao, “NMR metabonomics study on serum samples of young and middle-age hypertension patients with phlegm dampness retention syndrome,” Chinese Journal of Information on Traditional Chinese Medicine, vol. 23, no. 10, pp. 21–25, 2015.

[19] J. L. Chen, B. Y. Shi, H. Xiang et al., “1H NMR-based metabolic profiling of liver in chronic unpredictable mild stress rats with genipin treatment,” Journal of Pharmaceutical and Biomedical Analysis, vol. 115, no. 10, pp. 150–158, 2015.

[20] Z. Y. Li, L. L. Ding, J. M. Li et al., “1H-NMR and MS based metabolomics study of the intervention effect of curcumin on hyperlipidemia mice induced by high-fat diet,” PLoS One, vol. 10, no. 3, article e0120950, 2015.

[21] X. Song, J. Wang, P. Wang, N. Tian, M. Yang, and L. Kong, “1H NMR-based metabolomics approach to evaluate the effect of Xue-Fu-Zhu-Yu decoction on hyperlipidemia rats induced by high-fat diet,” Journal of Pharmaceutical and Biomedical Analysis, vol. 78-79, no. 5, pp. 202–210, 2013.

[22] Z. Zheng, S. Zhang, M. Yu et al., “An 1H NMR and UPLC-MS-based plasma metabonomic study to investigate the biochemical changes in chronic unpredictable mild stress model of depression,” Metabolomics, vol. 7, no. 3, pp. 413–423, 2011.

[23] H. Wu, P. Wang, M. Liu et al., “A 1H NMR-based metabonomic study on the anti-depressive effect of the total alkaloid of Corydalis Rhizoma,” Molecules, vol. 20, no. 6, pp. 10047–10064, 2015.

[24] B. Shi, J. Tian, H. Xiang et al., “A 1H-NMR plasma metabonomic study of acute and chronic stress models of depression in rats,” Behavioural Brain Research, vol. 241, no. 3, pp. 86–91, 2013.

[25] R. Laiatlussifa, A. Husni, and A. E. Nugroho, “Anti-stress activity of Sargassum polycystum extracts using a cold restraint stress model,” Food Science and Biotechnology, vol. 25, no. 2, pp. 589–594, 2016.

[26] C. S. Kim, Y. J. Jo, S. H. Park et al., “Anti-stress effects of ginsenoside Rs-G2-standardized ginseng extract in restraint stressed animals,” Biomolecules & Therapeutics, vol. 18, no. 2, pp. 219–225, 2010.

[27] W. Y. Wu, J. J. Hou, H. L. Long, W. Z. Yang, J. Liang, and D. A. Guo, “TCM-based new drug discovery and development in China,” Chinese Journal of Natural Medicines, vol. 12, no. 4, pp. 241–250, 2014.

[28] W. Peng, H. Shen, B. Lin et al., “Docking study and antiosteoporosis effects of a dibenzylbutylamine isolated from Lithsea cubeba targeting Cathepsin K and MEK1,” Medicinal Chemistry Research, vol. 27, no. 9, pp. 2062–2070, 2018.

[29] Y. T. Wang, Q. R. Tan, L. L. Sun et al., “Possible therapeutic effect of a traditional Chinese medicine, Sinisan, on chronic restraint stress related disorders,” Neuroscience Letters, vol. 449, no. 3, pp. 215–219, 2009.

[30] M. Moretti, J. Budni, D. B. dos Santos et al., “Protective effects of ascorbic acid on behavior and oxidative status of restraint-stressed mice,” Journal of Molecular Neuroscience, vol. 49, no. 1, pp. 68–79, 2013.

[31] Y. Cai, Y. F. Li, L. P. Tang et al., “A new mechanism of vitamin C effects on A/PR/1/47(H1N1) virus-induced pneumonia in restraint-stressed mice,” BioMed Research International, vol. 2015, Article ID 675149, 12 pages, 2015.

[32] A. Savini, A. Rossi, C. Pierro, L. Avigliano, and M. V. Catani, “SVCT1 and SVCT2: key proteins for vitamin C uptake,” Amino Acids, vol. 34, no. 3, pp. 347–355, 2008.

[33] A. Corti, A. F. Casini, and A. Pompella, “Cellular pathways for transport and efflux of ascorbate and dehydroascorbate,” Archives of Biochemistry and Biophysics, vol. 500, no. 2, pp. 107–115, 2010.

[34] B. Lee, B. Sur, J. Park et al., “Chronic administration of baicalin decreases depression-like behavior induced by repeated restraint stress in rats,” The Korean Journal of Physiology & Pharmacology, vol. 17, no. 5, pp. 393–403, 2013.

[35] S. H. Park, S. Jang, E. Son et al., “Polygonum aviculare L. extract reduces fatigue by inhibiting neuroinflammation in restraint-stressed mice,” Phytomedicine, vol. 42, no. 4, pp. 180–189, 2018.

[36] S. H. Park, S. Jang, S. W. Lee, S. D. Park, Y. Y. Sung, and H. K. Kim, “Akebia quinata Decaisne aqueous extract acts as a novel anti-fatigue agent in mice exposed to chronic restraint stress,” Journal of Ethnopharmacology, vol. 222, no. 8, pp. 270–279, 2018.

[37] Z. Zhang, X. Ma, Z. Xia et al., “NLRP3 inflammasome activation mediates fatigue-like behaviors in mice via neuroinflammation,” Neuroscience, vol. 358, no. 9, pp. 115–123, 2017.

[38] J. D. Johnson, J. Campisi, C. M. Sharkey et al., “Catecholamines mediate stress-induced increases in peripheral and central inflammatory cytokines,” Neuroscience, vol. 135, no. 4, pp. 1295–1307, 2005.

[39] R. L. Torres, I. S. Torres, G. D. Gamaro et al., “Lipid peroxidation and total radical-trapping potential of the lungs of rats submitted to chronic and subchronic stress,” Brazilian Journal of Medical and Biological Research, vol. 37, no. 2, pp. 185–192, 2004.

[40] A. Zaifer and N. Banu, “Antioxidant potential of flavoxetine in comparison to Curcuma longa in restraint-stressed rats,” Oxidative Medicine and Cellular Longevity
European Journal of Pharmacology, vol. 572, no. 1, pp. 23–31, 2007.

[41] W. Peng, X. Q. Qiu, Z. H. Shu et al., "Hepatoprotective activity of total iridoid glycosides isolated from Paederia scandens (lour.) Merr. var. tomentosa," Journal of Ethnopharmacology, vol. 174, no. 11, pp. 317–321, 2015.

[42] S. Kim, D.-S. Oh, J. Oh, T. Son, D. Yuk, and Y.-S. Jung, "Silymarin prevents restraint stress-induced acute liver injury by ameliorating oxidative stress and reducing inflammatory response," Molecules, vol. 21, no. 4, p. 443, 2016.

[43] L. Wu, X. H. Liu, T. H. Wang et al., "Protective effects of histone deacetylase inhibitor on stress-induced myocardial injury in rats," Chinese Journal of Applied Physiology, vol. 31, no. 3, pp. 193–196, 2015.

[44] H. Zardooz, S. Zahediasl, M. Gharibnaseri, and M. Hedayati, "Effect of chronic restraint stress on carbohydrate metabolism in rat," Physiology & Behavior, vol. 89, no. 3, pp. 373–378, 2006.