Effects of a combination of puerarin, baicalin and berberine on the expression of proliferator-activated receptor-γ and insulin receptor in a rat model of nonalcoholic fatty liver disease

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Abstract. Nonalcoholic fatty liver disease (NAFLD) is a prevalent disease, with a clinical spectrum ranging from simple fatty liver disease to nonalcoholic steatohepatitis and cirrhosis. Puerarin, baicalin and berberine are herbal products widely used in Asia, which are believed to possess therapeutic benefits for alleviating the symptoms of NAFLD. In the present study, a rat model of NAFLD, induced by a high-fat diet, was established and orthographical experimentation was used to investigate the effects of various combinations of puerarin, baicalin and berberine on the hepatic expression of proliferator-activated receptor (PPAR)-γ and insulin receptor (IR). The present study demonstrated that serum levels of total cholesterol, alanine transaminase and low-density lipoproteins were significantly decreased in the puerarin-dominated groups (P<0.05 vs. the model group), whereas the concentrations of tumor necrosis factor-α and interleukin-6 were significantly improved in the baicalin- and berberine-dominated groups (P<0.05 vs. the model group). Furthermore, as compared with the control group, the levels of PPAR-γ/IR mRNA and protein expression were significantly increased in the puerarin- and rosiglitazone-treated groups (P<0.01), and significantly decreased in the baicalin- and berberine-dominated groups (P<0.01). In conclusion, a combination of puerarin, baicalin and berberine induced favorable effects on NAFLD by upregulating hepatic PPAR-γ and IR expression levels, and different proportions of monomer compositions exerted variable positive effects on various stages of NAFLD.

Introduction

Nonalcoholic fatty liver disease (NAFLD) is characterized by an accumulation of lipids within hepatocytes, and is considered the hepatic manifestation of the metabolic syndrome (1). NAFLD develops progressively from simple steatosis to nonalcoholic steatohepatitis (NASH), fibrosis and cirrhosis (2,3). Previous studies have suggested a ‘two-hit’ theory for the pathogenesis of NAFLD, including insulin resistance, oxidative stress, lipid peroxidation and mitochondrial dysfunction (4,5), with insulin resistance considered the major factor.

Proliferator-activated receptor (PPAR)-γ is a member of the nuclear hormone receptor family and as such is activated by its ligand (6). Previous studies have demonstrated that this transcription factor is associated with insulin resistance, lipid metabolism and the regulation of inflammation in fatty liver disease (7-9). The insulin receptor (IR) is a well-studied receptor tyrosine kinase, which is associated with the regulation of peripheral glucose metabolism (10,11); therefore, a defect in IR may directly affect the role of insulin in the body, leading to insulin resistance and abnormal glucose tolerance (12). Previous studies have demonstrated that PPAR-γ and IR are closely associated with insulin resistance (13,14), and PPAR-γ agonists, such as thiazolidinediones (TZDs), are able to increase insulin sensitivity in adipose, liver and skeletal muscle tissue (15). However, TZDs possess numerous side effects, including significant weight gain and peripheral edema (16,17); therefore, it is necessary that novel agents that target PPAR-γ with reduced adverse effects be developed.

Previous studies have revealed the therapeutic effects of integrative medicine in treating patients with metabolic syndrome (18,19). Puerarin, baicalin and berberine (Fig. 1A-C), are three major constituents of the Chinese herbs: Puerariae Radix [roots of Pueraria lobata (Willd.) Ohwi, PR], Scutellariae Radix (roots of Scutellaria baicalensis Georgi SR) and Coptidis Rhizoma (rhizomes of Coptis chinensis Franch, CR), which have demonstrated therapeutic effects on rats models of NAFLD (20-22). It is believed that a combination of agents may effectively reduce side effects and improve adaptive resistance. In the present study, an orthogonal experimental design was used to investigate the various effects of puerarin, baicalin and berberine on rats with high-fat diet-induced NAFLD. The present
study aimed to elucidate the complex interactions of this multicomponent disease, and further clarify the potential for natural combination pharmacological therapies for patients with NAFLD.

Materials and methods

Experimental animals and modeling. A total of 96 adult male Sprague-Dawley rats weighing 150-180 g were obtained from the Animal Breeding Center of the Beijing Vital River Laboratories Company (Beijing, China). The rats were individually housed at 22±2°C with a relative humidity of 50±10% and a 12 h light/dark cycle. After 7 days of adaptive feeding, the rats were randomly divided into control (n=8), model (n=8), rosiglitazone treatment (n=8), and nine orthogonal experiment groups (n=8x9). The control group was fed a normal laboratory diet with free drinking water, whereas the model group, rosiglitazone treatment group (R) and orthogonal experiment groups (A-I) were fed a normal diet supplemented with 2% cholesterol and 10% lard, for 8 weeks. This study was approved by the Review Board of the Beijing University of Chinese Medicine (Beijing, China). All animal studies were conducted in accordance with the regulations and guidelines for the use and care of experimental animals of the Beijing University of Chinese Medicine (Beijing, China).

Therapeutic agents and reagents. Puerarin, baicalin, and berberine (Fig. 1) were purchased from Nanjing Zelang Medical Technology Co., Ltd. (Jiangsu, China). Rosiglitazone was purchased from Chengdu Hengrui Biotech Co., Ltd (Chengdu, China). Colorimetric kits for testing triglyceride (TG), total cholesterol (TC), high-density lipoprotein (HDL), low-density lipoprotein (LDL), alanine aminotransferase (ALT) and aspartate aminotransferase (AST) were purchased from BioSino Biotechnology and Science, Inc. (Beijing, China). ELISA kits for tumor necrosis factor (TNF)-α and interleukin (IL)-6 were purchased from Shanghai Yanji Biotechnology Co., Ltd. (Shanghai, China). Rabbit anti-PPARγ (ab27649, 1:2,000), mouse anti-IR (cat. no. ab69508, 1:5,000) and goat anti-GAPDH (ab9483, 1:20,000) antibodies were purchased from Abcam (Cambridge, UK). Additional reagents were obtained from Sigma-Aldrich (St. Louis, MO, USA).

Orthogonal experimental design and intervention. Based on single-factor experimental results, three independent parameters: Puerarin, baicalin and berberine doses (mg), were confirmed as the major influencing factors (23). A L9(34) orthogonal experimental design was conducted to evaluate the effects of the independent variables on the treatment of rats with NAFLD. The levels and ranges of the individual variables are outlined in Table I. The experiment design is presented in Table II, along with the experimental data. All experiments were performed in duplicate.

Measurement of serum biochemical parameters. Blood samples from the rats were obtained via abdominal aortic puncture and centrifuged at 644 x g for 10 min, in order to isolate the serum. Serum ALT, AST, TC, TG, LDL, and HDL concentrations were determined using a Vitros 250 automatic biochemical analyzer (Johnson & Johnson, Rochester, NY, USA).

Histological examination. All rats survived to the final analysis and rats were sacrificed by an intraperitoneal injection.

Table I. Factors and levels of orthogonal experimental design.

<table>
<thead>
<tr>
<th>Levels</th>
<th>Puerarin (mg)</th>
<th>Baicalin (mg)</th>
<th>Berberine (mg)</th>
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<tr>
<td>1</td>
<td>200</td>
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Table II. Orthogonal experimental design.

<table>
<thead>
<tr>
<th>Test</th>
<th>Puerarin</th>
<th>Baicalin</th>
<th>Berberine</th>
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Figure 1. Chemical structures of (A) puerarin, (B) baicalin, and (C) berberine.
of 10% chloral hydrate at the 8th week. In accordance with the frozen section technique, one fresh section of liver tissue from each rat was maintained in liquid nitrogen for 4-10 sec prior to staining with Oil Red O (Sigma-Aldrich). An additional liver tissue section was fixed by immersion in 10% buffered formalin for paraffin embedding, prior to staining with hematoxylin and eosin (Beijing CellChip Biotechnology Co. Ltd., Beijing, China). The samples were examined using an Olympus CK40 inverted microscope (Olympus, Tokyo, Japan).

Detection of TNF-α and IL-6 in the serum and liver tissue. The concentrations of TNF-α and IL-6 in the serum and liver tissue of the rats were analyzed using commercially available ELISA kits, according to the manufacturer's instructions. Briefly, the tissue was homogenized in an ice bath with ultrasonic irradiation. After 30 min, the homogenate was centrifuged for 15 min at 15,000 g at a temperature of 4°C. The supernatant was extracted in order to determine the concentration of protein. All samples were diluted to 1:10 and the absorbance was read at 450 nm using a microplate reader (Multiskan MK3; Thermo Fisher Scientific Inc., Rockford, IL, USA). Samples and standards were performed in triplicate.

Detection of the mRNA expression levels of PPAR-γ and IR by reverse transcription-quantitative polymerase chain reaction (RT-qPCR). Total RNA was isolated from the liver samples using TRIzol® reagent (Invitrogen Life Technologies, Carlsbad, CA, USA) and subsequently reverse-transcribed at 42°C for 60 min using the oligo(dT) primers and Super Script RT (Invitrogen Life Technologies). The primer sequences were as follows: PPAR-γ, forward 5'-TACACGGTTGATTCTTCT-3', reverse 5'-GCTCTACTTCTGATCGCATT-3'; IR, forward 5'-CTAAGCAGATGACGTTT-3', reverse 5'-GCTCCTTACACCATCATCAGC-3'; GAPDH, forward 5'-TGGAGTCTAGGCGCTCTT-3', reverse 5'-TGT CATATTTCCTGTTCCA-3'. All primers were synthesized by Shanghai Yingjun Biotechnology Company (Shanghai, China). Following cDNA synthesis, 3 µl cDNA was used in a 50-µl quantitative RT-qPCR reaction including 1X SYBR® Green Mix (Toyobo, Osaka, Japan) and 7.5 pmol of each specific primer. qPCR was conducted using a 7900HT Fast Real-Time PCR system (Applied Biosystems, CA, USA) with the following cycle conditions: 50°C for 2 min, 95°C for 10 min and then 40 cycles of 95°C for 5 min, 94°C for 15 sec, 60°C for 30 sec, 72°C for 30 sec and a final extension of 72°C for 5 min. The PCR products with the highest content were diluted by 1:1, 1:10, 1:100, 1:1,000, 1:10,000. PPAR-γ/IR mRNA expression levels were calculated using the ΔΔCt method (24) using Sequence Detection system 2.1 software (Eppendorf Co., Ltd., Hamburg, Germany). The relative PPAR-γ/IR mRNA expression levels were normalized to the expression of GAPDH in the samples. PCR amplifications were simultaneously performed in duplicate to avoid systematic errors.

Western blot analysis for the detection of PPAR-γ and IR protein expression. Proteins in the liver tissue homogenates were extracted using ice-cold tissue lysis buffer. Protein concentrations were determined using a bicinchoninic acid (BCA) protein assay kit (Promega, Madison, WI, USA). The liver tissue samples were homogenized in radioimmunoprecipitation assay lysis buffer (BioTeke, Co. Ltd., Beijing, China) containing a phenylmethylsulfonyl fluoride protease inhibitor (BioTeke, Co. Ltd.), and the total protein concentrations were subsequently determined using a BCA protein assay. Proteins (~50 µg) were loaded onto 5-10% SDS-PAGE gels and transferred to a polyvinylidene difluoride membrane (Bio-Rad Laboratories Inc., Hercules, CA, USA) for 70 min at 100 V. The membranes were then incubated in blocking buffer for 2 h prior to the addition of primary antibodies, including PPAR-γ and IR, which were then incubated at 4°C overnight. The membranes were immunoblotted with primary antibodies that recognized PPARγ (dilution, 1:2,000), IR (dilution, 1:5,000) and GAPDH (dilution, 1:5,000). The secondary antibodies [goat anti-rabbit immunoglobulin G (IgG; cat.no. 554020) and goat anti mouse IgG (cat.no. 551011) secondary antibodies; BD Biosciences, Franklin Lakes, NJ, USA] were diluted to 1:1,000 and incubated at room temperature for 2 h. Proteins were detected via enhanced chemiluminescence, and the intensity of protein bands was quantified using ImageJ version 1.37 software (National Institutes of Health, Bethesda, MD, USA).

Statistical analysis. All data were analyzed using SPSS software version 17.0 (SPSS, Inc., Chicago, IL, USA). Comparisons between the groups were analyzed for statistical significance using one-way analysis of variance and Wilcoxon's signed-rank test. P<0.05 was considered to indicate a statistically significant difference.

Results

Histological changes in the liver. Typical steatosis, balloon degeneration in hepatocytes and evident infiltration by inflammatory cells in the intercellular substance, were observed in the model group liver tissue samples (Fig. 2K), as compared with that of the control group (Fig. 2L). However, these changes in pathology were alleviated in the orthogonal experiment and rosiglitazone treatment groups (Figs. 2A-I and J, respectively). Furthermore, Oil Red O staining was used to detect the quantities of lipids in the hepatocytes, the results of which demonstrated that hepatocyte lipidosis was significantly increased in the model group (Fig. 3K), however, the deposition of lipid droplets in the hepatocytes was markedly reduced in the orthogonal experiment groups and rosiglitazone treatment group (Figs. 3A-I and J, respectively).

Effects of puerarin, baicalin and berberine on the biochemical parameters

Plasma HDL, LDL, TG and TC levels. Compared with the control group, the serum HDL levels in the model group were significantly decreased (P<0.01); whereas the levels of LDL, TG, and TC were significantly increased (P<0.01). However, no significant differences in the HDL levels of rats were determined between the orthogonal experiment groups (P>0.05, Fig. 4C), and the model group. The levels of TG in groups A, D, F, G and R were significantly decreased (P<0.05, Fig. 4A), as were the TC levels in groups A-C and E (P<0.05, Fig. 4B), and the levels of LDL in groups A-D and R (P<0.05, Fig. 4D), as compared with the model group.
Plasma ALT and AST levels. The serum levels of ALT and AST were significantly increased (P<0.01) in the model group, as compared with the control group. Furthermore, in groups F, G, H, I and R the serum levels of AST were significantly decreased (P<0.01, Fig. 5A), as were the levels of ALT in groups A, B, D, E, I and R (P<0.01, Fig. 5B), as compared with the model group.

Effects of puerarin, baicalin, and berberine on the levels of TNF-α and IL-6. Compared with the control group, the levels
of TNF-α and IL-6 in the serum and liver samples of the model group were significantly increased (P<0.05); whereas, compared with the model group, the serum levels of TNF-α in groups B, D, G and H were significantly decreased (P<0.05, Fig. 5C), and the serum levels of IL-6 were significantly decreased in groups A, B, D and F-H (P<0.05, Fig. 5D). Furthermore, in groups A, D, G, H and I, the levels of TNF-α in the liver were significantly decreased (P<0.05, Fig. 5E), and IL-6 levels in the liver were decreased in groups A, D, F and H (P<0.05, Fig. 5F), as compared with the model group.

mRNA expression levels of PPAR-γ and IR. In the model group, the mRNA expression levels of PPAR-γ and IR were significantly decreased (P<0.01), as compared with the control group. Furthermore, in groups A, D, G and R the mRNA expression levels of PPAR-γ were significantly increased (P<0.01, Fig. 6A), as were the IR mRNA expression levels in groups A, B, D, F and R (P<0.01, Fig. 6B), as compared with the model group. However, no significant changes were determined between the rosiglitazone treatment group and the orthogonal experiment groups (P>0.05).

PPAR-γ and IR protein expression levels, as determined by western blotting. Compared with the control group, the protein expression levels of PPAR-γ and IR in the model group were significantly decreased (P<0.01). Furthermore, the PPAR-γ protein expression levels in groups A, B, D, G, H and R were significantly increased, as compared with the model group (P<0.01, Fig. 7A); and in groups A-D, F and R the protein expression levels of IR were significantly increased (P<0.01, Fig. 7B). However, no differences were determined between the rosiglitazone treatment and orthogonal experiment groups (P>0.05).

Discussion

NAFLD, which has a clinical spectrum ranging from simple fatty liver disease to NASH and cirrhosis (25), is currently the most common type of liver disease (26). Although the etiology of NAFLD has yet to be fully elucidated, the ‘two-hit’ theory has been accepted as the most popular mechanism (27). The ‘first hit’ is associated with hepatic TG accumulation and insulin resistance, whereas the ‘second hit’ involves oxidative stress and the induction of inflammatory cytokines (28). PPAR-γ is a sequence-specific and ligand-dependent nuclear transcription factor. With its four isoforms: γ1, γ2, γ3 and γ4, PPAR-γ is associated with the control of lipid storage and the differentiation of adipocytes and macrophages (29). TZDs are high affinity synthetic PPAR-γ ligands that can stimulate adipocyte differentiation and improve insulin resistance via PPAR-γ activation (30). One TZD in particular, rosiglitazone, has been established as a widely used therapeutic agent in the treatment of insulin resistance. Rosiglitazone has previously been demonstrated to improve sensitivity to insulin as well as transaminases and liver histology (31). In the present study, rosiglitazone was used as a positive pharmacological control, in order to compare the effects of puerarin, baicalin, and berberine on rats with high-fat diet-induced NAFLD. IR was initially identified as a homodimer, with extrinsic disulfide bonds that generate the functional receptor. Each monomer of IR is composed of one α
The lack of IR may directly affect the distribution of insulin in the body, causing insulin resistance and abnormal glucose tolerance. Previous studies have demonstrated that PPAR-γ and IR may be important for the ‘first hit’ stage of NAFLD, since they are associated with insulin resistance (34,35). Increased
therapy through a small number of experiments that determine the best therapeutic effect, the orthogonal experiment was applied to the overall design, comprehensive comparison and statistical analysis of the present study.

In the present study, rats with high-fat diet-induced NAFLD were employed to evaluate the efficacies of puerarin, baicalin and berberine against NAFLD. The high-fat diet model has been widely used in previous NAFLD studies (39,40). The histological results of the present study observed NAFLD features in the model rats, indicating that an NAFLD model was successfully generated. Furthermore, as expected, the rosiglitazone treatment and orthogonal experiment groups significantly attenuated hepatic steatosis, inflammation and fibrosis in the rats.

The pharmaceutical compositions examined in the present study demonstrated that a combination of puerarin, baicalin and berberine may exert promising lipid-lowering, hepatoprotective and anti-inflammatory activities. The various orthogonal experiment groups had numerous effects on biochemical parameters and pro-inflammatory cytokines. The present study demonstrated that serum levels of TC, LDL and ALT were significantly reduced in the puerarin-dominated groups, whereas serum levels of TNF-α and IL-6 were improved in the baicalin- and berberine-dominated groups. Therefore, the results of the present study suggested that puerarin may preferentially affect lipid metabolism, whereas baicalin and berberine may impact the inflammatory response. Furthermore, PPAR-γ/IR protein and mRNA expression levels were also reduced in the model group, thus suggesting that insulin resistance had developed in response to decreased PPAR-γ and IR in the NAFLD model rats. Notably, the rosiglitazone treatment group and certain orthogonal experiment groups improved insulin resistance; however, no obvious trends in the various combinations of puerarin, baicalin and berberine were determined. In addition, no differences were demonstrated between the rosiglitazone treatment and orthogonal experiment groups. Therefore, we hypothesize that a combination of puerarin, baicalin and berberine may be used in the early pathological stage (‘first hit’) of NAFLD to improve insulin resistance via PPAR-γ and IR upregulation. Since puerarin treatment is associated with the regulation of lipid metabolism, puerarin may be used in the early stages of simple fatty liver disease; whereas baicalin and berberine may be used in NASH, as they may alleviate the inflammatory response. However, it was not possible to clarify the effects on liver cirrhosis in the present study due to a lack of detection of liver fibrosis.

In conclusion, the combination of puerarin, baicalin and berberine induced favorable therapeutic effects on rats with high-fat diet-induced NAFLD. Furthermore, variably positive effects were demonstrated in the various stages of NAFLD following treatment with numerous proportions of monomer compositions. This suggests that a therapeutic combination of puerarin, baicalin and berberine may regulate lipid metabolism by upregulating the expression of hepatic PPAR-γ and IR, leading to improvements in insulin resistance, which may be useful in the prevention and treatment of NAFLD, especially simple fatty liver disease and NASH. Further studies focusing on the effects of combination treatment with puerarin, baicalin and berberine on liver fibrosis are required.
Acknowledgements

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