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# The Chinese herbal medicine FTZ attenuates insulin resistance via IRS1 and PI3K in vitro and in rats with metabolic syndrome

Xuguang Hu<sup>†</sup>, Man Wang<sup>†</sup>, Weijian Bei, Zongyu Han and Jiao Guo<sup>\*</sup>

## Abstract

**Background:** Insulin resistance plays an important role in the development of metabolic syndrome (MS). Fu Fang Zhen Zhu Tiao Zhi formula (FTZ), a Chinese medicinal decoction, has been used to relieve hyperlipidemia, atherosclerosis and other symptoms associated with metabolic disorders in the clinic.

**Methods:** To evaluate the effect of FTZ on insulin resistance, HepG2 cells were induced with high insulin as a model of insulin resistance and treated with FTZ at one of three dosages. Next, the levels of glucose content, insulin receptor substrate1 (IRS1) protein expression and phosphatidylinositol 3-kinase (PI3K) subunit p85 mRNA expression were measured. Alternatively, MS was induced in rats via gavage feeding of a high-fat diet for four consecutive weeks followed by administration of FTZ for eight consecutive weeks. Body weight and the plasma levels of lipids, insulin and glucose were evaluated. Finally, the expression of PI3K p85 mRNA in adipose tissue of rats was measured.

**Results:** Our results revealed that FTZ attenuated glucose content and up-regulated the expression of PI3K p85 mRNA and IRS1 protein in insulin-resistant HepG2 cells in vitro. Moreover, FTZ reduced body weight and the plasma concentrations of triacylglycerol, cholesterol, fasting glucose and insulin in insulin resistant MS rats. FTZ also elevated the expression of PI3K p85 mRNA in the adipose tissues of MS rats.

**Conclusion:** FTZ attenuated MS symptoms by decreasing the plasma levels of glucose and lipids. The underlying mechanism was attenuation of the reduced expression of PI3K p85 mRNA and IRS1 protein in both insulin-resistant HepG<sub>2</sub> cells and MS rats.

**Keywords:** Metabolic syndrome, Insulin resistance, Fu Fang Zhen Zhu Tiao Zhi formula (FTZ)

## Background

Reaven noted that dyslipidemia, hypertension and hypertriglyceridemia often occurred together [1]. Consequently, this author proposed the concept of X Syndrome, regarding insulin resistance as the primary characteristic of X Syndrome. Based on the association of X syndrome with a variety of metabolic diseases, Zimmet et al. defined its symptoms as metabolic syndrome (MS). MS is related to excessive energy intake, a sedentary lifestyle, and

multiple genetic factors [2]. Many studies have demonstrated that patients with MS displayed an increased risk of developing diabetes, cardiovascular disease and other diseases. Insulin resistance (IR) has been suggested as the common pathophysiological basis of MS. Because there is no eutherapeutic Western medicine to treat MS, searching for an effective therapeutic among the traditional Chinese medicines (TCMs) or botanicals is urgently required.

Fu Fang Zhen Zhu Tiao Zhi formula (FTZ), for which a patent has been issued, is composed of eight medical herbs and has been proven clinically effective for the treatment of dyslipidemia [3,4]. Our previous study demonstrated that FTZ could reduce serum levels of total cholesterol (TC), triglycerides (TG) and low-density lipoprotein cholesterol (LDL-C) and increase the

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serum level of high-density lipoprotein cholesterol (HDL-C) in hyperlipidemic patients [5,6]. FTZ could reduce the viscosity of plasma and whole blood in hyperlipidemic patients [7]. In addition, our previous studies also demonstrated the antioxidant properties of FTZ, which reduced the oxidation of LDL-C [8]. FTZ also could improve serum lipid profile in high lipid diet induced hyperlipidemic rats by regulating the HMG-CoA reductase and CYP7A1 [9].

MS includes the symptoms dyslipidemia, hypertension and hypertriglyceridemia and is typically characterized by insulin resistance. Therefore, insulin resistance is suggested to be a new drug target for treatment of MS or dyslipidemia. Nevertheless, the effect and mechanisms of FTZ on MS and insulin resistance remain unclear.

To elucidate the effect of FTZ on insulin resistance and its mechanism of action, HepG2 cells with insulin resistance and MS rats were used, and the effect of FTZ on insulin-resistant HepG2 cells and MS rats was measured. Moreover, the resistance index, the gene expression level of PI3K and the protein expression level of IRS1 were also investigated.

## Methods

### Materials

HepG2 cells were purchased from the American Type Culture Collection (ATCC) bioresource center (Manassas, VA, USA). Dulbecco's modified Eagle's medium (DMEM) and fetal bovine serum (FBS) were purchased from Invitrogen (Carlsbad, CA, USA). IRS1 and GAPDH antibodies were from Abcam Inc. (Cambridge, MA, USA). Insulin and rosiglitazone (RGS) were purchased from Sigma (St. Louis, MO, USA). All other reagents were analytical grade.

### Preparation of FTZ extract

Herbs in FTZ [composed of *Ligustrum lucidum* W.T. Aiton, fructus; *Atractylodes macrocephala* Koidz., rhizoma; *Salvia miltiorhiza* Bunge, radix; *Coptis chinensis* Franch., rhizoma; *Panax notoginseng* (Burk.) F.H. Chen, radix; *Eucommia ulmoides* Oliv., cortex; *Cirsium japonicum* (Thunb.) Fisch. ex DC., radix; *Citrus medica* var. *sarcodactylus* (Siebold ex Hoola van Nooten) Swingle, fructus] were provided by Zhixin Chinese Herbal Medicine Co., Ltd. (Guangzhou, China) and authenticated by Professor Shuyan Li, pharmacognosist of School of Chinese Medicinal Sciences, Guangdong Pharmaceutical University. All of the raw materials in FTZ were examined according to the quality control criteria of Chinese Pharmacopoeia 2010 and were controlled as previously reported [10]. The FTZ extract was prepared via alcohol and water extraction of eight herbs according to the protocol (see Supporting Material for the protocol of FTZ preparation). FTZ was obtained from the Institute of Materia Medica, Guangdong Pharmaceutical University.

The voucher specimens were GDPZY 20110901-8 (see Supporting Material-Voucher specimen). Quality analysis of the FTZ extract was performed via HPLC fingerprinting, which was obtained using a HPLC unit (Waters, USA) with an Agilent HC-C18 column (4.6 mm × 250 mm, 5 μm). All assigned peaks were identified by performing a co-injection test with authentic samples and comparative analysis of the UV spectral data (see Supporting Material-HPLC conditions) [9].

### Cell culture

The human hepatocellular carcinoma cell line HepG2 was purchased from ATCC. Cells were cultured in DMEM supplemented with 10% heat-inactivated fetal FBS at 37°C in a 5% CO<sub>2</sub> atmosphere. In all experiments, cells grew to 80-90% confluence.

### Induction of insulin-resistance in HepG2 cells and glucose uptake experiments

Insulin resistance was induced in HepG2 cells as previously described [11-13]. In brief, HepG2 cells were seeded on 24-well plates at  $2 \times 10^5$  cells/well, incubated for 24 h to reach maximal confluence and serum-starved for another 24 h. The cells were then incubated for 36 h in serum-free DMEM containing 25 mmol/l d-glucose,  $10^{-6}$  mol/l insulin in the absence or presence of 1, 25 and 100 μg/ml FTZ or 10 μmol/l RGS. FTZ administration at 100, 25 and 1 μg/ml were defined as high, medium and low dosages, respectively. Next, cells were washed twice with PBS. The cells were then incubated for 24 h in serum-free DMEM without phenol red. The glucose content was quantified using a GOD-POD kit, measuring optical absorbance at 505 nm.

### Western blot analysis

Cells were washed with ice-cold PBS and lysed with lysis buffer (50 mmol/l Tris HCl, 1% Triton X-100, 0.5% sodium deoxycholate, 150 mmol/l NaCl, 1 mmol/l EDTA, 1 mmol/l PMSE, 1 mmol/l sodium orthovanadate, 1 mmol/l NaF and 0.2% protease inhibitor cocktail; pH7.2). For western blotting, protein samples (20 μg) of high insulin-induced insulin-resistant HepG2 cells were separated via 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE). The proteins were transferred to a PVDF membrane and incubated with primary antibody (anti-IRS1, or anti-GAPDH), followed by secondary antibody (horseradish peroxidase-conjugated anti-rabbit IgG). Intensity of the immunoblot signal was assayed using Western Bright™ ECL spray and analyzed quantitatively using GeneTools software from Syngene (Syngene, Cambridge, UK).

### RNA isolation and quantitative real-time PCR

Total RNA was isolated from HepG2 cells using TRIzol® reagent (Invitrogen, Carlsbad, CA, USA) dissolved in

DEPC-treated water according to the manufacturer's instructions. RNA (2 µg) was reverse-transcribed to cDNA using oligo(dT) primers and moloney murine leukemia virus reverse transcriptase (RT) in a final volume of 20 µl under the conditions recommended by the supplier (Invitrogen). The resulting cDNA was amplified using primers specific for PI3K or β-actin in a total volume of 10 µl. Quantitative real-time PCR was performed using a Roche LightCycler 480 PCR system using SYBR green fluorescence. The expression level of PI3K was normalized to that of β-actin, which was used as a specific endogenous control. Primer sequences used were as follows:

PI3K forward 5'-TGG ACG GCG AAG TAA AGC ATT-3', reverse 5'-AGT GTG ACA TTG AGG GAG TCG-3'; β-actin forward 5'-AGC CAT GTA CGT AGC CAT CC-3', reverse 5'- ACC CTC ATA GAT GGG CAC AG-3'.

#### **Induction of metabolic syndrome in rats and experiment protocol**

Male Sprague Dawley rats (180-220 g) were obtained from the Laboratory Animal Center of Guangzhou University of Chinese Medicine (Guangzhou, Republic of China). Rats were housed in a climate-controlled environment (25±1°C at 50% relative humidity) with 12-h light/12-h dark cycles. Water was available *ad libitum*. The model rats of metabolic syndrome (MS) were induced by fed a high-fat diet (HFD) consisting of 40% (wt/wt) fat for 12 consecutive weeks, while control rats were fed normal chow. Then the MS rats were randomly divided into 5 groups and treated with distilled water (Group 1), FTZ -H (Group 2), FTZ -M (Group 3), FTZ -L (Group 4) and aspirin (200 mg/kg) (Group 5), respectively. FTZ with 6 g/kg, 3 g/kg and 1.5 g/kg were defined as high, medium and low dosages, respectively. From the fifth week to twelfth week, rats of FTZ groups were treated by FTZ for 8 consecutive weeks. Then the rats were used to measure body weight (BW) and serum levels of total cholesterol (TC), triglycerides (TG) and high-density lipoprotein cholesterol (HDL-C). Subsequently, all the rats were sacrificed and their epididymis and kidney fat were excised for PI3K mRNA assay. The animal care and study protocols were maintained in accordance with the provisions and general recommendation of the Chinese Regulations for the Administration of Affairs Concerning Experimental Animals at Guangdong Pharmaceutical University.

#### **Determination of body weight and biochemical indexes**

During the animal experiments, body weight (BW) was recorded at 0, 4, 8 and 12 weeks. Serum levels of total cholesterol (TC), triglycerides (TG) and high-density

lipoprotein cholesterol (HDL-C) were analyzed using a blood chemistry analyzer (Hitachi 7040).

#### **Assessment of insulin resistance (IR)**

Fasting plasma glucose level was measured via glucose oxidase method. Fasting plasma insulin was measured using radioimmunoassay method. IR was assessed according to the homeostasis model assessment index (HOMA-IR) calculated using the following formula: (fasting plasma insulin level [µU/ml] × fasting plasma glucose [mmol/l]) / 22.5 [14].

#### **RNA isolation and RT-PCR**

Total RNA was isolated from adipose tissue using TRIzol® reagent (Invitrogen, Carlsbad, CA, USA) dissolved in DEPC-treated water according to the manufacturer's instructions. RNA (2 µg) was reverse transcribed to cDNA using oligo (dT) primers and moloney murine leukemia virus reverse transcriptase (RT) in a final volume of 20 µl under the conditions recommended by the supplier (Invitrogen). For polymerase chain reaction (PCR) amplification of PI3K and β-actin, 1 µl of cDNA template and the following specific primers were used: PI-3K p85 forward 5'-GAA GGC AAC GAG AAG GA-3', reverse 5'-CAC AAG TGT CAG CCA CAT-3' (213bp); β-actin forward 5'-AGA TCT GGC ACC ACA CCT TCT AC-3', reverse 5'-TCA GGA TCT TCA TGA GGT AGT CT-3' (388bp). The reaction cycle conditions were: denaturation at 90°C for 50 s, annealing at 56.1°C for 50 s and extension at 72°C for 50 s. The PCR products were resolved using a 2% agarose gel and visualized with ethidium bromide staining. The expression level of PI3K was normalized to that of β-actin, which was used as a specific endogenous control.

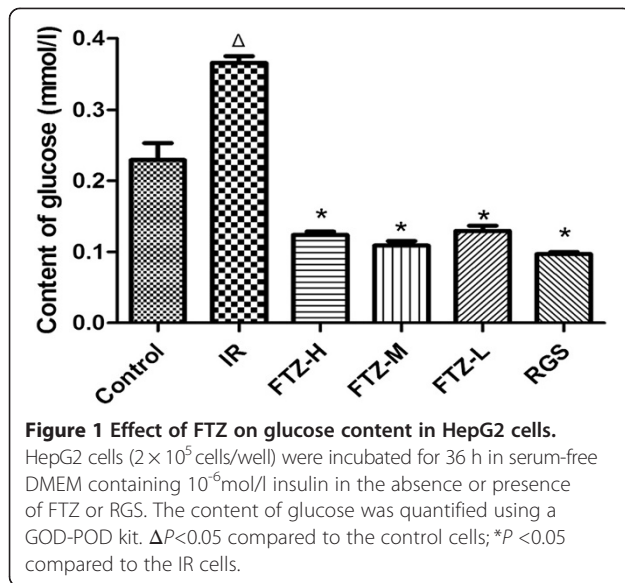
#### **Statistics**

Statistical analyses were conducted using SPSS16.0 software. All results are presented as the mean ± standard deviation (SD). Statistical analysis was performed via analysis of variance (one-way ANOVA) followed by the Student-Newman-Keuls test for significance. Differences were considered statistically significant at P < 0.05.

#### **Results**

##### **Effect of FTZ on glucose content in insulin-resistant HepG2 cells**

The glucose content in insulin-resistant HepG2 cells in culture medium significantly increased compared to that of control cells. After treatment with FTZ (1, 25 and 100 µg/ml), glucose content in the culture medium significantly decreased compared to that of IR cells (P < 0.05). RGS (10 µmol/l), used as a positive control drug, was also able to increase glucose content in the culture medium (Figure 1).

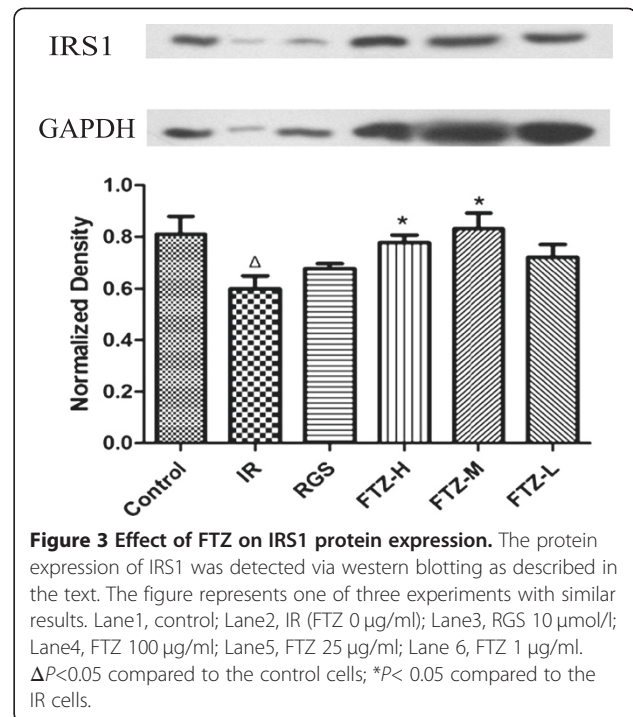
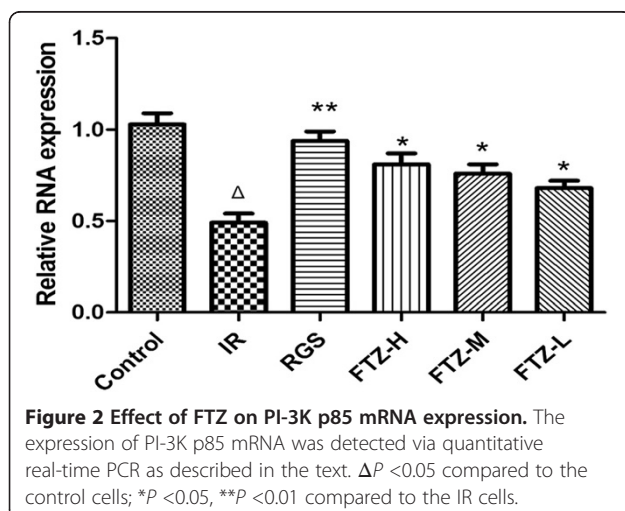


#### Effect of FTZ on PI-3K p85 mRNA expression in insulin-resistant HepG2 cells

To evaluate the effect of FTZ on PI-3K p85 mRNA expression, total RNA was extracted from HepG2 cells, and real-time PCR was performed. As shown in Figure 2, PI-3K p85 mRNA expression in HepG2 cells with IR was decreased compared to control cells ( $P < 0.05$  or  $P < 0.01$ ). After treatment with FTZ, PI-3K p85 mRNA expression significantly increased compared to IR cells ( $P < 0.05$ ). These results suggest that FTZ induces an insulin sensitizing effect on IR cells through the up-regulation of PI-3K p85 mRNA expression in HepG2 cells (Figure 2).

#### Effect of FTZ on IRS1 protein expression in HepG2 cells with insulin resistance

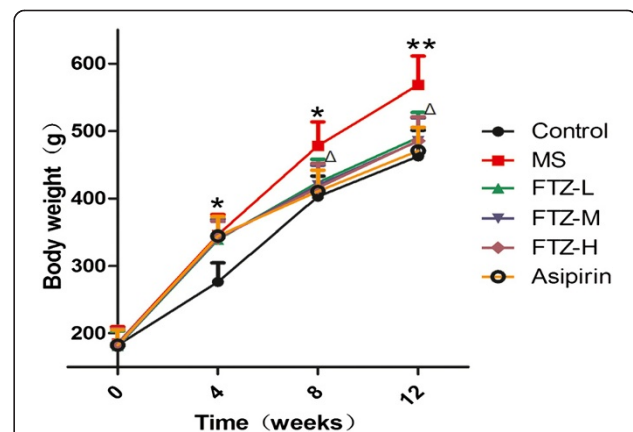
To elucidate the effects of FTZ, western blot analysis was used to measure IRS1 protein expression in HepG2

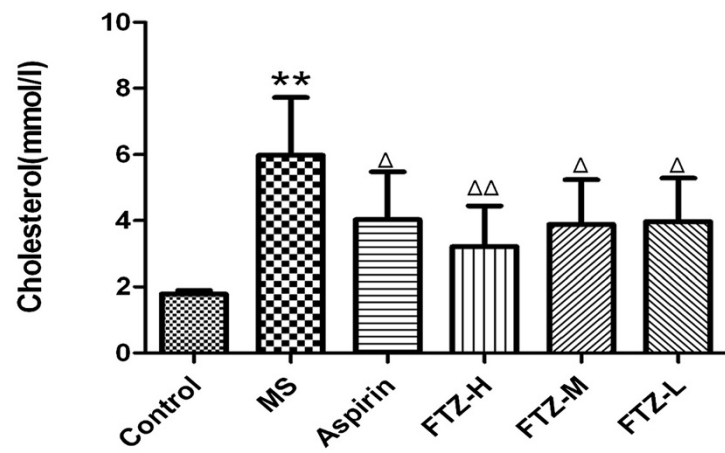


cells. As shown in Figure 3, IRS1 protein expression was significantly reduced compared to control cells ( $P < 0.05$ ). After treatment with FTZ, IRS1 protein expression was significantly increased compared to IR cells ( $P < 0.05$ ) (Figure 3).

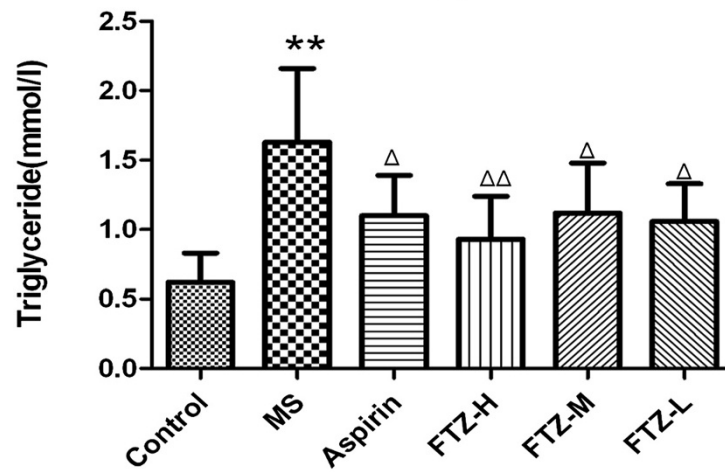
#### Effect of FTZ on body weight of MS rats

After the rats were fed a high-fat diet for 12 continuous weeks, our results indicated that the body weight of

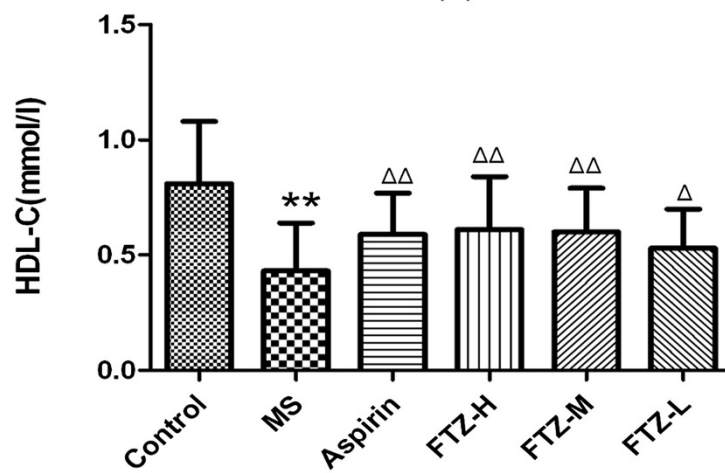




(a)



(b)



(c)

Figure 5 (See legend on next page.)

(See figure on previous page.)

**Figure 5 Blood biochemical indexes (cholesterol, triglyceride and HDL-C) of MS rats.** Blood samples were collected from the orbital sinus, and cholesterol (a), triglyceride (b), HDL-C (c) levels were measured following administration of FTZ at 6 g/kg/day, 3 g/kg/day or 1.5 g/kg/day, aspirin (AS) at 10 ml/kg/day or saline (control rats) for 8 consecutive weeks in high-fat diet-treated rats. \* $P < 0.05$ , \*\* $P < 0.01$  compared to the control rats;  $\Delta P < 0.05$ ,  $\Delta\Delta P < 0.01$  compared to the MS rats.

MS rats was significantly higher than that of control rats ( $P < 0.05$ ,  $P < 0.01$ ). After high, medium or low dosage FTZ treatment for 4 consecutive weeks (equivalent to the rats fed a high-fat diet for 8 weeks), the body weight of the rats in the FTZ groups was significantly reduced compared to that of MS rats. After FTZ treatment for 8 consecutive weeks (equivalent to the rats fed a high-fat diet for 12 weeks), the high, medium and low dosage FTZ-treated groups all possessed body weights which were significantly reduced compared to the MS rats ( $P < 0.05$ ,  $P < 0.01$ ) (Figure 4).

#### Effect of FTZ on blood biochemical index of MS rats

Compared to the control rats, the levels of serum TG and TC in MS rats were significantly increased ( $P < 0.05$  or  $P < 0.01$ ), while the level of serum HDL-C was significantly reduced ( $P < 0.05$  or  $P < 0.01$ ). After treatment with FTZ, the serum levels of TG and TC in MS rats were significantly reduced ( $P < 0.05$  or  $P < 0.01$ ), and the serum level of HDL-C was significantly increased ( $P < 0.05$  or  $P < 0.01$ ). These results revealed that FTZ could alter the serum levels of lipids in MS rats (Figure 5a-c).

#### Effect of FTZ on fasting glucose, insulin and HOMA-IR index in MS rats

Compared to the control rats, levels of fasting glucose and insulin, as well as the HOMA-IR index, were significantly higher in MS rats ( $P < 0.05$  or  $P < 0.01$ ). After treatment with the high dosage of FTZ or aspirin, the levels of fasting glucose and insulin and HOMA-IR index were all significantly reduced compared to the MS rats ( $P < 0.05$  or  $P < 0.01$ ). After treatment with the medium dosage of FTZ, the fasting glucose level and the HOMA-IR index were significantly reduced compared to the MS rats ( $P < 0.05$  or  $P < 0.01$ ). After treatment with the low dosage of FTZ, the HOMA-IR index was significantly reduced compared to the MS rats ( $P < 0.05$  or  $P < 0.01$ ) (Figure 6).

#### Effect of FTZ on PI-3K p85 mRNA expression in adipose tissue

To evaluate the effect of FTZ on PI-3K p85 mRNA expression, we performed RT-PCR in the adipose tissue of rats. As shown in Figure 7, compared to the control rats, the MS rats produced a lower expression level of PI-3K p85 mRNA ( $P < 0.05$  or  $P < 0.01$ ). Administration of either

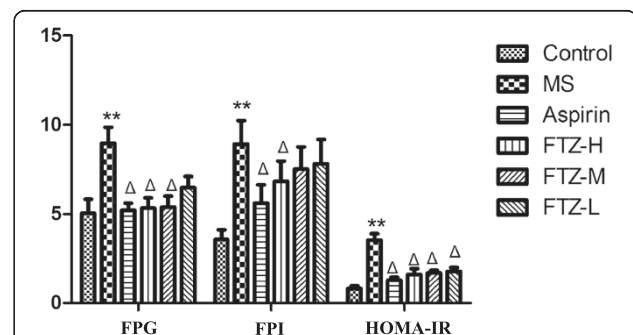
aspirin or FTZ, the expression of PI-3K p85 mRNA was increased compared to the MS rats (Figure 7).

#### Discussion

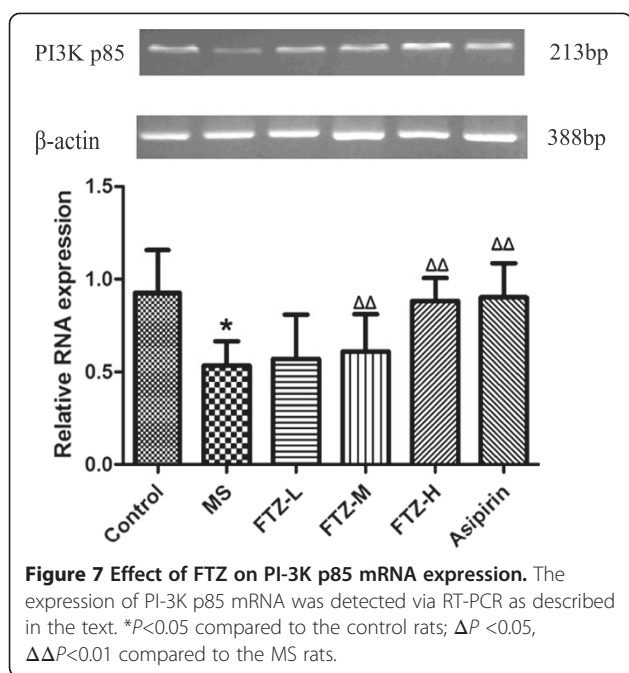
This study revealed that the Chinese herbal formula FTZ could attenuate MS symptoms by decreasing the content of glucose and lipids, and well as the insulin resistance index. Additionally, its effects were possibly mediated through increased expression of PI-3Kp85 mRNA and IRS1 protein in insulin-resistant HepG2 cells and MS rats. Insulin resistance has been suggested as an underlying cause of MS, including hyperglycemia, dyslipidemia and type 2 diabetes mellitus.

In our study, HepG2 cells were used as an insulin resistance model to investigate the effect of FTZ on glucose metabolism and insulin signaling. HepG2 cells express PI-3Kp85 and IRS1 genes, which are involved in the insulin signaling pathway [15,16]. Therefore, these cells have been widely used to analyze glucose metabolism, lipid metabolism, and insulin resistance [17,18].

Defects in the insulin signaling cascade, which lead to impaired glucose utilization, were believed to play a key role in the pathogenesis of insulin resistance [19]. It is conceivable that IRS-1 tyrosine phosphorylation in response to insulin stimulation generally increased the association of IRS-1 with PI 3-kinase, resulting in increased PI 3-kinase activity, which in turn led to activation of serine/threonine kinase protein B (PKB or Akt) and, ultimately, to an



**Figure 6 Other blood biochemical indexes (fasting glucose, insulin and HOMA-IR index) of MS rats.** Fasting plasma glucose (FPG) level was measured via the glucose oxidase method. Fasting plasma insulin (FPI) in rats was measured using a radioimmunoassay method. To quantify the insulin resistance index, the following formula was used:  $HOMA-IR = \{(FPG * FPI) / 22.5\}$ . \*\* $P < 0.01$  compared to the control rats;  $\Delta P < 0.05$  compared to the MS rats.



enhancement in insulin-stimulated glucose disposal [20]. Our research results revealed that the insulin receptor was impaired, producing an insulin-resistant state in HepG2 cells under high insulin conditions. The expression of the IRS-1 protein and IRS-1-associated PI-3K activity in HepG2 cells were significantly decreased. After treatment with FTZ, the expression of IRS-1 protein and PI-3K mRNA were partially restored. Here, we revealed that the FTZ-mediated recovery of insulin action was related to the improvement of the IRS-1/PI 3-kinase signaling pathway in insulin-resistant HepG2 cells. It appears that a FTZ-mediated improvement in post-receptor insulin signaling may have induced the subsequent increase in insulin sensitivity.

In our study, MS model rats were induced via high-fat diet feeding for 4 weeks. This model exhibited hyperinsulinemia, obesity, decreased insulin sensitivity, dyslipidemia and other features [21]. In our study, the MS rats exhibited increased body weight, levels of serum TG and total cholesterol, fasting glucose and plasma insulin, as well as an increased insulin resistance index. This was consistent with previous studies, such as I-Min Liu et al. [22]. After treatment with FTZ, body weight, levels of serum TG and TC, fasting glucose and plasma insulin and the insulin resistance index were significantly reduced compared to MS rats. FTZ treatment also enhanced the activity of PI3K in adipose tissue compared to MS rats.

Our study suggested that FTZ might ameliorate insulin resistance and treat MS. This effect might be associated with the compounds which it contained. It has

been reported that oleanolic acid (OA) in *Ligustrum lucidum* W.T. Aiton decreased serum triglyceride, total cholesterol, LDL and free fatty acids, increased serum HDL and reduced hepatic lipid accumulation. Furthermore, inflammation in db/db mice was improved by OA, as evidenced by decreased levels of IL-1  $\beta$ , IL-6, and TNF- $\alpha$  in the circulation and in the liver. These results suggested that OA improved hepatic insulin resistance via inhibition of mitochondrial ROS, hypolipidemia and anti-inflammatory effects [23]. Ginsenoside Re in *Panax notoginseng* (Burk.) F.H. Chen reduced insulin resistance via activation of the PPAR- $\gamma$  pathway by directly increasing the expression of PPAR- $\gamma$ 2 and its responsive genes, adiponectin, IRS-1 and ap2, inhibiting TNF- $\alpha$  production and facilitating the translocation of GLUT4 to promote glucose uptake and disposal in 3T3-L1 adipocytes [24]. Berberine in *Coptis chinensis* Franch. improved insulin-induced tyrosine phosphorylation of IRS-1 and the recruitment of p85 to IRS-1. The ameliorated insulin signal transduction was related to berberine-mediated inhibition of mTOR, which attenuated serine phosphorylation of IRS-1. These results suggested that berberine might ameliorate insulin resistance by modulating key molecules in the insulin signaling pathway, leading to increased glucose uptake in insulin-resistant cells [25]. Therefore, we suspect that these ingredients might explain the role of FTZ in ameliorating insulin resistance.

## Conclusion

In conclusion, our study indicated that FTZ could decrease serum triglyceride, total cholesterol and fasting blood glucose and increase serum HDL-C, thereby reactivating the insulin-stimulated IRS1/PI3K pathway in insulin-resistant HepG2 cells and up-regulating PI3K expression in adipose tissue. Therefore, the beneficial effects of FTZ on insulin resistance suggest that this decoction may be a promising therapeutic for MS and insulin resistance.

## Abbreviations

FTZ: Fu Fang Zhen Zhu Tiao Zhi formula; MS: Metabolic syndrome; IR: Insulin resistance; IRS1: Insulin receptor substrate-1; PI3K: Phosphatidylinositol 3-kinase; TG: Triglyceride; TC: Total cholesterol; HDL-C: HDL-cholesterol; FPG: Fasting plasma glucose; FPI: Fasting plasma insulin; HOMA-IR: Homeostasis model assessment- insulin resistance index.

## Competing interests

The author(s) declare that they have no competing interests.

## Authors' contributions

Dr. J.Guo and Xuguang Hu designed the study. Man Wang carried out experiments. Bei WJ and Wang LY, participated in the design of study, interpretation of results, and drafted the manuscript. Mr. Shuyan Li, Zongyu Han, Xiuteng Zhou, Le Cao, Hu Yinming, Ms. Wei He, Junhui Peng and Duosheng Luo have took part in the research projects. All authors have read and approved the final manuscript.

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