

A novel partial agonist of PPAR γ with excellent effect on insulin resistance and type 2 diabetes

Hui-juan Liu, Cheng-yu Zhang, Fei Song, Ting Xiao, Jing Meng, Qiang Zhang, Cai-li Liang, Shan Li, Jing Wang, Bo Zhang, Yan-rong Liu, Tao Sun and Hong-gang Zhou

Tianjin Key Laboratory of Molecular Drug Research, Tianjin International Joint Academy of Biomedicine, Tianjin, People's Republic of China (H.-j.L., C.-y.Z., F.S., T.X., J.M., Q.Z., C.-l.L., S.L., J.W., B.Z., Y.-r.L.).

The State Key Laboratory of Medicinal Chemical Biology, College of Pharmacy, Nankai University, Tianjin, People's Republic of China (T.S., H.-g.Z.).

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Corresponding author

Dr. Hong-gang Zhou, The State Key Laboratory of Medicinal Chemical Biology,
College of Pharmacy, Nankai University, Tianjin 300071, People's Republic of China,
Email: hgzhou964@163.com, Tel.: +8613602042014.

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Abbreviations: PPAR γ , peroxisome proliferator-activated receptor γ ; TZDs, thiazolidinediones; DEX, dexamethasone; HepG2, human hepatocarcinoma cells; MDCK, Madin-Darby canine kidney; DMEM, dulbecco's minimum essential medium; FBS, fetal bovine serum; GOD-POD, glucose oxidase-peroxidase; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; DMSO, dimethyl sulfoxide; MS, mass spectrometry, LBD, ligand-binding domain; OGTT, oral glucose tolerance test; LDH, lactate dehydrogenase; GS, glycogen synthase; GSK, glycogen synthase kinase; LC-MS, liquid chromatography-mass spectrometry; FSG, fasting serum glucose; TC, total cholesterol; TG, triglyceride; LDL-c, low-density lipoprotein

cholesterol; HDL-c, high-density lipoprotein cholesterol.

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ABSTRACT:

Partial agonists of peroxisome proliferator-activated receptor- γ (PPAR γ) reportedly reverse insulin resistance in patients with type 2 diabetes mellitus. In this work, a novel non-thiazolidinedione partial PPAR γ ligand MDCCCL1636 was investigated. The compound displayed partial agonist activity in biochemical and cell-based transactivation assays, and reversed insulin resistance. MDCCCL1636 showed a potential anti-diabetic effect on an insulin-resistance model of HepG2 human hepatoma cells. High-fat diet-fed streptozotocin (HFD-STZ)-induced diabetic rats treated with MDCCCL1636 for 56 d displayed reduced fasting serum glucose and reversed dyslipidemia and pancreatic damage without significant weight gain. Furthermore, MDCCCL1636 had lower toxicity in vivo and in vitro than pioglitazone. MDCCCL1636 also potentiated glucose consumption and inhibited the impairment in insulin signaling targets, such as AKT, GSK3 β and glycogen synthase, in HepG2 human hepatoma cells. Overall, our results suggest that MDCCCL1636 is a promising candidate for the prevention and treatment of type 2 diabetes mellitus.

1. Introduction

Peroxisome proliferator-activated receptor γ (PPAR γ) is a ligand-dependent nuclear receptor that regulates gene expression associated with glucose homeostasis and insulin sensitization (Evans et al., 2004; Lehrke et al., 2005; McKenna et al., 2009). PPAR γ is expressed in the liver, fat, and skeletal muscle. PPAR γ gene deletion leads to insulin resistance and diet-induced obesity in mice, indicating that PPAR γ plays a regulatory role in lipid and glucose homeostases and tissue inflammation.

PPAR γ is a valid molecular target for metabolic syndrome and type 2 diabetes (Walczak et al., 2002; Waki et al., 2007). Several chemically diverse full agonists of PPAR γ , such as thiazolidinediones (TZDs), have been used as insulin-sensitizing drugs in type 2 diabetes. “Partial” agonists have recently been developed to reduce the side effects (e.g., weight gain, risk of heart attack and edema) of TZDs (Berger et al., 2005). “Full” and “partial” PPAR γ agonists are categorized by their transcriptional activities in the cell-based reporter assay (Reginato et al., 1998). Partial PPAR γ agonists possess higher safety margins than full PPAR γ agonists (Berger et al., 2005). Recent studies have exerted considerable efforts to design partial PPAR γ agonists that retain their insulin-sensitizing efficacy without significant side effects.

Insulin resistance, which results from the defective utilization of metabolites in insulin-targeted tissues, is the major cause of type 2 diabetes and obesity (Leclercq et al., 2007). The insensitivity of the liver to the biological effects of insulin leads to a decrease in insulin-induced glucose transport, and hence, Human hepatocarcinoma cells (HepG2) are a good experimental system to investigate insulin resistance and the

metabolic disorder (Postic et al., 2004). Dexamethasone (DEX) induces insulin resistance (Niels et al., 2009) by accelerating hepatic glucose production and proteometabolism and suppressing peripheral glucose transport and utilization. In this work, an in vitro insulin resistance cell model was established to screen active compounds that can reverse the insulin resistance (Sangeetha et al., 2010).

We searched for novel PPAR γ agonists in a library of structurally diverse organic compounds and identified a novel non-thiazolidinedione partial PPAR γ ligand, N-(4-hydroxyphenethyl)-3-mercapto-2-methylpropanamide (Fig. 1A, also known as MDCCCL1636). This ligand was synthesized in our laboratory (Li NN et al., 2014) and its partial activation of PPAR γ and effect on insulin resistance reversal were assessed in vitro. The cytotoxicity and the developmental toxicity in zebrafish embryos treated with MDCCCL1636 were evaluated and compared with those treated with pioglitazone. MDCCCL1636 effectively improved glucose tolerance and total plasma cholesterol level in high-fat diet-fed streptozotocin (HFD-STZ)-induced diabetic rats. We also investigated the major markers involved in insulin signaling, such as phosphatidylinositol 3-kinase (PI3K) signaling, to elucidate the mechanism by which MDCCCL1636 reverses insulin resistance.

2. Materials and methods

2.1 Cells and animals

human hepatocarcinoma HepG2, HEK-293T, HEK-293, NIH-3T3, and Madin-Darby canine kidney (MDCK) cells were obtained from KAIJI Company (Nanjing, China). These cells were maintained in dulbecco's minimum essential medium (DMEM)/high glucose with 10% fetal bovine serum (FBS) at 37 °C in a humidified 5% CO₂ environment.

AB strain zebrafish (*Danio rerio*) was obtained from Tianjin International Joint Academy of Biomedicine. Normoglycemic Wistar albino rats were obtained from Vital River Laboratory Animal Technology Co., Ltd (Beijing, China).

2.2 Antibodies and reagents

Antibodies to phosphoAkt (Ser⁴⁷³), phosphoGSK-3 β (Ser⁹) and phospho glycogen synthase (Ser⁶⁴⁵) were purchased from Affinity (USA).

3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) and dimethyl sulfoxide (DMSO) were purchased from Sangon Biotech (Shanghai, China). Glucose assay kit was purchased from Leadman Group Co. Ltd. (Beijing, China). STZ was purchased from Dalian Meilun Biotech Co., Ltd.

2.3 Methods

2.3.1 Transcriptional transactivation assay

HEK-293T cells cotransfected with pGal4-DBD (pBIND)-PPAR γ -ligand-binding domain (LBD) fusion and pG5luc vector were used to measure the transcriptional transactivation activities of the compounds through luciferase reporter

assay (Elke et al., 2006). Approximately 204-505 residues of the human PPAR γ -LBD were cloned into the pBIND to generate fusion proteins with the DNA-binding domain of GAL4. HEK-293T cells were grown in DMEM with 10% FBS at 37°C in 5% CO₂. HEK-293T cells were seeded in 96-well plates at a concentration of 2×10^4 cells per well. Lipofectamine 2000 transfection reagent was used for transfection. The recombinant plasmid pBIND-LBD (0.05 μ g/well) and pG5luc vector (0.15 μ g/well) were cotransfected into HEK-293T cells for 6 h. Subsequently, the cells were treated with various concentrations of MDCCCL1636 or pioglitazone for 24 h. The firefly luciferase activity was measured by using the ONE-GloTM Luciferase Assay System (Promega) in accordance with the manufacturer's protocol.

2.3.2 Mass spectrometry (MS) assay

The purified PPAR γ -LBD protein [13] (methods for expression and purification of PPAR γ -LBD are available in the supplementary materials) was diluted to 50 μ M in 10 mM ammonium acetate and incubated with MDCCCL1636 in a 1:3 molar excess for 15 min. The PPAR γ -LBD-MDCCCL1636 complex was analyzed with a Waters SYNAPT G1 High-Definition Mass Spectrometry.

2.3.3 Development of insulin-resistant model by using HepG2 cells

HepG2 cells were seeded in 96-well plates at a concentration of 1×10^4 cells per well and then treated in serum-free DMEM/high glucose overnight. Subsequently, the cells were induced with 100 nM, 1 μ M, 5 μ M, and 10 μ M DEX for 24, 48, and 72 h. At selected time points, the cells were washed thrice with PBS and stimulated for 24 h with 1 nM insulin. Glucose uptake was detected by using a glucose assay kit. The kit

was used to detect the glucose level in the medium via the glucose oxidase-peroxidase (GOD-POD) method. Subsequently, we calculated the consumption of glucose. The best time point for the insulin-resistant model was determined by measuring the maximum inhibition in glucose uptake (Yuan et al., 2003). The insulin-resistant model was validated using pioglitazone. The HepG2 cells were induced with 100 nM DEX for 24 h and then treated with varying concentrations of pioglitazone for 24 h with 1 nM insulin. Glucose uptake was detected by using a glucose assay kit.

2.3.4 Effect of MDCCCL1636 on glucose uptake in insulin-resistant HepG2 cells

MDCCCL1636 and pioglitazone were used to treat the normal control cells and insulin-resistant cells for 24 h to assess the effect of MDCCCL1636 and pioglitazone on glucose uptake. A glucose assay kit was used for the glucose uptake experiments.

2.3.5 Cytotoxicity assessment

MDCCCL1636 and pioglitazone were assessed for their cytotoxic effects on HEK-293, HepG2, NIH-3T3 and MDCK cells by using an MTT reagent. The assay was performed 24 h after treatment with varying concentrations of MDCCCL1636 and pioglitazone. Formazan concentration, which is directly proportional to cell viability, was measured at 492 nm.

2.3.6 Developmental toxicity assay in zebrafish embryos

AB line zebrafish was maintained in accordance with the standard procedures (M. Westerfield, 1994). The night before breeding, adult male and female zebrafish were placed in a breeding tank and then separated from each other with a mesh screen. The embryos were generated through natural mating the next morning, after turning on the

light and withdrawing the mesh screen. The embryos were collected within 30 min after spawning, rinsed thrice, transferred into Petri dishes containing the embryo medium, and then cultured at 28.5 °C.

The protocol described by Mark *et al.* (2004) was adopted in this experiment. The normal embryos were selected and transferred into a multiwell microplate with one embryo per well in 300 µL treatment solution. A total of 30 embryos at 4 h postfertilization were treated with various concentrations of MDCCCL1636 and pioglitazone, and the control group was treated with 0.1% DMSO. Developmental phenotypes were observed every 24 h for 96 h. All experiments were repeated thrice. The mortality and malformation rate of embryos were calculated at the end of the experiment. The study was performed in accordance with the national and institutional guidelines for animal welfare.

2.3.7 In vivo efficacy of MDCCCL1636 in HFD-STZ-induced type 2 diabetic rats

Normoglycemic male Wistar albino rats weighing 180–200 g were used. All rats, except for those in the normal control group, were fed with HFD (20% glucose, 10% egg yolk powder, 10% lard, 0.2% bile salts, 1.5% cholesterol, and 58.3% normal commercial pellet diet). After 10 d of HFD, the rats were fasted overnight for 12h and then given a single injection of freshly prepared solution of STZ (40 mg/kg) in citrate-phosphate buffer (0.1 M, pH 4.2) (Ashok et al., 2011). Hyperglycemia in the rats was assessed by measuring fasting serum glucose (FSG) levels, after 72 h of STZ administration. Rats with an FSG levels higher than 13.89 mmol/L were selected for the subsequent experiments (Ashok et al., 2011). The rats were randomly divided into

six groups of six animals each: (1) normal control group (rats without any treatment); (2) diabetic control group (diabetic rats treated with vehicle); (3) positive control group (diabetic rats treated with pioglitazone at 30 mg/kg/day); (4) high dose group of MDCCCL1636 (diabetic rats treated with MDCCCL1636 at 15 mg/kg/day); (5) medium dose group of MDCCCL1636 (diabetic rats treated with MDCCCL1636 at 7.5 mg/kg/day); (6) lower dose group of MDCCCL1636 (diabetic rats treated with MDCCCL1636 at 3.75 mg/kg/day). The treatment was continued for 56 days.

In the oral glucose tolerance test (OGTT) assay, 12 h fasted rats were administered with vehicle or drugs after FSG measurement, and then with an oral bolus of glucose (2 g/kg). Subsequently, the blood glucose level was measured at 0, 0.5, 1, and 2 h.

On the last day, FSG levels were measured and then blood was withdrawn from the heart. Serum was analyzed for triglyceride (TG), total cholesterol (TC), high-density lipoprotein cholesterol (HDL-c), and low-density lipoprotein cholesterol (LDL-c) after being separated by centrifugation at $3000 \times g$ for 10 min. The animals were sacrificed, and the pancreas was subjected to histopathological studies (Kamalakkannan et al. 2004). The lipid profiles were assessed using commercial kits. Formalin-fixed pancreatic tissue was cut in 4 μ m thick sections and then stained with hematoxylin and eosin (H&E) (Ashok et al., 2011). The stained sections were blindly examined using a light microscope (Nikon, Tokyo, Japan). The study was performed in accordance with the national and institutional guidelines for animal welfare.

2.3.8 Effect of MDCCCL1636 on the expression of insulin signaling markers in

PI3K signaling

Insulin-sensitive cells, non-treated insulin-resistant cells, and MDCCCL1636 or pioglitazone-treated insulin-resistant cells were lysed at 4°C with RIPA lysis buffer. The cell lysates were subjected to SDS-PAGE and then Western blot analysis using anti-phospho antibodies for AKT, GSK3 β , and GS.

2.3.9 Statistical analysis.

Datas were reported as mean \pm SD. All data were analyzed using one-way analysis of variance (ANOVA) followed by Bonferroni post-hoc test (SPSS software package version 17.0). Statistical significance was considered at $p < 0.05$.

3 Results

3.1 Identification of the partial PPAR γ agonist MDCCCL1636

MDCCCL1636 (Molecular Weight: 239.1 Da) was identified as a lead hit among a class of partial PPAR γ ligands through transcriptional transactivation assay of a PPAR γ agonist library. In HEK-293T cells, MDCCCL1636 (EC_{50} = 7.092 μ M) was a partial agonist with 40% efficacy compared with pioglitazone (EC_{50} = 3.36 μ M) at 100 μ M (Fig. 1B).

3.2 Verifying the interaction of MDCCCL1636 and PPAR γ -LBD

MS data revealed peaks corresponding to the molecular weight of the protein PPAR γ -LBD and the complex PPAR γ -LBD-MDCCCL1636. The molecular weights of the PPAR γ -LBD protein the complex were 32910 Da (Fig. 2A) and 33146 Da (Fig. 2B), respectively. The difference in molecular weight between PPAR γ -LBD protein and the PPAR γ -LBD-MDCCCL1636 complex was 236 Da, which was approximately equal to the molecular weight of MDCCCL1636. This result demonstrated that the MDCCCL1636 strongly interacted with the ligand binding domain of human PPAR γ .

3.3 Effect of MDCCCL1636 on the insulin-resistant model of HepG2 cells

3.3.1 Time and concentration course analyses of DEX induction on HepG2 cells

Time and concentration course analyses of DEX treatment at 100 nM, 1 μ M, 5 μ M, and 10 μ M concentrations in 24, 48, and 72 h were performed. Insulin resistance of DEX was also assessed based on the inhibition of glucose uptake. Insulin-stimulated glucose uptake was reduced maximally to approximately 26.83% at 24 h of 100 nM DEX treatment compared with the control (Fig. 3A). Thus, treatment

with 100 nM DEX for 24 h was the optimal conditions for the establishment of insulin-resistant model.

Cell viability was assayed at different time points (24, 48, and 72 h) of DEX induction to eliminate the influence of DEX toxicity on the inhibition of glucose uptake. DEX exhibited no significant cytotoxicity to the cells at 24 h, which confirmed the insulin-resistant inducing effect of DEX on HepG2 cells (Fig. 3B).

3.3.2 Validation of DEX-induced insulin-resistant model

Pioglitazone was used as a positive control to determine the validity and the reversal properties of the DEX-induced insulin-resistant model. The induced HepG2 cells were treated with varying concentrations of pioglitazone for 24 h. Glucose uptake analysis showed that pioglitazone reversed the DEX-induced insulin resistance in a dose-dependent manner (Fig. 3C).

3.3.3 Glucose uptake potential of MDCCCL1636 on the DEX-induced insulin-resistant model

The effects of MDCCCL1636 and pioglitazone on insulin resistance were assessed and compared using the DEX-induced insulin-resistant model. MDCCCL1636 effectively restored the DEX-induced glucose uptake inhibition in a dose-dependent manner (Fig. 3D). MDCCCL1636 maximally augmented glucose uptake to approximately 45.13% at 150 μ M in the DEX-induced insulin-resistant model. The response of MDCCCL1636 at 75 μ M (approximately 41.45%) to insulin resistance reversal was higher than that of pioglitazone (34.82%).

3.4 Cytotoxicity of MDCCCL1636 and pioglitazone on HEK-293, HepG2,

NIH-3T3, and MDCK cells

The cytotoxic effects of MDCCCL1636 and pioglitazone on HEK-293, HepG2, NIH-3T3, and MDCK cells were assessed via a cytotoxicity assay. MDCCCL1636 showed negligible effect on cell viability, whereas pioglitazone exhibited about 25% (Figs. 4A-C) and 50% (Fig. 4D) toxic effects at its highest concentration. This result confirmed that MDCCCL1636 had lower cytotoxicity than pioglitazone.

3.5 Toxic effects of MDCCCL1636 and pioglitazone on embryonic development of zebrafish

The mortality rate of zebrafish embryos treated with various concentrations of MDCCCL1636 and pioglitazone for 96 h is presented in Fig. 5A. Treatment with 1 mM of pioglitazone for 96 h resulted in 95% embryo death compared with 65% embryo death after MDCCCL1636 treatment. MDCCCL1636 and pioglitazone showed dose-dependent lethal effects. The lethal concentration of MDCCCL1636 that caused 50% mortality (LC_{50}) in embryos was 742.569 μ M, whereas that of pioglitazone was 390.226 μ M. Exposure experiments revealed that the effective concentrations of MDCCCL1636 and pioglitazone to induce 50% malformations (EC_{50}) of the embryos was 916.044 μ M and 420.741 μ M (Fig.5B), respectively.

Pericardial edema and axial malformation were observed in embryos exposed to 50 mM pioglitazone, whereas no obvious malformation was observed in embryos exposed to less than 250 mM MDCCCL1636. Embryos only subjected to more than 500 mM MDCCCL1636, would showed pericardial edema, and those exposed to more than 750 mM MDCCCL1636 would have axial malformations (Fig. 6). These

results indicated that MDCCCL1636 exerted lower toxicity on zebrafish embryos than pioglitazone.

3.6 Hypoglycemic effect of MDCCCL1636 on HFD-STZ-induced type 2 diabetic rats

Administration of MDCCCL1636 at 15 mg/kg/day and pioglitazone at 30 mg/kg/day for 56 d resulted in a significant decrease in FSG compared with the diabetic control rats with a reduction of 20.72% and 18.73%, respectively.

As shown in Table 1, the HFD-STZ-induced diabetic rats had higher serum TG, TC, and LDL-c and significantly lower HDL-c than the normal control rats. Treatment with 7.5 mg/kg MDCCCL1636 and 30 mg/kg pioglitazone significantly reduced serum TG (53.5% and 31.7%, respectively) and TC (56.86% and 43.4%, respectively) compared with diabetic control rats. In addition, the rats treated with MDCCCL1636 and pioglitazone had significantly lower LDL-c (65.4% and 45.6%, respectively, $p < 0.001$) while significantly higher HDL-c (16.1% and 9.7%, respectively) than the diabetic controls. Administration of pioglitazone at 30 mg/kg for 56 d resulted in a significant increase in body weight (12.13%), whereas MDCCCL1636-treated groups did not show significant changes in body weight in comparison with the diabetic control rats.

OGTT was performed in HFD-STZ-induced type 2 diabetic rats on day 42 of the treatment period. After being challenged with an oral bolus of glucose, the MDCCCL1636-treated animals showed lower glucose excursion compared with the vehicle-treated diabetic animals. The results indicated that MDCCCL1636 improved

the impaired glucose tolerance of type 2 diabetic rats (Fig. 7).

Fig. 8A showed the histopathological changes in the pancreatic islet of the experimental groups. The number and area of islets in the pancreas (Figs. 8B-C) were also analyzed. Neither β -cell damage nor inflammatory changes were observed in the normal architecture of pancreatic islet (Fig. 8A1). However, the β -cells were obviously damaged and the area of pancreatic islets was significantly reduced in the HFD-STZ induced diabetic rats (Fig. 8A2). Damage to the pancreatic islets and β -cell in the diabetic rats was restored by MDCCCL1636 and pioglitazone, as evidenced by their protective effects on β -cell damage (Figs. 8A3-4). Compared with pioglitazone, MDCCCL1636 also more effectively improved the fatty degeneration of renal tubular epithelial cells of the diabetic rats (supporting data are provided in the supplementary materials).

3.7 Effect of MDCCCL1636 and pioglitazone on insulin signaling in insulin-resistant cells

AKT Ser⁴⁷³ phosphorylation was reduced in insulin-resistant cells compared with control cells. The insulin-resistant cells showed 73.9% lower AKT Ser⁴⁷³ phosphorylation than control cells. Both pioglitazone and MDCCCL1636 significantly increased AKT Ser⁴⁷³ phosphorylation by approximately 1.86-fold and 3.68-fold, respectively, (Fig. 9A). AKT expression was similar in the normal control and insulin resistant cells. Both pioglitazone and MDCCCL1636 increased AKT Ser⁴⁷³ phosphorylation, although MDCCCL1636 showed better effects than pioglitazone.

GSK-3 β Ser⁹ phosphorylation in the insulin-resistant cells was reduced by approximately 19.1% compared with that in the normal control cells. Pioglitazone and MDCCCL1636 increased GSK-3 β phosphorylation in the insulin-resistant cells (Fig. 9B). GSK-3 β expression was similar in the normal control and insulin-resistant cells. MDCCCL1636 and pioglitazone upregulated GSK-3 β Ser⁹ phosphorylation by 0.53-fold and 0.21-fold, respectively.

Glycogen synthase Ser⁶⁴⁵ phosphorylation was higher in the insulin-resistant cells than in the normal control cells. DEX-treated insulin-resistant cells showed lower activity than the insulin-sensitive cells. Pioglitazone and MDCCCL1636 increased glycogen synthase activity by 62% and 70%, respectively, in the insulin-resistant cells compared with the normal control cells (Fig. 9C). GS expression was similar in the normal control and the insulin resistant cells. The insulin-resistant cells treated with MDCCCL1636 had higher GS activity than those treated with pioglitazone.

4. Discussion

This study introduced the partial PPAR γ activator MDCCCL1636. Transactivation assay data confirmed that the propanamide derivative functions as a partial PPAR γ agonist that strongly combines with PPAR γ -LBD, as determined by MS.

Insulin-sensitizing agents could function as drugs to treat various metabolic disorders caused by insulin resistance (Kahn et al., 2000). An in vitro model that mimics insulin resistance was established to screen insulin-sensitive agents (Olivares-Reyes et al., 2009; Sakoda et al., 2000). The model mimicked the exact in vivo clinical insulin-resistant conditions, and it can be reversed by insulin-sensitizing drug treatment (Sakoda et al., 2000). In this work, a model of HepG2 cells was developed using DEX and the model was used to assess the potential of MDCCCL1636 in insulin resistance reversal. The extent of insulin resistance of the model was measured through the inhibition of glucose uptake. In this work, the maximum desensitization (approximately 26.83%) of insulin-stimulated glucose uptake was achieved after 24 h of 0.1 μ M DEX treatment.

In this study, the TZDs pioglitazone was used to validate the developed resistance model. TZDs as insulin-sensitizing drugs reportedly reverse DEX-induced insulin resistance by suppressing the adverse effects of DEX on insulin sensitivity and glucose tolerance (Samuel, 2011; Willi et al., 2002). Concentration course analysis of the effect of pioglitazone on the insulin-resistant model revealed that TZD reversed DEX-induced insulin resistance in a dose-dependent manner. This study determined

whether or not MDCCCL1636 could also reverse the DEX-induced impairment in glucose uptake. Similar to pioglitazone, MDCCCL1636 showed potential reversal of DEX-induced insulin resistance as evidenced by the restoration of glucose uptake. MDCCCL1636 and pioglitazone showed similar effects on insulin resistance, however, MDCCCL1636 had lower toxicity on cells and zebrafish embryos than pioglitazone. This result may be ascribed to the partial PPAR γ agonist activity of MDCCCL1636.

Rats fed with HFD may develop insulin resistance (Kraegen et al., 1991). In addition, low-dose STZ leads to a mild β cell dysfunction and impairment of insulin secretion, which result in hyperglycemia (Mythili et al., 2004). Thus, feeding rats with HFD and treating them with low-dose STZ (40 mg/kg) can mimic type II diabetes (Reed et al., 2000; Parveen et al., 2010). In the present work, the HFD-STZ-treated type 2 diabetic rats showed significantly increased serum glucose and β -cell dysfunction, as well as decreased body weight and dyslipidemia. Our results showed that MDCCCL1636 reduced the serum glucose level and improved the impaired glucose tolerance in type 2 diabetes.

In the HFD-STZ-induced diabetic rats, dyslipidemia was manifested in the increased levels of LDL-c, TG, and TC, and in reduced HDL-c levels. The hypercholesterolemia and hypertriglyceridemia observed in the model resulted from the increased absorption of TGs and TC from the HFD and from the elevated concentrations of very-low-density lipoproteins, which consequently increased LDL-c and reduced HDL-c levels (Zammit et al., 2001; Taskinen et al., 2003). In the present

work, MDCCCL1636 treatment significantly reduced the TG, TC, and LDL-c levels while increased the HDL-c level in the HFD-STZ-induced diabetic rats. MDCCCL1636 improves dyslipidemia possibly by inhibiting hepatic TG secretion or increasing peripheral TG clearance (Kamalakkannan et al., 2005). MDCCCL1636 reverses the insulin resistance, which consequently improves the lipid metabolism.

In the present study, MDCCCL1636 upregulated PI3K signaling. Our data suggested that DEX induced HepG2 cells insulin resistance by impairing the phosphorylation of insulin signaling proteins of AKT, GSK-3 β , or GS. DEX treatment reduced insulin-stimulated AKT and GSK-3 β phosphorylation and blocked the dephosphorylation and activation of GS.

In this study, DEX treatment reduced insulin-stimulated AKT Ser⁴⁷³ phosphorylation by approximately 73.9%. Both MDCCCL1636 and pioglitazone increased AKT Ser⁴⁷³ phosphorylation in the insulin-resistant cells, but the effect of MDCCCL1636 was 0.64-fold higher than that of pioglitazone. The insulin-resistant cells also exhibited 19.1% lower GSK-3 β Ser⁹ phosphorylation than the normal control cells. MDCCCL1636 can reverse such a reduction. DEX inhibited the glycogen synthase activity. MDCCCL1636 and pioglitazone reduced GS phosphorylation by 70% and 62%, respectively. DEX induction significantly desensitized HepG2 cells toward insulin-stimulated glucose uptake (Pereira et al., 2008). The possibility of this desensitization can also be attributed to impairment in the insulin signaling targets, including AKT, GSK3 β , and GS. Our results showed that MDCCCL1636 effectively restored DEX-induced desensitization by restoring AKT

and GSK3 β phosphorylation and GS activity.

In conclusion, the novel compound MDCCCL1636 can partially activate PPAR γ . MDCCCL1636 can also reduce FSG and reverse the dyslipidemia and damage of the pancreas without significantly increasing body weight. Moreover, MDCCCL1636 was less toxic in vivo and in vitro than pioglitazone. Therefore, MDCCCL1636 is a potential treatment for type 2 diabetes.

Authorship Contributions

Participated in research design: Sun, and Zhou;

Conducted experiments: Song, H.-j.Liu, C.-y.Zhang, Xiao, Meng, Liang, and Q. Zhang;

Performed data analysis: Song, H.-j.Liu, C.-y.Zhang, Li and Wang;

Wrote or contributed to the writing of the manuscript: Song, H.-j.Liu, C.-y.Zhang, B.Zhang, and Y.-r.Liu.

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Footnotes

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Hui-juan Liu, Cheng-yu Zhang, and Fei Song contributed equally to this study.

Figure Legends

Fig. 1. (A) Structure of N-(4-hydroxyphenethyl)-3-mercapto-2-methylpropanamide. (B) Representative concentration–response curves of reporter gene transactivation by MDCCCL1636 compared with pioglitazone. Stand and error bars represent three independent experiments, and each experiment was performed in triplicate. All values are expressed as mean \pm SD. Significance level was determined by one-way ANOVA followed by Bonferroni post-hoc test. ^{##}*p* < 0.05, ^{###}*p* < 0.001 vs. normal control.

Fig. 2. (A) Mass spectra of PPAR γ -LBD protein, (B) PPAR γ -LBD-MDCCCL1636 complex.

Fig. 3. Insulin-resistant cell model and drug screening. (A) Time and concentration course analyses of DEX induction in HepG2 cells. (B) Effect of DEX induction on cytotoxicity. (C) Validation of DEX-induced insulin-resistant model. (D) Comparative assessment of glucose uptake potential of MDCCCL1636 and pioglitazone on the insulin-resistant model of HepG2 cells. Stand and error bars represent three independent experiments and each experiment was conducted in triplicate. ^{**}*p* < 0.05 and ^{***}*p* < 0.01 vs. insulin resistance control; ^{##}*p* < 0.05, ^{###}*p* < 0.001 vs. normal control.

Fig. 4. Cytotoxicity of MDCCCL1636 and pioglitazone. Cytotoxicity of MDCCCL1636 and pioglitazone on (A) HEK-293, (B) HepG2, (C) NIH-3T3, and (D) MDCK cells. Stand and error bars represent three independent experiments, and each experiment was performed in triplicate.

Fig. 5. Cytotoxicity of MDCCCL1636 and pioglitazone on the embryonic

development of zebrafish. (A) Lethality curve of embryos exposed to MDCCCL1636 or pioglitazone. The LC_{50} of MDCCCL1636 and pioglitazone were 742.569 μ M ($r^2=0.993$) and 390.226 μ M ($r^2=0.985$), respectively. (B) Malformation curve of embryos exposed to MDCCCL1636 or pioglitazone. The EC_{50} of MDCCCL1636 and pioglitazone were 916.044 μ M ($r^2=0.971$) and 420.741 μ M ($r^2=0.999$), respectively.

Fig. 6. Concentration-dependent developmental toxicity endpoints in zebrafish.

Embryos were exposed to either water or to the indicated MDCCCL1636 or pioglitazone concentrations. Lateral view of 96 hpf embryos revealing numerous malformations. Axial malformation, AM; pericardial edema, PE.

Fig. 7. Effect of MDCCCL1636 and pioglitazone on oral glucose tolerance in normal and HFD-STZ induced diabetic rats. (A) Blood glucose curves in the oral glucose tolerance test. (B) Area under the curve in the oral glucose tolerance test. All values are expressed as mean \pm SD. Significance was determined by one-way ANOVA followed by Bonferroni post-hoc test, ** $p < 0.01$ vs. diabetic control.

Fig. 8. Histopathological changes in the pancreatic islets of rats in the experimental groups. (A1) Light micrograph of normal control rats showing normal β -cells in the pancreatic islet (H & E, 200 \times), (A2) HFD-STZ-induced diabetic control rats showing pancreatic islet damage (H&E, 200 \times), (A3) MDCCCL1636 at 7.5 mg/kg/day reversed pancreatic islet damage (H&E, 200 \times), (A4) pioglitazone at 30 mg/kg/day reversed pancreatic islet damage (H&E, 200 \times). (B) Islets number analysis of experimental groups. (C) Islets area analysis of experimental groups. All values are expressed as mean \pm SD. Significance was determined by one-way ANOVA followed by

Bonferroni post-hoc test, ** $p < 0.01$ vs. normal control.

Fig. 9. Effect of MDCCCL1636 and pioglitazone on insulin signaling in insulin-resistant cells. HepG2 cells were incubated with or without 100 nM DEX for 24 h and in 1 nM insulin with or without the drugs (pioglitazone or MDCCCL1636) for another 24 h. Cell lysates were separated by SDS-PAGE and subjected to Western blot analysis with (A) anti-phosphoAkt (ser473) antibody, (B) anti-phosphoGSK-3 β (Ser⁹) antibody, and (C) anti-phospho glycogen synthase (Ser⁶⁴⁵) antibody. Data are expressed as a percentage of the normal control. ** $p < 0.05$, *** $p < 0.001$ vs. insulin resistance control; ## $p < 0.05$, ### $p < 0.001$ vs. normal control.

Table 1. Effects of MDCCCL1636 on body weight and biochemical parameters of normal and diabetic rats. All values are expressed as mean \pm SD. Significance level was determined by one-way ANOVA followed by Bonferroni post-hoc test. ^ap < 0.001 vs. normal control; ^bp < 0.05, ^cp < 0.001 vs. diabetic control.

Biochemical parameter	Normal	Diabetic	Pioglitazone		MDCCCL1636	
	Control	Control	30 mg/kg	3.75 mg/kg	7.5 mg/kg	15 mg/kg
FSG (mmol/L)	5.6 \pm 0.9	25.1 \pm 2.9 ^a	20.4 \pm 2.8 ^c	21.8 \pm 2.3 ^b	20.9 \pm 1.9 ^c	19.9 \pm 1.1 ^c
TG (mmol/L)	0.81 \pm 0.10	1.01 \pm 0.27 ^a	0.69 \pm 0.19 ^c	0.74 \pm 0.26 ^b	0.47 \pm 0.15 ^c	0.42 \pm 0.12 ^c
TC (mmol/L)	1.47 \pm 0.18	8.88 \pm 3.64 ^a	5.03 \pm 1.27 ^b	4.80 \pm 0.76 ^b	3.83 \pm 1.11 ^c	4.40 \pm 0.78 ^c
HDL-C (mmol/L)	0.78 \pm 0.01	0.62 \pm 0.12 ^a	0.68 \pm 0.13 ^b	0.67 \pm 0.13 ^b	0.72 \pm 0.07 ^c	0.77 \pm 0.10 ^c
LDL-C (mmol/L)	0.26 \pm 0.03	2.98 \pm 1.44 ^a	1.62 \pm 1.27 ^b	1.46 \pm 0.56 ^c	1.03 \pm 0.49 ^c	1.21 \pm 0.70 ^c
Body Weight (g)	553.85 \pm 22.71	365.09 \pm 12.53 ^a	409.37 \pm 16.39 ^b	379.94 \pm 22.46	376.27 \pm 21.24	380.51 \pm 7.49

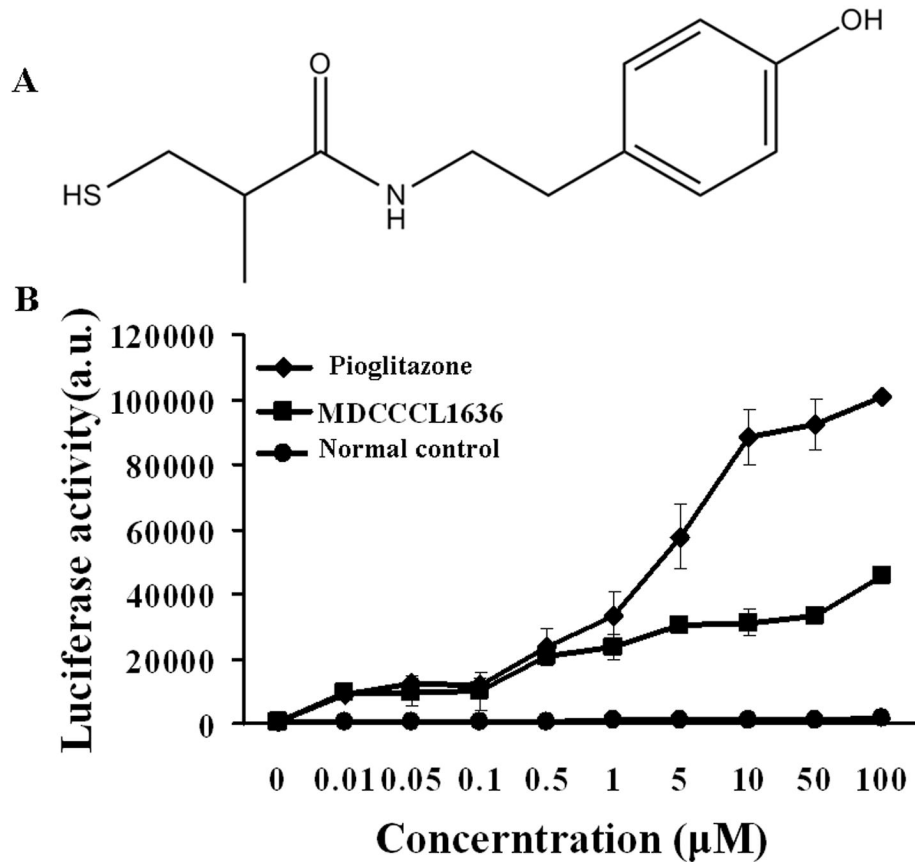


Fig.1

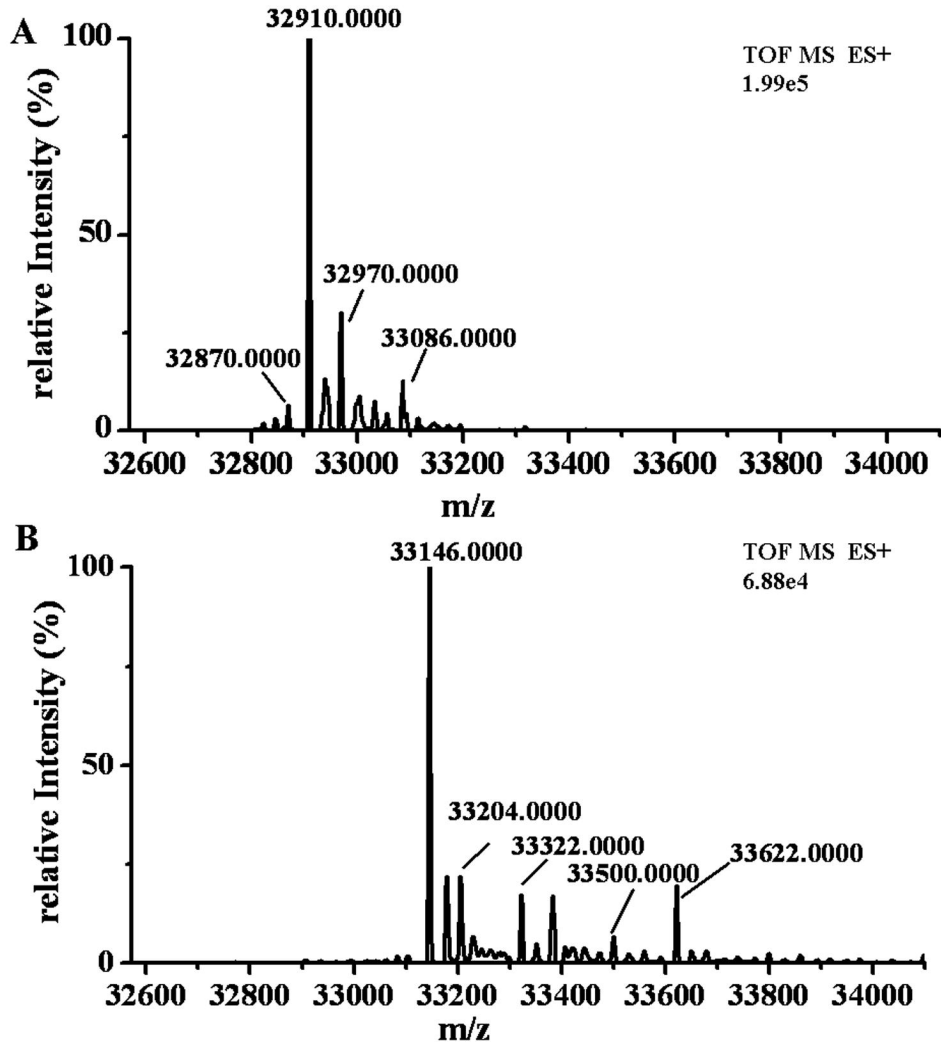


Fig. 2

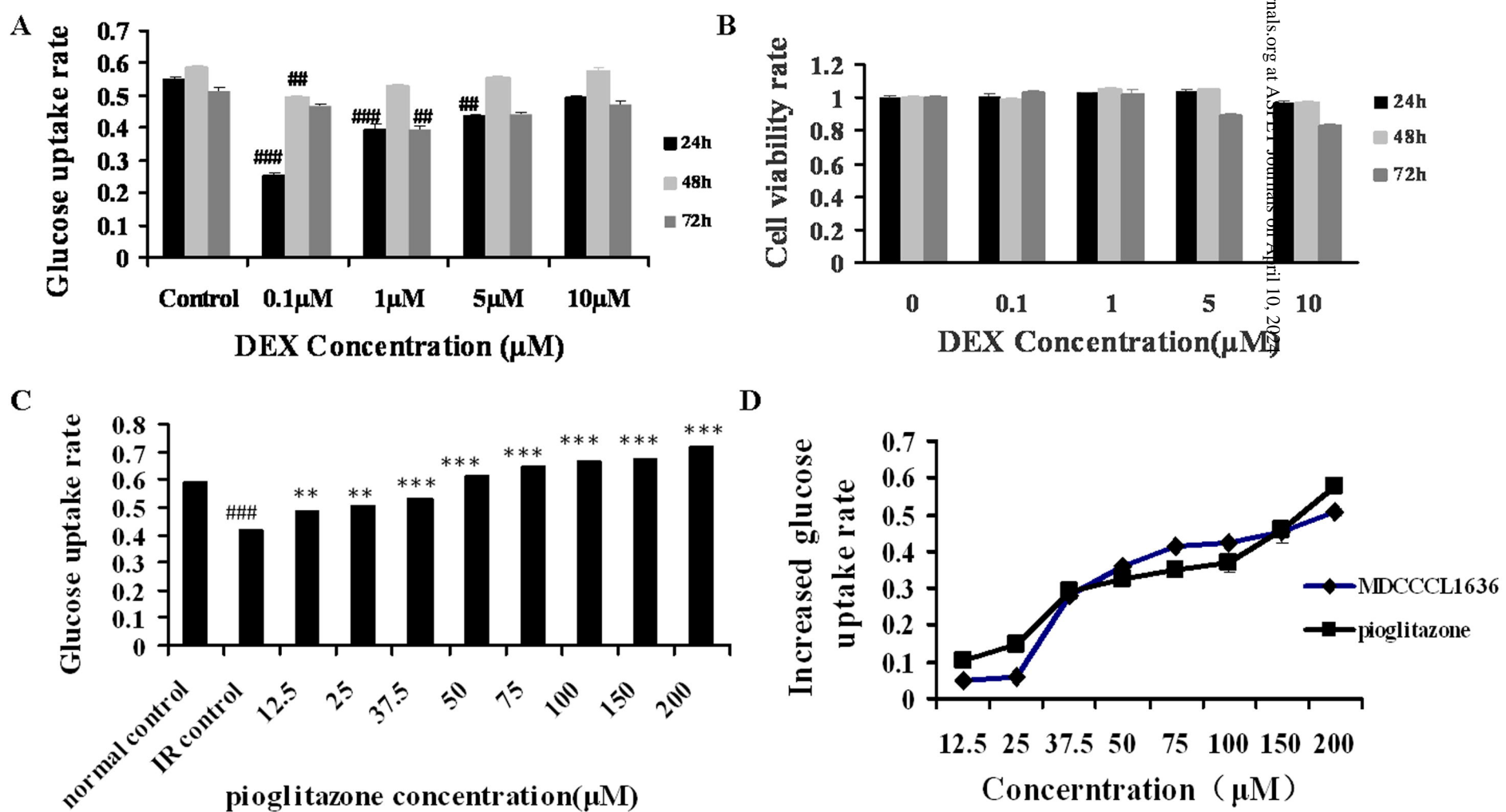


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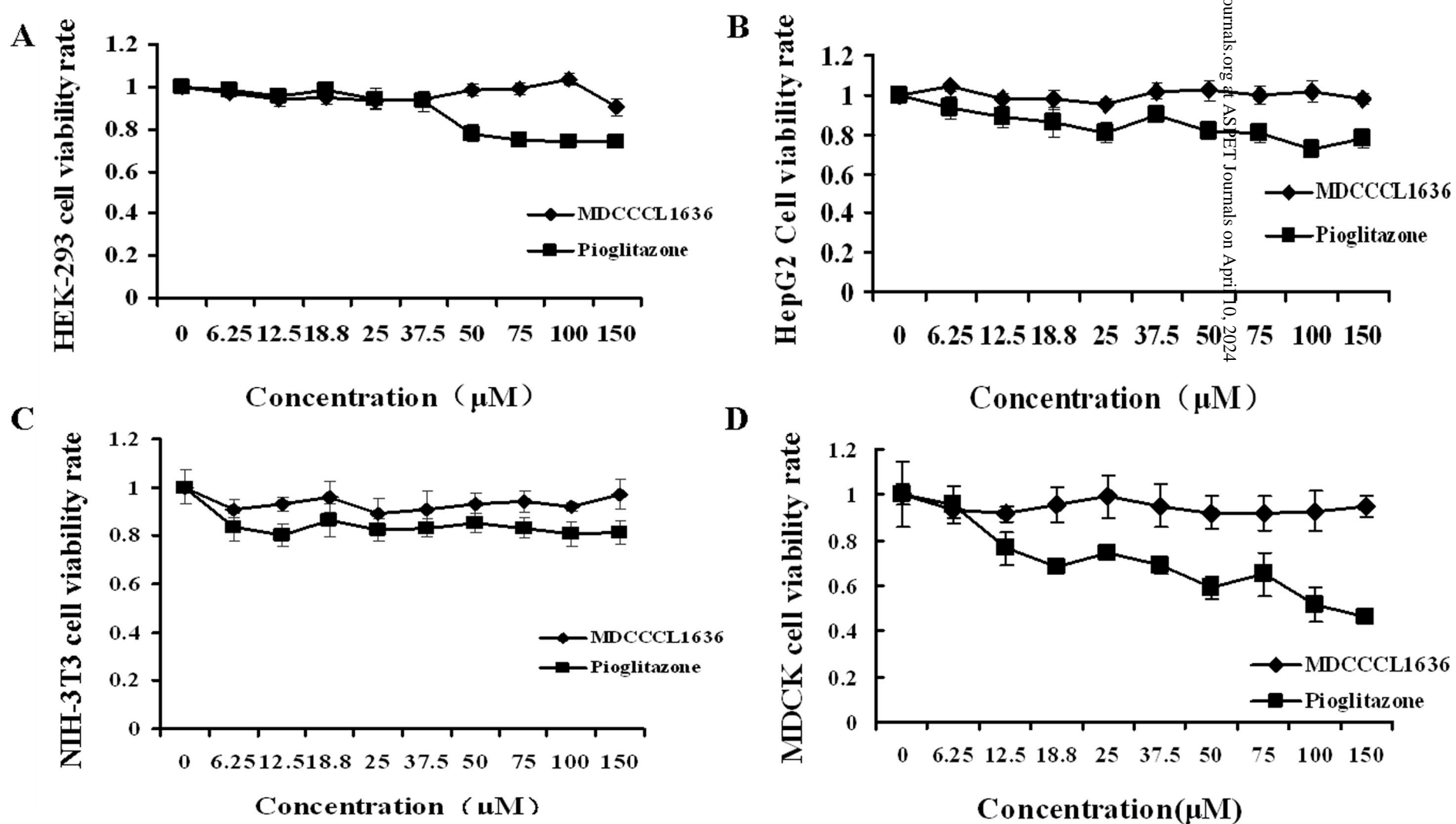


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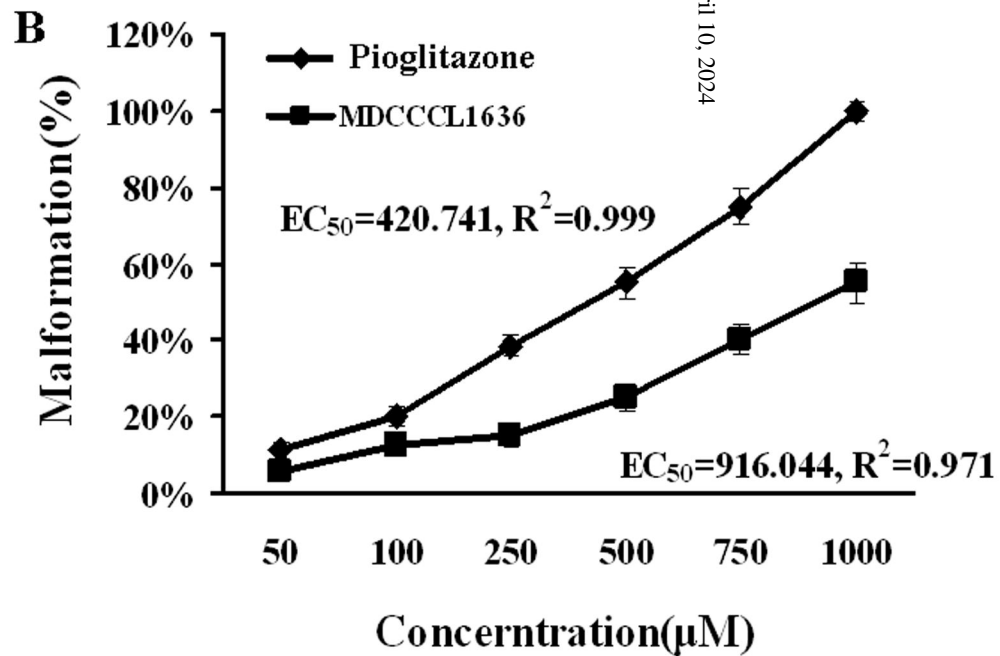
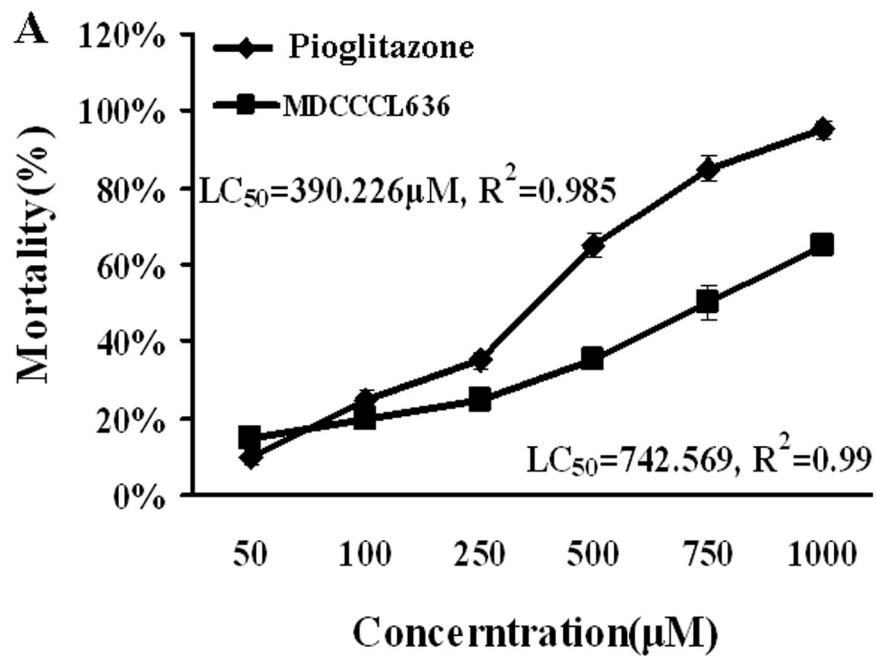


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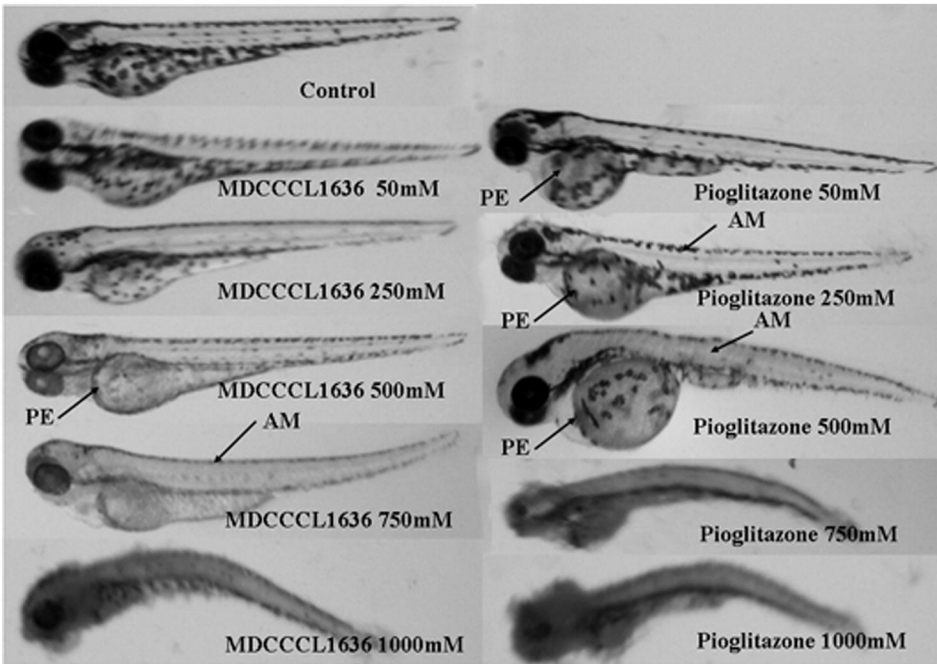


Fig. 6

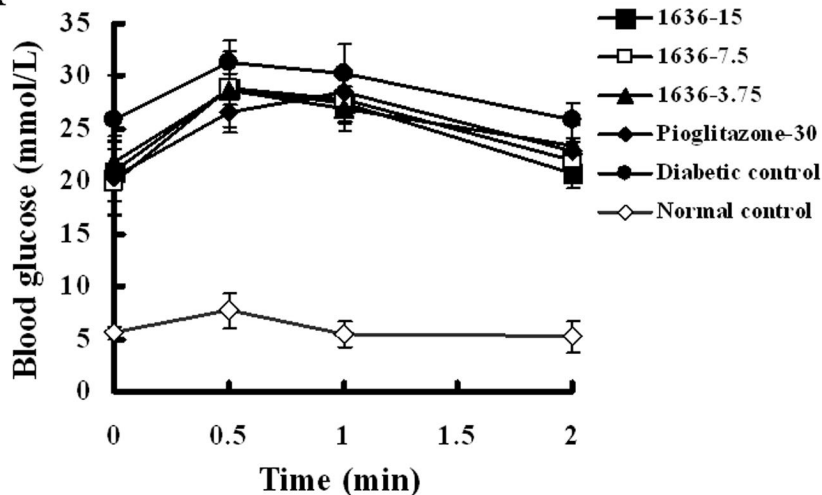
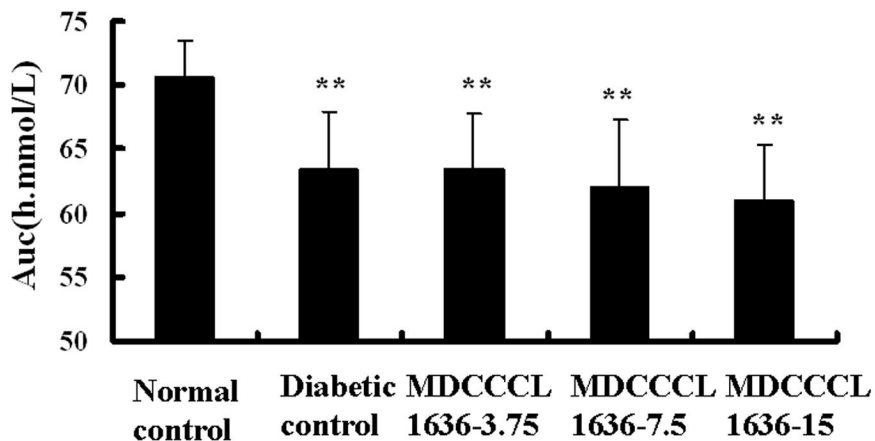
A**B**

Fig. 7

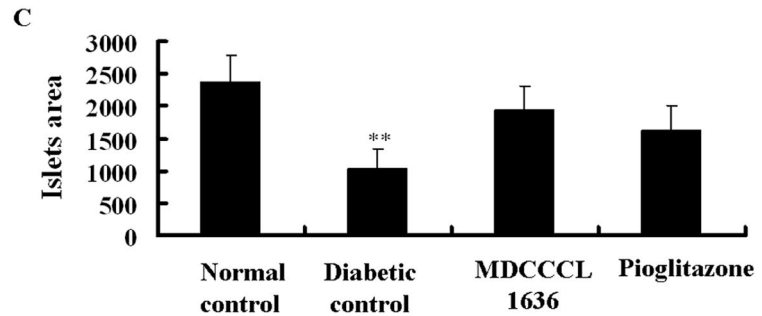
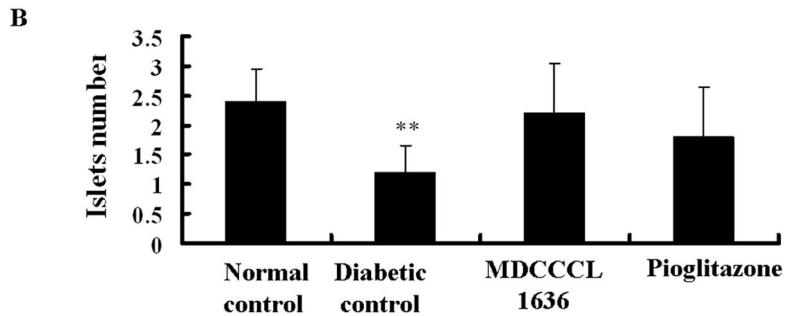
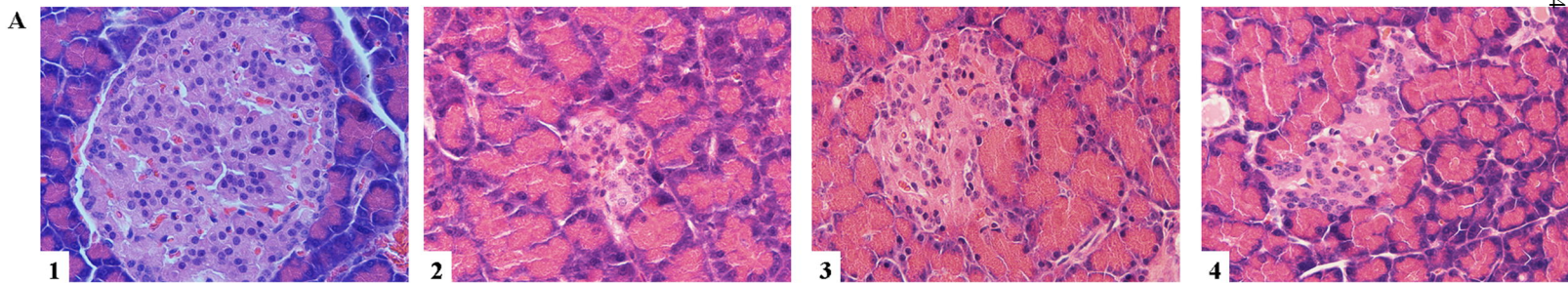


Fig. 8

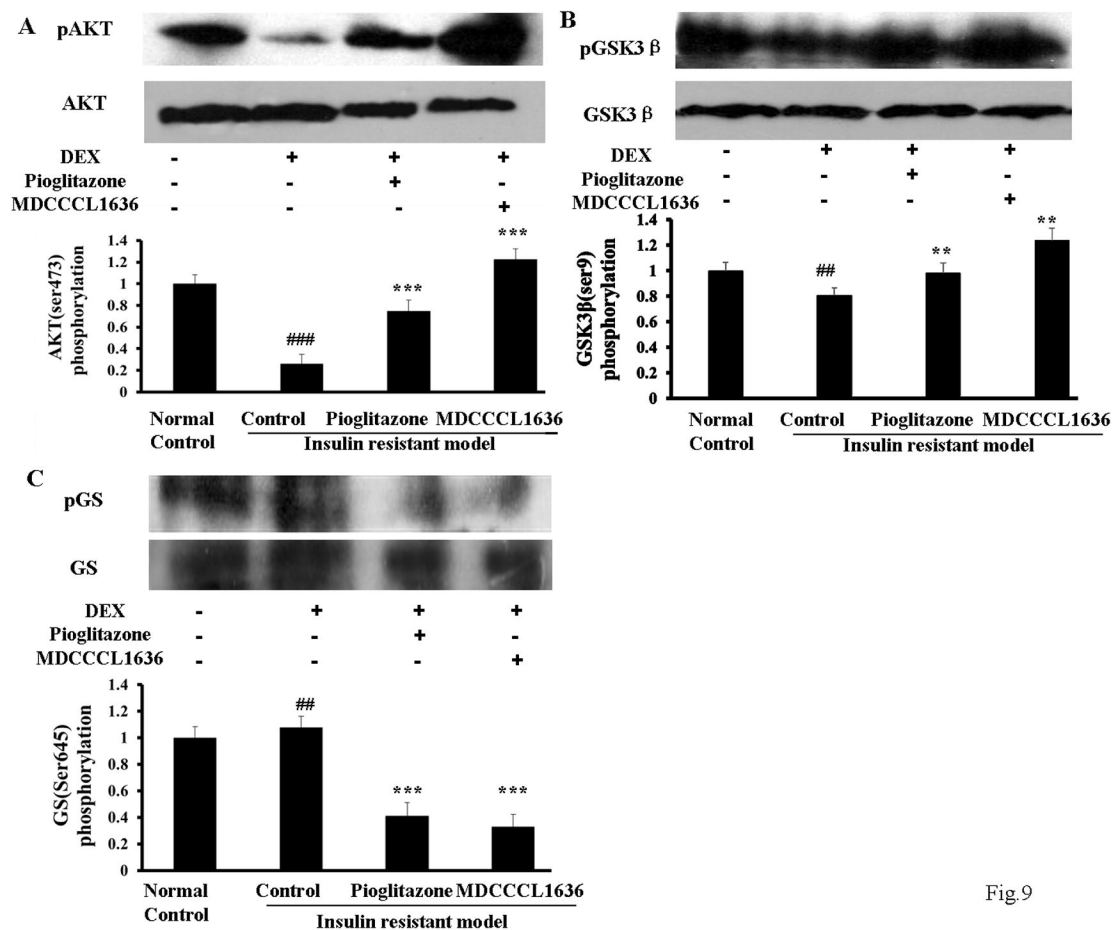


Fig.9

A novel partial agonist of PPAR γ with excellent effect on insulin resistance and type 2 diabetes

Hui-juan Liu, Cheng-yu Zhang, Fei Song, Ting Xiao, Jing Meng, Qiang Zhang, Cai-li Liang, Shan Li, Jing Wang, Bo Zhang, Yan-rong Liu, Tao Sun and Hong-gang Zhou

1. Methods for expression and purification of PPAR γ -LBD

The PPAR γ -LBD (the ligand binding domain of human PPAR γ , residues 206–477) was cloned into the expression vector pET15b. *E.coli* BL21 (DE3) host cells transformed with this expression plasmid were grown in LB at 37 °C to an A₆₀₀ of 1 and induced with 0.1 mM IPTG at 16 °C [13]. The cells were harvested, resuspended in extract buffer (50 mM Tris, pH 8.0, 250 mM NaCl, and 10% glycerol), and lysed by French press. The lysate was centrifuged at 12,000 rpm for 30 min. The supernatant was loaded onto a Ni-affinity column. The column was washed with extract buffer, and the protein was eluted with wash buffer (50 mM Tris, pH 8.0, 250 mM NaCl, 10% glycerol, and 100 mM imidazole). The protein was further purified on a Superdex 200 column.

Fig. S1

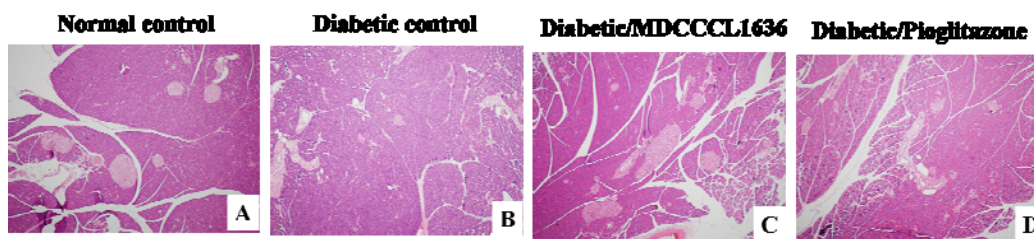


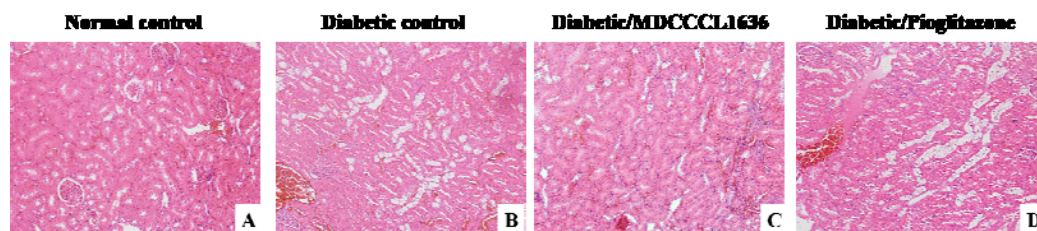
Fig.S2**Figure legends:**

Fig. S1. Histopathological changes in islets of pancreas of experimental groups. (A) Light micrograph of normal control rats showing normal β -cells in pancreatic islet (H & E, 40x), (B) HFD-STZ induced diabetic control rats showing pancreatic islet damage (H & E, 40x), (C) MDCCCL1636 at 7.5 mg/kg/day resulted in the reversal of pancreatic islet damage (H & E, 40x), (D) pioglitazone at 30 mg/kg/day resulted in the reversal of pancreatic islet damage (H & E, 40x).

Fig. S2. Histopathological changes in renal tubular epithelial cells of experimental groups. (A) Light micrograph of normal control rats showing normal renal tubular epithelial cells in pancreatic islet (H & E, 40x), (B) HFD-STZ induced diabetic control rats showing the fatty degeneration of renal tubular epithelial cells (H & E, 40x), (C) MDCCCL1636 at 7.5 mg/kg/day resulted in the reversal of the fatty degeneration of renal tubular epithelial cells (H & E, 40x), (D) pioglitazone at 30 mg/kg/day showing no effect on the fatty degeneration of renal tubular epithelial cells (H & E, 40x).