Introduction

Obesity, characterized as a state of low-level inflammation, is associated with insulin resistance [1, 2]. In adipose tissue of obese patients, recruited macrophages can activate the inflammatory response in neighboring adipocytes by releasing proinflammatory cytokines such as TNF-α and interleukin-6 (IL-6) [3–5]. These inflammatory molecules impair insulin receptor substrate-1 (IRS-1) function and downstream insulin/PI3K signaling to block glucose uptake, thereby leading to insulin resistance in adipocytes [6]. In addition, AMP-activated protein kinase (AMPK) regulates glucose and lipid homeostasis [7]. Emerging evidence demonstrates that anti-inflammatory action of AMPK is implicated in the reduction of insulin resistance [8, 9].

Berberine is a major isoquinoline alkaloid present in the Chinese herb *Rhizoma coptidis* and has a wide range of pharmacological actions [10, 11]. Recently, its antidiabetic action and related mechanisms have led to an increased interest among people. Due to the low bioavailability of berberine in vivo [12], a higher number of derivatives have been developed [13]. Nandinine is a benzylisoquinoline alkaloid derived from berberine, and this derivative contains an unsaturated heterocycle ring in the skeleton and a hydroxyl group instead of the methoxyl group in the C-9 position as compared to berberine. Although it was first isolated more than 80 years ago, few studies reporting its pharmacological effect have been published. In this study, we found that both berberine and nandinine attenuated insulin resistance by inhibiting inflammation in an AMPK-dependent manner in vivo and in vitro. A parallel investigation would present new insights on the development of derivatives from berberine for the management of diabetes and insulin resistance.
Results

In the present study, the production of TNF-α and IL-6 was increased after macrophage-derived conditioned medium (Mac-CM) treatment in 3T3-L1 adipocytes. Pretreatment with berberine and nandinine (Fig. 1) effectively reversed these phenomena (Fig. 2A–D). Adiponectin is a novel adipocyte-specific protein and an anti-inflammatory factor [14]. Mac-CM showed a reduction in adiponectin secretion. However, berberine and nandinine pretreatment obviously restored the production of adiponectin (Fig. 2E, F). Salicylate, a positive control, also showed activities similar to those of berberine and nandinine.

Dysregulated secretion of adipokines is the response to inflammation [15], and interleukin-β (IL-β) could modulate proinflammatory cytokine expression by activating nuclear factor-kappa B (NF-κB) in many types of cells [16–18]. In the present research, stimulation of Mac-CM enhanced the phosphorylation of IKKβ (Fig. 3A), the interaction of IκBα and IKKβ (Fig. 3B, C), and the translocation of activated NF-κB into the nucleus (Fig. 3D), whereas these phenomena were reversed by berberine, nandinine, and salicylate (Fig. 3).

IRS-1 is the main link between inflammation and insulin resistance [9]. After serine phosphorylation (S307) of IRS-1 is activated by IKKβ, the reduced tyrosine phosphorylation (Tyr) of IRS-1 will impair the insulin pathway to block glucose uptake in adipocytes [19, 20]. In this study, serine phosphorylation (S307) of IRS-1 was activated by Mac-CM-induced IKKβ (Fig. 4A). Meanwhile, the reduction of the tyrosine phosphorylation (Tyr) of IRS-1 (Fig. 4B) significantly decreased PI3K and Akt phosphorylation (Fig. 4C, D), and finally blocked insulin-stimulated glucose transporter-4 (GLUT-4) translocation and glucose uptake (Fig. 4E, F). Berberine, nandinine, and salicylate restored tyrosine phosphorylation to activate the insulin pathway, and stimulated GLUT-4 translocation as well as glucose uptake in the presence of insulin in adipocytes.

Activation of AMPK improves inflammation and insulin resistance in adipose tissues [21]. Results from our study showed that berberine, nandinine, and the AMPK activator 5-aminoimidazole-4-carboxamide ribonucleotide (AICAR) increased AMPK activity in both normal and Mac-CM-induced adipocytes (Fig. 5A, B). Furthermore, activated AMPK effectively inhibited IKKβ activation (Fig. 5C) to restore PI3K expression (Fig. 5D) and GLUT-4 translocation (Fig. 5E, F), which were impaired by Mac-CM. All of the beneficial effects from berberine, nandinine, and AICAR could be obviously abolished by AMPK inhibitor compound C pretreatment. These results suggested that AMPK was involved in the regulation of insulin resistance in adipocytes.

Finally, we investigated the influence of berberine and nandinine on glucose tolerance under inflammatory conditions in vivo. As an inflammatory challenge, Mac-CM treatment induced glucose intolerance in mice as evidenced by reduced glucose disposal. Similar to salicylate, the oral administration of berberine and nandinine (from 100 mg/kg to 200 mg/kg) obviously restored glucose disposal in the presence of Mac-CM (Fig. 6A). An increase of total glucose under AUC (AUC-G) was also reduced by both berberine and nandinine treatment (Fig. 6B). In addition, Mac-CM-induced insulin resistance was also effectively attenuated by berberine and nandinine treatment as evidenced by the decreased homeostasis model assessment of the insulin resistance (HOMA-IR) index (Fig. 6C).

Discussion

Inflammation is involved in the initiation and development of insulin resistance [22, 23]. Many inflammatory molecules such as TNF-α and free fatty acids could induce insulin resistance in many different cell types [13, 24]. AMPK is a metabolic sensor that helps maintain cellular energy homeostasis and it can exert significant anti-inflammatory effects in vitro [25–27]. In the present study, we showed that both berberine and nandinine reduced insulin resistance in an AMPK-dependent manner in adipocytes.

Mac-CM was derived from activated macrophages rich in inflammatory mediators, including TNF-α and IL-6, which can better mimic the pathology of insulin resistance in diabetes and obesity. In our study, Mac-CM stimulation obviously evoked inflammation in adipocytes, as evidenced by enhanced IKKβ phosphorylation and elevated levels of TNF-α and IL-6. However, these phenomena were reversed by berberine and nandinine. Besides, adiponectin is a novel adipocyte-specific protein and plays a role in the development of insulin resistance [28]. Berberine and nandinine also effectively increased adiponectin production, which was reduced by Mac-CM in adipocytes. Therefore, the beneficial effects of berberine and nandinine on the regulation of inflammatory cytokine well demonstrated their anti-inflammatory potency.

IRS-1 is a key control linking inflammation to insulin resistance [24]. Increased serine phosphorylation (S307) of IRS-1 could disturb Tyr of IRS-1, thereby leading to the impairment of the insulin signaling pathway [29]. Besides, IKKβ is required for activating serine phosphorylation of IRS-1 [30]. In the study, as expected, Mac-CM inhibited tyrosine phosphorylation of IRS-1 mediated by increased IKKβ to impair the PI3K/Akt pathway and finally blocked glucose uptake. Berberine and nandinine effectively attenuated insulin resistance by targeting IKKβ.

AMPK is a crucial regulator of energy metabolism and controls many characteristics of cellular stress resistance [31]. Accumulated evidence demonstrated that AMPK has a close relationship with insulin resistance [32]. Berberine reportedly suppresses proinflammatory responses through AMPK activation in macrophages [33]. Hence, we investigated whether berberine and nandinine regulated insulin signaling in an AMPK-dependent manner in 3T3-L1 adipocytes. In our study, both berberine and nandinine
increased AMPK activation in normal and Mac-CM-damaged adipocytes. Additionally, berberine, nandinine, and AICAR (AMPK activator) effectively reversed Mac-CM-induced phosphorylation of IKKa to restore PI3 K signaling, thereby leading to GLUT-4 translocation onto membranes and glucose uptake in the presence of insulin. These results suggested that berberine and nandinine ameliorated insulin resistance by activating AMPK.

To clarify the effects of berberine and nandinine on the actions of insulin in vivo, we detected the effects of them on glucose tolerance in mice. Glucose load stimulates insulin secretion from pancreatic islets, and then insulin, in turn, promotes glucose uptake [34]. In our study, the administration of berberine and nandinine effectively ameliorated glucose intolerance and induced the insulin sensitivity index in mice, thereby hinting that berberine and nandinine ameliorated glucose intolerance by restoring insulin sensitivity.

In conclusion, berberine and nandinine inhibited IKKa/NF-κB activation in adipocytes. Cells were pretreated with berberine (10 µM), nandinine (10 µM), or salicylate (5 mM) for 1 h and then incubated with Mac-CM for 24 h. Amounts of the cytokines in the culture media were quantified by ELISA (A, C, and E) and RT-PCR (B, D, and F). Data are presented as the mean ± SD of three independent experiments. *P < 0.05 vs. Mac-CM, **p < 0.01 vs. Mac-CM, ***p < 0.001 vs. Mac-CM. Mac-CM, group with Mac-CM treatment alone; NDN, nandinine; BBR, berberine; Sal, salicylate.

Materials and Methods

Chemicals and biochemicals

Lipopolysaccharide (Escherichia coli serotype 055:B5, LPS), compound C (purity ≥ 98% by HPLC analysis), protein A/G PLUS-Agarose, AICAR (purity ≥ 98% by HPLC analysis), and insulin were obtained from Sigma-Aldrich. Anti-IRS-1 (R301; BS1408); anti-Phospho-IRS-1 (Ser307; BS4725), anti-Akt (A444; BS1810), anti-Phospho-Akt (T308; BS4008), anti-IKKB (F182; BS1407), anti-Phospho-IKKB (Y199; BS4320), anti-Na+/K+-ATPase-α (-BS1436), GAPDH (AP0063), HRP-conjugated anti-rabbit, and anti-mouse IgG antibodies were purchased from Bioworld Technology. PY99 (sc-7020) was purchased from Santa Cruz Biotechnology. Histone H3, anti-AMPKα, anti-phospho-AMPKα (T172), and anti-GLUT-4 were purchased from Cell Signaling Technology. 2-NBD-glucose (N13195, lot: 873337) was purchased from Invitrogen. Sodium 205
Salicylate (purity ≥ 99.5%) was purchased from Tianjin Kemiou Chemical Agent Center. Fraction-PREP Cell Fractionation Kit was purchased from BioVision.

**Animals**

The male ICR mice (6–8 weeks of age), used all throughout the experiments, were supplied by the Laboratory Animal Center of Nanjing Qinglongshan. The care and treatment of these mice were maintained in accordance with the Provisions and General Recommendation of Chinese Experimental Animals Administration Legislation. The animal protocol was approved by Animal Ethics Committee of School of Chinese Materia Medica, China Pharmaceutical University (Permission Number: 2012–2–20) and the exact date of approval was February 25, 2012.

**Plant material**

Berberine (Kunming Fengshanjian Medical Research Co., Ltd.; purity > 98%) and nandinine (a gift from Dr. Zhang Jian, Department of Complex Prescription of TCM, China Pharmaceutical University; purity > 98%) were dissolved in DMSO, and the final concentration of DMSO in the medium or water was < 0.1%. The structures are shown in Fig. 1. Salicylate (purity ≥ 99.5%) was a product of Tianjin Kemiou Chemical Agent Center. Berberine and nandinine modulated the serine/tyrosine phosphorylation of IRS-1 and restored insulin-mediated GLUT-4 translocation and glucose uptake. Cells were pretreated with berberine (10 µM), nandinine (10 µM), or salicylate (5 mM) for 1 h and then incubated with Mac-CM for 30 min. Serine phosphorylation of IRS-1 was determined by Western blot analysis (A). After stimulation with Mac-CM, adipocytes were treated with insulin (100 nM) for 10 min. IRS-1 tyrosine, phosphorylation of Akt and PI3 K, and GLUT-4 protein expression on the membrane were determined by Western blot analysis (B–E), and glucose uptake was detected by flow cytometry (F). Results are expressed as the mean ± SD of three independent experiments. *P < 0.05 vs. Mac-CM, **p < 0.01 vs. Mac-CM, ***p < 0.001 vs. Mac-CM. Mac-CM, group with Mac-CM treatment alone; NDN, nandinine; BBR, berberine; Sal, salicylate. 

### Preparation of macrophages-derived conditioned medium

Mice were killed by cervical dislocation and injected intraperitoneally with 5 ml of PBS. After gentle abdominal massage, the PBS containing peritoneal macrophages was collected and cultured in 6-well plates (2 × 10⁶/well) for 2 h. Adherent cells were cultured in serum-free DMEM and stimulated by 5 µg/ml lipopolysaccharide (LPS) for 24 h. The supernatant was collected by centrifuge at 4 °C as a macrophage-derived medium (Mac-CM). TNF-α and IL-6 in the supernatant were measured with ELISA kits (R&D). When levels of TNF-α and IL-6 were greatly elevated compared with the control, the macrophages were activated. Mac-CM was filtered through a 0.22-µm filter and stored at −70°C.

### 3T3-L1 Cell culture and differentiation

The 3T3-L1 cell line, which is a cell line of preadipocytes, was obtained from the cell bank of the Chinese Academy of Sciences, and cultured in DMEM with 10% FBS at 37 °C in a 5% CO₂ atmosphere until cells grew to confluence. At 2 days after full confluence, cells were differentiated via incubation in DMEM containing 0.5 mM isobutylmethylxanthine (IBMX), 1 µm dexamethasone, 10 µg/ml insulin for 48 h, and then for 2 days in DMEM (10% FBS) containing 10 µg/ml insulin alone. Cells were maintained and refed every 2 days with DMEM, 25 mM glucose, and 10% FBS. After 8–12 days, over 80% of the cells exhibited the adipocyte phenotype with large lipid droplets in the cytoplasm.

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Determination of TNF-α, IL-6, and adiponectin production in adipocytes

Cells were plated in 24-well cell culture plates at a density of 2×10^5 cells/well and fasted for 24 h. Cells were pretreated with tested agents at determined concentrations for 1 h and then incubated in diluted Mac-CM (1:1, v/v) for 24 h. TNF-α, IL-6, and adiponectin in the medium were quantified with an ELISA kit (eBioscience).

AMPK and PI3K activity assay in adipocytes

Adipocytes (1×10^6 per well) were incubated for 4 h in serum-free DMEM pretreated with berberine (10 µM), nandinine (10 µM), compound C (20 µM), or AICAR (100 µM) for 1 h. Subsequently, these adipocytes were exposed to Mac-CM for another 30 min followed or not by a 10-min treatment of insulin (100 nM). After washing with ice-cold PBS, cells were lysed in cell lysis buffer (Tris–HCl 50 mM, pH 7.2, containing 1% sodium deoxycholate, 0.1% sodium dodecyl sulfate, 1% NP-40, NaCl 0.15 M, sodium orthovanadate 1 mM). The lysates were centrifuged at 13,000 g for 40 min at 4°C. The supernatant was collected for the AMPK activity assay or for the quantification of PIP3, the product of PI3K with ELISA Kits (Dizhao Biotech), respectively.

Western blotting

Cells were collected and lysed with the sample buffer (62.5 mM Tris–HCl, pH 6.8, 2% sodium dodecyl sulfate, 10% glycerol, 50 mM dithiothreitol, 0.1% bromphenol blue). Total proteins (30–50 µg) were resolved by 10% SDS-PAGE and then transferred electrophoretically to PVDF membranes. After blocking and washing, the membranes were detected by chemiluminescence (ECL system) and exposed by autoradiography, followed by immunoblotting with first and second antibodies.

2-(N-[7-Nitrobenz-2-oxa-1,3-diazol-4-yl] amino)-2-deoxyglucose assay

To analyze glucose uptake, the fluorescent glucose analog 2-(N-[7-nitrobenz-2-oxa-1,3-diazol-4-yl] amino)-2-deoxyglucose (2-NBDG), which allows for direct quantification of glucose incorporation in living cells by flow cytometry, was used [35]. After pretreatment, cells were incubated with the 2-NBDG probe (10 µM) for 40 min and insulin (100 nM) for 10 min. Then, cells were washed with PBS thrice, and fluorescence density was measured by a flow cytometry using a FACSCanto™ system (BD Biosciences).
Real-time PCR

Total RNA was extracted using TRIzol reagent. Quantitative real-time PCR (RT-PCR) was performed using SYBR Green PCR reagents. The specific primers for TNF-α were 5′-CTGTAGCCCACGTCGTA GC-3′ (forward) and 5′-CTGTAGCCCACGTCGTAGC-3′ (reverse). For IL-6, the primers were 5′-TCCAGTTGCCTTCTTG -3′ (forward) and 5′-GTGTAATTAAGCCTCCGACTTG -3′ (reverse). For adiponectin, the primers were 5′-TGTTCCTCTTAATCCTGCCCA -3′ (forward) and 5′-CCAACCTGCACAAGTTCCCTT-3′ (reverse). For GAPDH, the primers were 5′-CATGACCACAGTCCATGCATCAC -3′ (forward) and 5′-TGAGGTCCACCACCCTGTTGCTGT-3′ (reverse).

Briefly, cDNA (1 µL) from the RT reaction was added to 10 µL of the RT-PCR mixture containing 5 µL of Master Mix and 0.2 µM forward and reverse primers. The samples were incubated at 95 °C for 2 min, followed by 40 cycles at 95 °C for 15 s and at 60 °C for 1 min. The samples were assessed by 2−ΔΔCt relative quantitative analysis to determine the expression differences.

Immunoprecipitation

After determination of the protein concentrations, the cell extract was incubated with anti-IKKβ antibody (2 µg) for 2 h at 4 °C and then incubated with 20 µL of protein A/G plus-agarose beads overnight with constant shaking. Afterward, the beads were washed thrice with ice-cold radio immunoprecipitation assay buffer. The bound protein was extracted by adding 40 µL of 2 × SDS sample buffer and boiling for 5 min. The complexes were subjected to SDS-PAGE followed by Western blot.

Preparation of nuclear fractions

The nuclear proteins were isolated using the Fraction-PREP Cell Fractionation Kit according to the manufacturer’s instructions.

Glucose and insulin tolerance test in mice

Mice were fasted for 12 h and orally administrated berberine, nandinine, or salicylate at the given concentrations. After 1 h, mice were intraperitoneally injected with 0.2 ml of diluted macrophages-CM (1 : 2, v/v). After 30 min, mice were orally administered glucose solution (2 g/kg). Blood was collected from the orbital sinus at indicated times. Blood glucose was determined with a commercial kit based on the glucose oxidase peroxidase (GOD-POD) method. To determine insulin tolerance, mice were administered insulin (0.5 U/kg, s.c) 30 min after Mac-CM treatment. AUC (AUC-G) for blood glucose was calculated as follow: 0.5 × [Bg0 + Bg30]/2 + 0.5 × [B g30 + Bg60]/2 + 1 × [B g60 + B g120]/2 (Bg, blood glucose). To determine insulin sensitivity, blood was collected 30 min after oral glucose load, and blood glucose and insulin concentrations were measured simultaneously (ELISA kit for insulin assay). Homeostasis model assessment of the insulin resistance (HOMA-IR) index was calculated according to the following formula: HOMA-IR index = blood glucose (mmol/L) × blood insulin (mIU/L)/22.5 [36].
Statistical analysis

Data were expressed as means ± SD. Differences between groups were analyzed using Prism 5.0 (GraphPad Software Inc.), and statistical analysis was performed using one-way ANOVA, followed by the Student-Newman-Keuls test. A value of p < 0.05 was considered statistically significant.

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Conflict of Interest

The authors declare no conflict of interest.

References