Salidroside and isorhamnetin attenuate urotensin II-induced inflammatory response *in vivo* and *in vitro*: Involvement in regulating the RhoA/ROCK II pathway

CHENJING WANG¹, XIAODONG NAN², SHUYAN PEI¹, YU ZHAO¹, XIAOKUN WANG¹, SHIJIE MA¹ and GUOYAN MA¹

¹Department of Pharmacology, School of Basic Medical Sciences, Northwest Minzu University Health Science Center, Lanzhou, Gansu 730030; ²Intensive Care Unit, Gansu Provincial Corps Hospital of Chinese People's Armed Police Force, Lanzhou, Gansu 730050, P.R. China

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Abstract. Urotensin II (UII), a vital vasoconstrictor peptide, causes an inflammatory response in the pathogenesis of atherosclerosis. Previous studies have reported that the Ras homolog gene family, member A (RhoA)/Rho kinases (ROCK) pathway modulates the inflammatory response of the atherosclerotic process. However, to the best of our knowledge, whether the RhoA/ROCK pathway mediates the inflammatory effect of UII has not been previously elucidated. Salidroside and isorhamnetin are two early developed antioxidant Tibetan drugs, both displaying cardioprotective effects against atherosclerosis. Therefore, the aim of the present study was to investigate the protective effects of salidroside, isorhamnetin or combination of these two drugs on the UII-induced inflammatory response in vivo (rats) or in vitro [primary vascular smooth muscle cells (VSMCs)], as well as to examine the role of the RhoA/ROCK pathway in these processes. The levels of inflammatory markers were measured via ELISA. The mRNA and protein expression levels of RhoA and ROCK II were detected using reverse transcription-quantitative PCR assay and western blot analysis. It was demonstrated that salidroside, isorhamnetin and both in combination decreased the levels of the serum pro-inflammatory cytokines TNF- α and IL-1 β , as well as increased the levels of the anti-inflammatory cytokine IL-10 and macrophage migration inhibitory factor in rats with subacute infusion of UII and in the culture supernatant from primary VSMCs-exposed to UII. Moreover, salidroside,

Correspondence to: Dr Chenjing Wang, Department of Pharmacology, School of Basic Medical Sciences, Northwest Minzu University Health Science Center, 1 XibeiXincun, Chengguan, Lanzhou, Gansu 730030, P.R. China E-mail: wangcj01002@163.com

isorhamnetin and both in combination attenuated the mRNA and protein expression levels of RhoA and ROCK II *in vivo* and *in vitro*, at concentrations corresponding to human therapeutic blood plasma concentrations. Thus, these drugs could inhibit the RhoA/ROCK II pathway under UII conditions. The combination of salidroside and isorhamnetin did not display a stronger inhibitory effect on the inflammatory response and the RhoA/ROCK II pathway compared with salidroside and isorhamnetin in isolation. Collectively, the results indicated that salidroside, isorhamnetin and both in combination inhibited the RhoA/ROCK II pathway, which then attenuated the inflammatory response under UII-induced conditions, resulting in cardioprotection in atherosclerosis.

Introduction

Atherosclerosis is the leading cause of acute myocardial infarction and stroke, resulting in large global health and economic burdens (1). It has been reported that atherosclerosis is a chronic vascular wall-related inflammatory disease that occurs within the arterial wall (2). Inflammatory factors are mainly divided into three categories: i) Chemokines, including monocyte chemoattractant protein-1 (MCP-1), fractalkine/CX3CR1 and macrophage colony stimulating factor, whose effects are inhibited by macrophage migration inhibition factor (MIF); ii) pro-inflammatory factors, including C-reactive protein, IL-6, IL-1 and TNF-a; and iii) anti-inflammatory factors, including IL-10 and TGF- β (3). Pharmacological studies have reported that the imbalances between the pro-atherogenic inflammatory response and atheroprotective anti-inflammatory responses serve a key role in the initiation and progression of atherosclerosis (4). Previous in vivo and in vitro experimental data have revealed the key signaling pathways, such as programmed death ligand-1/programmed cell death protein 1 axis, brain-derived neurotrophic factor/tyrosine kinase B signaling pathway (5), Nod-like receptor protein 3 inflammasome/IL-1/IL-18/IL-6 pathway (6) and Toll-like receptor pathways (7), that mediate the inflammatory response, which may be research hotspots and provide potential preventive targets for atherosclerosis.

Key words: salidroside, isorhamnetin, urotensin II, inflammatory response, atherosclerosis, Ras homolog gene family, member A/Rho kinases II pathway

Therefore, the treatment of atherosclerosis using anti-inflammatory drugs may be an attractive strategy (8-10).

It has been reported that the Ras homolog gene family, member A (RhoA)/Rho kinases (ROCK) pathway is an important signal transduction system involved in cell proliferation, endothelial dysfunction, oxidative stress, inflammation, vascular remodeling and atherosclerosis (11-13). Previous studies have suggested that various multiple risk factors and pathological mediators of atherosclerosis can activate the RhoA/ROCK pathway to different degrees (14-16), and the inhibition of the RhoA/ROCK pathway could be a therapeutic potential target in the treatment of atherosclerosis (16). Furthermore, the regulatory effect of the RhoA/ROCK pathway on the inflammatory response of the atherosclerotic process has been confirmed by previous finding (17,18). The study from Shimada and Rajagopalan (19) revealed that ROCK mediates lysophosphatidic acid (an inflammatory mediator that is elevated in multiple inflammatory diseases)-induced IL-8 and MCP-1 production in human endothelial cells. The ROCK pathway also contributes to hyperglycemia-activated macrophages, which results in a pro-inflammatory phenotype and eventually contributes to atherosclerosis (17). However, the underlying potential RhoA/ROCK-regulated signaling pathways in inflammatory response under atherosclerosis remain to be elucidated.

Urotensin II (UII), a vasoactive cyclic peptide, and its high-affinity G-protein-coupled receptor UT are both highly expressed in the human cardiovascular system, and UII is involved in the development of cardiovascular homeostasis disease (20,21). Clinical and experimental studies have identified a positive correlation between increased UII levels and the development of atherosclerosis (22-24). UII can enhance the development of aortic atherosclerotic lesions and destabilizes atherosclerotic plaques (25). UII also exerts a pro-inflammatory effect on vascular wall cells in atherosclerosis (24,26). Previous studies have shown that the RhoA/ROCK pathway mediates UII-induced migration of endothelial progenitor cells and the formation of macrophage derived foam cells, suggesting that the RhoA/ROCK pathway may contribute to the UII-induced inflammatory response (27). However, to the best of our knowledge, there are no previous reports on the roles of the RhoA/ROCK pathway in UII-derived inflammatory effects.

Salidroside and isorhamnetin are two early used antioxidant Tibetan drugs, possessing a variety of biological activities such as anti-apoptosis, anti-oxidative stress and anti-inflammation effects (28,29). It has also been reported that salidroside and isorhamnetin exert a cardioprotective effect on the development of atherosclerosis, which is partly dependent on their anti-inflammatory ability (26,30,31). However, the anti-inflammatory or protective properties of salidroside and isorhamnetin against UII-derived inflammatory effects in atherosclerosis, as well as the underlying molecular mechanisms, are yet not fully understood.

Therefore, the present study aimed to investigate the role of the RhoA/ROCK II pathway in the UII-induced inflammatory response, as well as to identify the effects of salidroside and isorhamnetin treatment on the UII-induced inflammatory response and their potential mechanism *in vivo* and *in vitro*.

Materials and methods

Chemicals and reagents. UII (cat. no. U4753), salidroside (cat. no. 05410590) and isorhamnetin (cat. no. 17794) were purchased from Sigma-Aldrich (Merck KGaA). DMEM (cat. no. 11965092) and FBS (cat. no. 16140071) were purchased from Gibco (Thermo Fisher Scientific, Inc.). Anti-RhoA (cat. no. 2117s), anti-ROCK II (cat. no. 9029s) and anti-GAPDH (cat. no. 5174) antibodies were purchased from Cell Signaling Technology, Inc.

Rat modeling and animal treatment. All the experimental procedures were performed in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals (32), and were approved by the Experimental Animal Administration Committee of the School of Basic Medical Sciences, Northwest Minzu University Health Science Center (approval no. XBMZ-YX20130101; March 1, 2013). In total, 120 healthy male Wistar rats (weight, 180-200 g; age, 8 weeks) were provided by Jiangning Qinglongshan Animal Cultivation Farm and were housed under laboratory conditions (temperature $22\pm 2^{\circ}$ C with a relative humidity of 40-50% and natural light-dark cycle time of 12/12 h) with free access to food and water.

For the preparation of the rat model, rats were anaesthetized with pentobarbital sodium [60 mg/kg; intraperitoneal (i.p.)] and osmotic mini-pumps (Alzet Model 2006D; Durect Corporation) were loaded with either UII or saline alone (vehicle). The rats were randomly divided into eight groups: i) Normal control (saline alone, equal amount as the UII group, n=15); ii) UII group [rats were subcutaneously injected with UII (10 ng/kg/min) for 7 consecutive days, n=15]; ii) Salidroside (12 mg/kg) + UII group, (n=15); iv) Salidroside (24 mg/kg) + UII group, (n=15); v) Isorhamnetin (12 mg/kg) +UII group, (n=15); vi) Isorhamnetin (24 mg/kg) + UII group, (n=15); vii) Salidroside + isorhamnetin (Both, 12 mg/kg) + UII group, (n=15); and viii) Salidroside + isorhamnetin (Both, 24 mg/kg) + UII group, (n=15). The doses of salidroside and isorhamnetin in vivo and in vitro were chosen in accordance with the previous literatures (33-36).

Isolation and identification of primary vascular smooth muscle cells (VSMCs). A total of 8 adult male Wistar rats (age, 10 weeks; weight, 250-350 g) housed under the same aforementioned laboratory conditions were heparinized (4 IU/g; i.p.) and then euthanatized via pentobarbital sodium (100 mg/kg; i.p.) administration. The procedure for VSMCs isolation was performed in accordance with a previously described protocol (31). The aorta was immediately collected, placed into 75% (v/v) alcohol, dissected into sections (length, 3 cm) and subsequently placed in PBS. After removing fibroblasts that were present in the tunica externa (the external third of the vessel wall thickness) using forceps, the residual vessels were longitudinally cut and the tunica interna was scraped off, leaving the tunica media. Then, the tunica media was washed with DMEM, cut into sections (1 mm³) and maintained in DMEM containing 10% (v/v) FBS at 37°C in a Heraeus 5% CO₂ incubator (Thermo Fisher Scientific, Inc.). The morphology of cultured VSMCs was observed by phase contrast microscopy at passages 3-8 at room temperature. The identification of VSMCs

was performed using a Histostain-streptavidin-peroxidase kit (rabbit; cat. no. SP-0023; Beijing Biosynthesis Biotechnology Co., Ltd.) according to the manufacturer's protocol. Briefly, the third generation of VSMCs was inoculated into a 100-mm culture. When the cells grew to near fusion state, the cover glass was removed and the cells were fixed with 4% freshly prepared cooled neutral paraformaldehyde buffer at room temperature for 15 min and washed with PBS. Subsequently, after blocking non-specific binding sites with 5% BSA Blocking Reagent included in the aforementioned immunohistochemical kit at 37°C for 30 min, rabbit anti-a smooth muscle actin polyclonal antibodies (cat. no. bs-0189R; Beijing Biosynthesis Biotechnology Co., Ltd.) were incubated with the cells overnight at 4°C. Next, the slides were incubated with secondary antibodies for 1 h at room temperature and subsequently stained with a-smooth muscle actin antibody included in the immunohistochemical kit for 10 min at room temperature. The cells were visualized using a laser confocal microscope (Leica Microsystems GmbH; magnification, x400).

Cell culture and treatment. VSMCs were cultured in DMEM supplemented with 10% FBS in a humidified atmosphere with 5% CO₂ at 37°C. To investigate the effects of UII on VSMCs, cells were treated with different concentration of UII (10⁻⁹, 10⁻⁸, 10⁻⁷ and 10⁻⁶ mol/l) for 24 hat 37°C. To demonstrate the impacts of salidroside and isorhamnetin on VSMCs exposed to UII, cells were pretreated with salidroside (1, 3 or 10 μ M), isorhamnetin (1, 3 or 10 μ M) or both salidroside (3 or 10 μ M) and isorhamnetin (3 or 10 μ M) for 1 h, followed by treatment with UII (10⁻⁶ mol/l) for 24 hat 37°C.

Measurement of inflammatory markers in the culture supernatant using ELISA. VSMCs were seeded into a 6-well plate at a density of 1x10⁶ cells/ml. After incubation for 24 h as aforementioned, the culture supernatants were collected, centrifuged at 1,000 x g for 5 min at room temperature, and used to assess the levels of TNF- α (Nanjing Jiancheng Bioengineering Research Institute; cat. no. H052), IL-1β (Nanjing Jiancheng Bioengineering Research Institute; cat. no. H002), IL-10 (Nanjing Jiancheng Bioengineering Research Institute; cat. no. H009) and MIF (BioLegend® Legend Max TM Human Active MIF; cat. no. 438408; BioLegend, Inc.) with ELISA kits, according to manufacturer's instructions. The supernatants (500 μ M) were seeded into enzyme labelling 96-well plate and anti-TNF- α , IL-1 β , IL-10 and MIF antibodies (200 μ M) included in the kits were added, respectively. Following incubation for 2 h at room temperature, horseradish peroxidase (HPR)-labelled secondary antibody included in the kit (200 μ M) was added to for 1 h at room temperature. The absorbance value at 450 nm was determined using a Multiskan Microplate reader (Thermo Fisher Scientific, Inc.).

Reverse transcription-quantitative PCR (RT-qPCR) assay. The total RNA in each group was extracted with TRIzol[®] reagent (Invitrogen; Thermo Fisher Scientific, Inc.; cat. no. 15596018), and reverse-transcribed into cDNA using a PrimeScript RT Reagent kit (Takara Bio, Inc.; cat. no. RR037B). The RT conditions were 10 min at 25°C, 45 min at 48°C and a final step for 6 min at 95°C. RT-qPCR was performed using Platinum SYBR Green qPCR SuperMix (Invitrogen; Thermo Fisher

Scientific, Inc.; cat. no. 11744100) on an ABI Prism 7500 system (MP Biomedicals) according to the manufacturer's instructions. The amplification conditions were as follows: Initial denaturation at 95°C for 1 min, followed by 40 cycles of 95°C for 5 sec, 60°C for 15 sec and 72°C for 30 sec. GAPDH was used as the endogenous control. The mRNA expression levels of RhoA and ROCK II were normalized to GAPDH using the $2^{-\Delta\Delta Cq}$ equation (37). The primer sequences used were as follows: RhoA forward, 5'-TCGGAATGATGAGCA CACAA-3' and reverse, 5'-GCTTCACAAGATGAGGCAC-3'; ROCK II forward, 5'-CAGCAACTTTGACGACATTGAGG-3' and reverse, 5'-AGATTTGCACTTCTGTTCCAGC-3'; and GAPDH forward, 5'-ACGGCAAGTTCAACGGCACAG-3' and reverse, 5'-GACGCCAGTAGACTCCACGACA-3'.

Western blot analysis. After the indicated treatments, VSMCs were harvested, lysed in RIPA buffer (Beyotime Institute of Biotechnology; cat. no. P0013K) containing 1% (V/V) PMSF (Beyotime Institute of Biotechnology; cat. no. ST506) on ice for 30 min, and then centrifuged at 13,000 x g for 10 min at 4°C. The protein concentration was confirmed using a BCA Protein Assay kit (Beyotime Institute of Biotechnology; cat. no. P0012). Equal amounts of protein (30 μ g/lane) were separated on 12% SDS-PAGE, transferred onto PVDF membranes (EMD Millipore; cat. no. IPVH00010) and blocked with blocking buffer [0.1% Tween-20 in TBS (TBS-T) supplemented with 5% fat-free milk] for 2 h at room temperature. After washing with TBS-T buffer, the membranes were incubated with anti-RhoA, anti-ROCK II and anti-GAPDH antibodies (1:2,000) overnight at 4°C. GAPDH was used as a loading control. Then, the membranes were incubated with HRP-conjugated secondary antibody (1:5,000; cat. no. 7077; Cell Signaling Technology, Inc.) for 2 h at room temperature. The immunoreactive bands were visualized using a chemiluminescence imaging analysis system (cat. no. 32106; Pierce; Thermo Fisher Scientific, Inc.). The densities of protein expression were semi-quantified using Bio-Rad ChemiDoc XRS (version 4.3.0;Bio-Rad Laboratories, Inc.).

Statistical analysis. Data are presented as the mean \pm SD from \geq 3 different experiments. Data were analyzed using SPSS.18 software (SPSS, Inc.). Comparison among multiple relevant groups was performed using a one-way ANOVA followed by Bonferroni's multiple comparison test. P<0.05 was considered to indicate a statistically significant difference.

Results

Salidroside, isorhamnetin and both in combination decrease the levels of serum inflammatory cytokines after subacute infusion of UII in rats. The results of ELISA suggested that compared with the control group, after subacute infusion of UII (10 ng/kg/min) for 7 days, the levels of pro-inflammatory cytokines TNF- α (Fig. 1A) and IL-1 β (Fig. 1B) were significantly upregulated, while the levels of anti-inflammatory cytokine IL-10 (Fig. 1C) and MIF (Fig. 1D) were significantly downregulated in rat serum. However, compared with UII group, salidroside (12 and 24 mg/kg), isorhamnetin (24 mg/kg) and the combination of salidroside (12 or 24 mg/kg) and isorhamnetin (12 or 24 mg/kg) significantly attenuated the

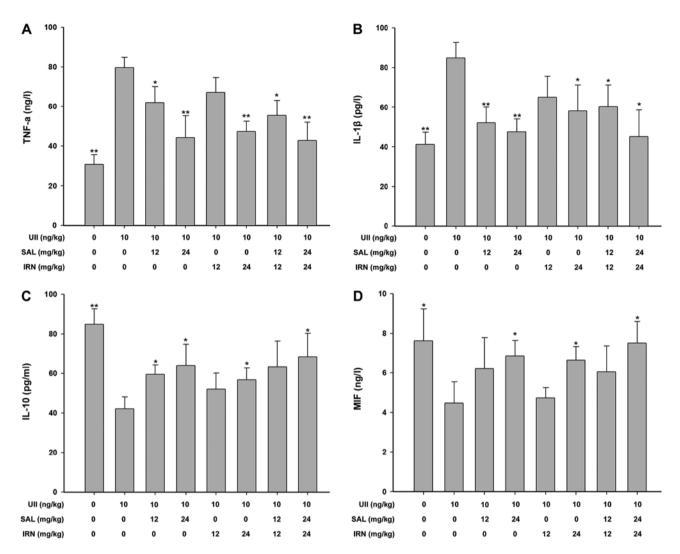


Figure 1. Effects of SAL and IRN on inflammatory cytokines in the serum of rats following UII treatment. Levels of (A) TNF- α , (B) IL-1 β , (C) IL-10 and (D) MIF in the serum of rats were analyzed via ELISA. Data are from \geq 3 independent experiments and presented as the mean \pm SD. *P<0.05, **P<0.01 vs. UII group. MIF, macrophage migration inhibitory factor; UII, Urotensin II; SAL, salidroside; IRN, isorhamnetin.

levels of TNF- α (Fig. 1A) and IL-1 β (Fig. 1B). Moreover, salidroside (12 and 24 mg/kg), isorhamnetin (24 mg/kg) and the combination of salidroside (24 mg/kg) and isorhamnetin (24 mg/kg) significantly increased the level of IL-10 compared with UII group (Fig. 1C). It was also identified that salidroside (24 mg/kg), isorhamnetin (24 mg/kg) and the combination of salidroside (24 mg/kg) and isorhamnetin (24 mg/kg) significantly promoted the level of MIF compared with UII group (Fig. 1D) in the serum of rats. The combination of salidroside (24 mg/kg) and isorhamnetin (24 mg/kg) did not exert a higher effect compared with salidroside or isorhamnetin alone (one-way ANOVA followed by Bonferroni's multiple comparison test). These results indicated that salidroside, isorhamnetin and both in combination protected rats against the UII-induced inflammatory response.

Salidroside, isorhamnetin and both in combination inhibit the RhoA/ROCK II pathway in the thoracic aorta of rats following subacute infusion of UII. Previous studies have reported that the RhoA/ROCK pathway can be activated by various atherosclerosis-related risk factors to different degrees, and may participate in the pathogenesis of atherosclerosis (11,38). Thus,

the present study investigated the effects of salidroside and isorhamnetin on the RhoA/ROCK II pathway in UII-treated rats. The results demonstrated that subacute infusion of UII significantly increased the mRNA expression levels of RhoA (Fig. 2A) and ROCK II (Fig. 2B) in thoracic aorta compared with control group. Furthermore, administration of salidroside (12 and 24 mg/kg), isorhamnetin (24 mg/kg) and the combination of salidroside (12 or 24 mg/kg) and isorhamnetin (12 or 24 mg/kg) significantly reduced the mRNA expression level of RhoA (Fig. 2A), while administration of salidroside (24 mg/kg), isorhamnetin (24 mg/kg) and the combination of salidroside (12 or 24 mg/kg) and the combination of salidroside (12 or 24 mg/kg) and isorhamnetin (12 or 24 mg/kg) significantly decreased the mRNA expression level of ROCK II (Fig. 2B), compared with the UII group.

Western blot analysis results further indicated that UII significantly increased the protein expression levels of RhoA (Fig. 2C) and ROCK II (Fig. 2D) in thoracic aorta of rats, compared with the control group. However, these impacts were mitigated by salidroside (12 or 24 mg/kg), isorhamnetin (24 mg/kg), both salidroside (12 mg/kg) and isorhamnetin (12 mg/kg), and both salidroside (24 mg/kg) and isorhamnetin (24 mg/kg) in combination. When the drug concentration

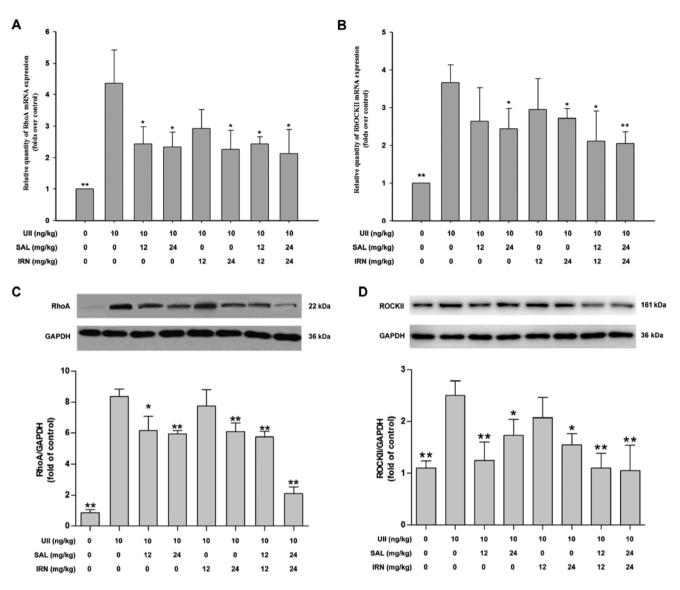


Figure 2. Effects of SAL and IRN on the RhoA/ROCK II pathway in the thoracic aorta of rats following subacute infusion of UII. mRNA expression levels of (A) RhoA and (B) ROCK II were measured via reverse transcription-quantitative PCR. Protein expression levels of (C) RhoA and (D) ROCK II were detected using western blot analysis. Data are from \geq 3 independent experiments and presented as the mean \pm SD. *P<0.05, **P<0.01 vs. UII group. RhoA, Ras homolog gene family, member A; ROCK, Rho kinases; UII, Urotensin II; SAL, salidroside; IRN, isorhamnetin.

was low, the combination of the two drugs could inhibit the RhoA/ROCK II pathway under UII-induced conditions. Moreover, the inhibitory effects of the combination of salidroside and isorhamnetin on the RhoA/ROCK II pathway were not more effective compared with salidroside or isorhamnetin alone (one-way ANOVA followed by Bonferroni's multiple comparison test). These results suggested that salidroside, isorhamnetin and both in combination inhibited the RhoA/ROCK II pathway, which may lead to the inhibition of UII-induced inflammation.

UII promotes the inflammatory response and the RhoA/ROCK II pathway in primary VSMCs. VSMCs were isolated from male Wistar rats. Under a phase contrast microscope, a small number of cells emerged from the surrounding of the tissue block (rat thoracic aorta) after 4-5 days of culture (Fig. 3A-1). After 10-12 days of culture, the cells in the local bundles were arranged in parallel, and some cells overlapped in multiple layers, demonstrating typical ups and downs of

'peaks' and 'valleys' (Fig. 3A-2). In the third generation of VSMCs, actin immunocytochemical staining identified that >99% of the cells were positive; the cytoplasm was brownish yellow and the nucleus was not stained (Fig. 3A-3). These results indicated that VSMCs were successfully extracted.

Subsequently, the effects of UII on the inflammatory response in VSMCs were detected via ELISA. With the increased concentration of UII (10⁻⁹, 10⁻⁸, 10⁻⁷ and 10⁻⁶ mol/l) stimulation for 24 h, the levels of the anti-inflammatory cyto-kines IL-10 (Fig. 3B) and MIF (Fig. 3C) were significantly decreased. In addition, RT-qPCR results demonstrated that UII (10⁻⁸, 10⁻⁷ and 10⁻⁶ mol/l) treatment for 24 h induced significant increases in the mRNA expression levels of RhoA (Fig. 3D) and ROCK II (Fig. 3E) in VSMCs. Western blot analysis (Fig. 3F) results identified that UII (10⁻⁸, 10⁻⁷ and 10⁻⁶ mol/l) stimulation for 24 h significantly increased the expression of RhoA (Fig. 3G), and UII (10⁻⁷ and 10⁻⁶ mol/l) significantly increased the expression of ROCK II (Fig. 3H) in VSMCs. As UII (10⁻⁶ mol/l) treatment for 24 h significantly upregulated the

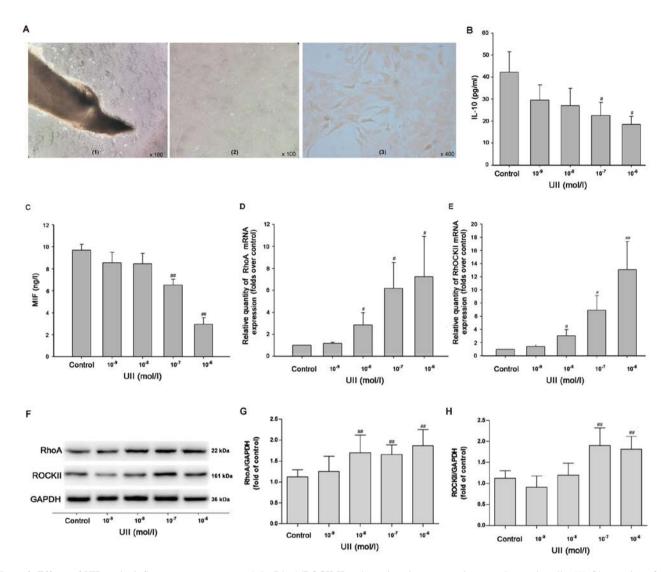


Figure 3. Effects of UII on the inflammatory response and the RhoA/ROCK II pathway in primary vascular smooth muscle cells. (A) Observation of cell morphology. (A-1) After 4-5 days of culture, cell morphology was observed under the phase contrast microscope (magnification, x100). (A-2) After 10-12 days of culture, cell morphology was observed under an invert microscope (magnification, x100). (A-3) α -actin immunocytochemical staining (magnification, x400). Inflammatory markers (B) IL-10 and (C) MIF in the culture supernatant were measured via ELISA. mRNA expression levels of (D) RhoA and (E) ROCK II were detected using reverse transcription-quantitative PCR. Protein expression levels of (G) RhoA and (H) ROCK II were detected using (F) western blot analysis. Data are from \geq 3 independent experiments and presented as the mean \pm SD. [#]P<0.05, ^{##}P<0.01 vs. control. RhoA, Ras homolog gene family, member A; ROCK, Rho kinases; UII, Urotensin II; MIF, macrophage migration inhibitory factor; SAL, salidroside; IRN, isorhamnetin.

protein expression levels of RhoA and ROCK II in VSMCs, this treatment regimen was selected for subsequent experiments. Taken together, it was suggested that UII stimulation promoted the inflammatory response by enhancing the RhoA/ROCK II pathway in primary VSMCs.

Salidroside, isorhamnetin and both in combination attenuate the UII-induced inflammatory response and inhibit the RhoA/ROCK II pathway in VSMCs. Next, in order to demonstrate the inhibitory effects of salidroside and isorhamnetin on inflammatory responses in VSMCs stimulated with UII, VSMCs were pretreated with salidroside, isorhamnetin or both in combination for 1 h followed by treatment with UII (10⁻⁶ mol/l) for 24 h. Compared with the control group. UII treatment significantly increased the TNF- α (Fig. 4A) and IL-1 β (Fig. 4B) levels, and decreased the IL-10 level (Fig. 4C). Moreover, compared with the UII treatment group, pretreatment with salidroside (3 and 10 μ M), isorhamnetin (3 and 10 μ M) and both in combination reduced TNF- α level (Fig. 4A), while pretreatment with salidroside (10 μ M), isorhamnetin (10 μ M) and both in combination reduced IL-1 β level (Fig. 4B) in UII-treated VSMCs. It was also found that pretreatment with salidroside (3 and 10 μ M), isorhamnetin (1, 3 and 10 μ M) and both in combination increased the IL-10 level (Fig. 4C) in UII-treated VSMCs. The low concentration of salidroside (1 μ M) had no effect on the level of inflammatory-related factors in UII-treated VSMCs.

The results suggested that pretreatment with salidroside (3 and 10 μ M), isorhamnetin (10 μ M) and a combination of salidroside and isorhamnetin attenuated the UII-induced increases in the mRNA expression level of RhoA (Fig. 4D). In addition, pretreatment with salidroside (3 and 10 μ M), isorhamnetin (3 and 10 μ M) and both in combination attenuated the UII-induced the increases in the mRNA expression levels of ROCK I (Fig. 4E) and ROCK II (Fig. 4F). Western blotting results (Fig. 4G) also identified that salidroside (10 μ M), isorhamnetin (10 μ M) and

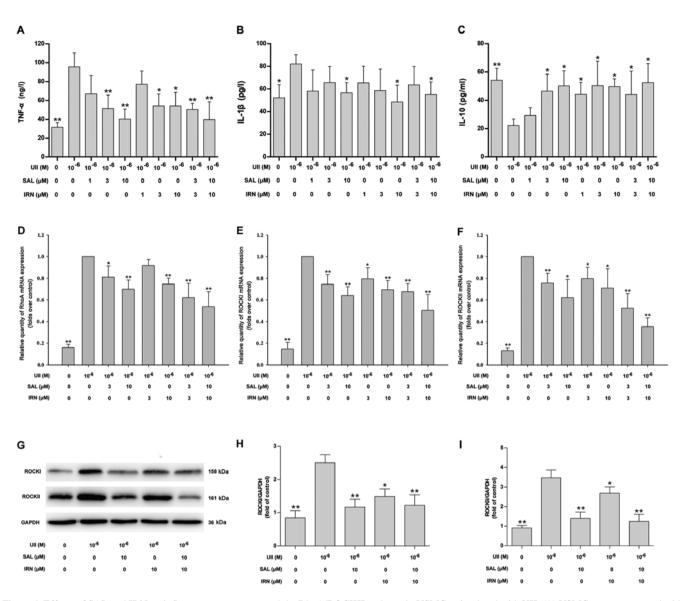


Figure 4. Effects of SAL and IRN on inflammatory response and the RhoA/ROCKII pathway in VSMCs stimulated with UII. (A) VSMCs were pretreated with SAL or IRN (1, 3 and 10 μ M) alone or SAL and IRN (3 and 10 μ M) in combination for 1 h followed by treatment with UII (10⁻⁶ mol/l) for 24 h. Quantitative analysis of (A) TNF- α , (B) IL-1 β and (C) IL-10 levels via ELISA. mRNA expression levels of (D) RhoA, (E) ROCK I and (F) ROCK II were detected via reverse transcription-quantitative PCR. (G) Western blot analysis of the protein expression levels of (H) ROCK I and (I) ROCK II were measured using. Data are from \geq 3 independent experiments and presented as the mean \pm SD. *P<0.05, **P<0.01 vs. UII group. MIF, macrophage migration inhibitory factor; UII, Urotensin II; RhoA, Ras homolog gene family, member A; ROCK, Rho kinases; SAL, salidroside; IRN, isorhamnetin; VSMCs, vascular smooth muscle cells.

both in combination reversed UII-induced upregulation of ROCK I (Fig. 4H) and ROCK II (Fig. 4I) protein expression levels in VSMCs. There were no statistically significant differences between the combination of drugs and with salidroside or isorhamnetin alone (one-way ANOVA followed by Bonferroni's multiple comparison test). Collectively, it was indicated that salidroside, isorhamnetin and both in combination mitigated the UII-induced inflammatory response in VSMCs partly by inhibiting the RhoA/ROCK pathway.

Discussion

The present findings indicated that UII stimulation resulted in an inflammatory response along with an increase in the RhoA/ROCK pathway activation *in vivo* and *in vitro*. It was also demonstrated that salidroside, isorhamnetin and both in combination could elicit inhibitory effects on the UII-induced inflammatory response, at least partially by attenuating the RhoA/ROCK pathway.

Previous studies have reported that UII exerts vasculopathic and vasculoprotective effects, and contributes to the pathogenesis of atherosclerosis (26,39,40). UII promotes VSMCs proliferation by activating reactive oxygen species (ROS) and MAPK signaling pathways, which results in vascular remodeling (41). UII also participates in macrophage activation via UII receptor/ROS/Akt pathways in RAW264.7 macrophages (26). Furthermore, UII can promote NAD(P)H oxidase-induced reactive oxygen production from a variety of inflammatory cells, resulting in activated NF- κ B (42,43), which is the most important downstream event involved in the signal transduction pathway of various inflammatory factors in the process of vascular injury and atherosclerosis (44-46). These results provide evidence for the inflammatory effect of UII in cardiovascular disease. Notably, to the best of our knowledge, the present results were the first to demonstrate that UII stimulation results in an inflammatory response, as shown by the increases in the levels of pro-inflammatory cytokines (TNF- α and IL-1 β) and the decreases in the levels of anti-inflammatory cytokines (IL-10 and MIF) in the serum of rats and the culture supernatant of VSMCs. Therefore, it was indicated that inflammatory factors induced by UII led to an inflammatory response that is involved in the pathological process of atherosclerosis.

RhoA and its downstream effector ROCK, which exists in two isoforms, ROCK1 and ROCK2, serve significant roles in multiple cellular processes, such as proliferation, apoptosis and migration (47). Abnormal activation of the RhoA/ROCK pathway has been reported to be involved in various types of diseases including diabetes, osteoarthritis, cardiovascular and cerebrovascular diseases (11-13). In recent years, the regulatory effect of the RhoA/ROCK pathway on the inflammatory process of atherosclerosis has been revealed. For instance, upregulation of the RhoA/ROCK signaling cascade has been identified in atherosclerosis (11). Moreover, RhoA-mediated NF-kB signaling pathways lead to vascular endothelial dysfunction in diabetes (48). ROCK pathways also contribute to hyperglycemia-activated macrophages, which result in a more pro-inflammatory phenotype and eventually lead to atherosclerosis (17). Previous studies have shown that ROCK I is predominantly increased in the process of macrophage adherence, and ROCK1-deficiency decreases atherosclerosis in bone marrow-derived cells (49), indicating that ROCK I serves an important role in the development of atherosclerosis. However, whether ROCK II inhibition could also be beneficial in attenuating atherosclerosis remain to be investigated. In line with these previous results, the present findings suggested that UII stimulation promoted the RhoA/ROCK pathway in rats and VSMCs, implying the involvement of the RhoA/ROCK II pathway in UII-induced inflammatory response.

It has been revealed that salidroside, an early antioxidant Tibetan medicine, has anti-inflammatory effects in atherosclerosis (50,51). Li *et al* (50) reported that salidroside can decrease the generation of inflammatory cytokines, such as IL-6, IL-1 β and MCP-1, in TNF- α -induced cardiac microvascular endothelial cells, alleviating vascular inflammation and atherosclerosis. Another previous study revealed that salidroside attenuated endothelial cellular senescence via reducing the expression of inflammatory cytokines, thus mitigating the pathogenesis of atherosclerosis (51). In addition, isorhamnetin, a flavonoid monomer extracted from seabuckthorn fruit, has been shown to possess anti-cancer, anti-oxidant, anti-inflammatory and anti-atherosclerotic activities (34,52,53). However, there are few studies on the anti-inflammatory and anti-atherosclerotic effects of salidroside, isorhamnetin and both in combination.

To the best of our knowledge, the present study was the first to demonstrate that salidroside, isorhamnetin and both in combination decreased TNF- α and IL-1 β levels, and increased IL-10 level in the serum of UII-treated rats and in the culture supernatant of UII-stimulated VSMCs, at concentrations corresponding to human therapeutic blood plasma concentrations, thus eliminating the UII-induced inflammatory response. High concentrations of salidroside and isorhamnetin both reversed the UII-induced inflammatory response *in vivo* and *in vitro*. However, the inhibitory effects of low concentration

of salidroside and isorhamnetin on inflammatory response were inconsistent. The combination of low and high concentrations of salidroside and isorhamnetin both eliminated the UII-induced inflammatory response *in vivo* and *in vitro*. Furthermore, the anti-inflammatory effect of the combination of salidroside and isorhamnetin was not significantly different compared with the single drug alone during UII conditions in atherosclerosis.

Subsequently, based on the role of the RhoA/ROCK pathway in UII-induced inflammatory response, the present study further investigated the effects of salidroside, isorhamnetin and both in combination on the RhoA/ROCK pathway. It was found that salidroside, isorhamnetin and both in combination inhibited the RhoA/ROCK pathway in the thoracic aorta of rats following subacute infusion of UII and in UII-stimulated VSMCs. Consistent with the *in vivo* results, the inhibitory effects of low concentrations of salidroside and isorhamnetin on the RhoA/ROCK pathway were inconsistent, while the high concentrations of salidroside and isorhamnetin both reversed UII-induced the enhancement of the RhoA/ROCK pathway in vivo and in vitro. The combination of low and high concentrations of salidroside and isorhamnetin both attenuated the RhoA/ROCK pathway under UII in vivo and in vitro. There was no significant difference between salidroside and isorhamnetin in isolation and in combination. Thus, the results suggested that, to a certain extent, the RhoA/ROCK pathway contributed to the anti-inflammatory effects of salidroside, isorhamnetin and both in combination under UII simulation in atherosclerosis.

However, there are a few limitations to the present study. First, gain-of-function and loss-of function experiments were not used to investigate the role of the RhoA/ROCK pathway in UII-induced inflammatory or the anti-inflammatory effects of salidroside, isorhamnetin and both in combination. This will require further examination in future studies. In addition, the potential mechanism of salidroside, isorhamnetin and both in combination in the inhibition of the RhoA/ROCK pathway under UII has not yet been elucidated. The present study also did not examine the comparison of the two drugs with currently, widely-used anti-atherosclerosis drugs and other reported anti-inflammatory drugs in atherosclerosis, which may be interesting to evaluate in the subsequent experiment plan.

In conclusion, the present findings indicated that UII stimulation resulted in an inflammatory response, accompanied by the enhancement of the RhoA/ROCK pathway in vivo and in vitro. In addition, to the best of our knowledge, the present results provided the first evidence that salidroside, isorhamnetin and both in combination attenuated the UII-induced inflammatory response, which was partly dependent on inhibition of the RhoA/ROCK pathway. Moreover, there is no significant difference between salidroside and isorhamnetin, both in isolation or in combination. The present study may provide a novel theoretical basis for the separate used of the two drugs or their combination in the treatment against atherosclerosis. Furthermore, the present results may provide additional theoretical guidance for the clinical combined use of the Tibetan medicines salidroside and isorhamnetin in the prevention against atherosclerosis-related cardiovascular diseases.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

CW and XN participated in the design of the study and conducted the experiment. CW, SP and YZ wrote the manuscript and analyzed data. XW, SM and GM collaborated in the analysis and interpretation of data. All authors read and approved the final manuscript.

Ethics approval and consent to participate

All animal experimental procedures were performed in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals, and were approved by the Experimental Animal Administration Committee of the School of Basic Medical Sciences, Northwest Minzu University Health Science Center (approval no. XBMZ-YX20130101; March 1, 2013).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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