

Original Article

Paeonol protects rat vascular endothelial cells from ox-LDL-induced injury *in vitro* via downregulating microRNA-21 expression and TNF- α release

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Aim: Paeonol (2'-hydroxy-4'-methoxyacetophenone) from *Cortex moutan* root is a potential therapeutic agent for atherosclerosis. This study sought to investigate the mechanisms underlying anti-inflammatory effects of paeonol in rat vascular endothelial cells (VECs) *in vitro*.

Methods: VECs were isolated from rat thoracic aortas. The cells were pretreated with paeonol for 24 h, and then stimulated with ox-LDL for another 24 h. The expression of microRNA-21 (miR-21) and PTEN in VECs was analyzed using qRT-PCR. The expression of PTEN protein was detected by Western blotting. TNF- α release by VECs was measured by ELISA.

Results: Ox-LDL treatment inhibited VEC growth in dose- and time-dependent manners (the value of IC₅₀ was about 20 mg/L at 24 h). Furthermore, ox-LDL (20 mg/L) significantly increased miR-21 expression and inhibited the expression of PTEN, one of downstream target genes of miR-21 in VECs. In addition, ox-LDL (20 mg/L) significantly increased the release of TNF- α from VECs. Pretreatment with paeonol increased the survival rate of ox-LDL-treated VECs in dose- and time-dependent manners. Moreover, paeonol (120 μ mol/L) prevented ox-LDL-induced increases in miR-21 expression and TNF- α release, and ox-LDL-induced inhibition in PTEN expression. A dual-luciferase reporter assay showed that miR-21 bound directly to PTEN's 3'-UTR, thus inhibiting PTEN expression. In ox-LDL treated VECs, transfection with a miR-21 mimic significantly increased miR-21 expression and inhibited PTEN expression, and attenuated the protective effects of paeonol pretreatment, whereas transfection with an miR-21 inhibitor significantly decreased miR-21 expression and increased PTEN expression, thus enhanced the protective effects of paeonol pretreatment.

Conclusion: miR-21 is an important target of paeonol for its protective effects against ox-LDL-induced VEC injury, which may play critical roles in development of atherosclerosis.

Keywords: paeonol; atherosclerosis; microRNA-21; PTEN; vascular endothelial cells; oxidized low density lipoprotein; inflammatory reaction

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Introduction

Atherosclerosis (AS) is a leading cause of morbidity and mortality worldwide and involves multiple genes and other factors^[1]. Dysfunction of and injury to vascular endothelial cells (VECs) are the driving forces in the initiation and development of AS, whereas ox-LDL is a well-recognized risk factor for VEC damage^[2]. Paeonol (2'-hydroxy-4'-methoxyacetophenone, C₉H₁₀O₃) is one of the main active compounds in *Cortex moutan* root, which has been used in traditional Chinese medicine for its anti-inflammatory, anti-thrombotic and antioxidant properties^[3, 4] (Figure 1). A large number of studies have demonstrated that paeonol is effective for preventing and

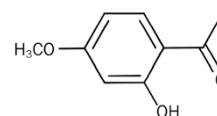


Figure 1. The chemical structure of paeonol from *Cortex Moutan*.

treating of AS in experimental models by modulating inflammatory reactions during the initiation and development of AS. For example, paeonol inhibits tumor necrosis factor (TNF)- α -induced vascular cell adhesion molecule-1 (VCAM-1) expression, which reduces monocyte adhesion to VECs^[5]. Moreover, paeonol suppresses lipopolysaccharide-induced inflammatory cytokine release from macrophage cells and protects mice from lethal endotoxin shock^[6]. Although the anti-inflamma-

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tory effect of paeonol is well recognized, the exact mechanisms are still unclear.

MicroRNAs (miRNAs) constitute a broad class of short noncoding RNAs that have been extensively studied in the context of cancer. Recently, increasing evidence has indicated that miRNAs are involved in the development of AS^[7]. miRNAs can modulate the expression of various cytokines or inflammatory molecules by binding to the complementary target sites in the 3' untranslated regions (3'UTR) of mRNA sequences and inducing mRNA degradation or translational repression. MicroRNA-21 (miR-21) is selectively expressed in vascular cells and modulates proliferation, invasion, apoptosis and inflammation^[8]. In the peripheral arterial system, miR-21 is upregulated after acute vascular injury, and knockdown of miR-21 has been shown to reduce neointima formation^[9]. Therefore, we were interested in determining whether miR-21 is involved in modulating the therapeutic effect of paeonol on AS, as well as the downstream target of miR-21 in VECs.

In this study, we investigated the mechanisms underlying the protective effect of paeonol in a VEC injury model. We assessed the expression of miR-21 and its downstream target genes, such as PTEN, in injured VECs in the presence or absence of paeonol treatment.

Materials and methods

Animals

Healthy male Sprague-Dawley rats weighing 150–170 g were provided by the Experimental Animal Center of Anhui Medical University (Hefei, China). The animal experiments were approved by the local institutional Animal Care and Use Committee.

Chemicals

Paeonol (98% purity) was obtained from Baicao Plants Biotech Co, Ltd (Anhui, China). The cell culture materials and fetal bovine serum (FBS) were obtained from GIBCO BRL Co, Ltd (Gaithersburg, USA). Ox-LDL (YB-002-1) was purchased from Yiyuan Biotechnologies Co, Ltd (China). HiPerFect transfection reagent, miR-21 inhibitor and mimic, MiRNeasy Mini Kit (Cat.217004) and MiScript PCR Starter Kit (Cat.218193) were purchased from Qiagen (Germany). The dual-luciferase reporter system (Cat.E1910) was purchased from Promega (USA). The BCA Protein Assay kit was obtained from Shanghai Haoran Bio Technologies Co, Ltd (China). Antibodies against β -actin and PTEN were purchased from Bioworld Technology Co, Ltd (USA).

Isolation and culture of VECs

VEC's were isolated from rat thoracic aortas as previously described^[10]. In short, thoracic aortas isolated from Sprague-Dawley rats were turned over to expose the luminal surface, under sterile conditions. The aortas were then tied tightly at both ends and digested with 0.2% collagenase I. After incubation at 37°C in 5% CO₂ for 1 h, the aortas were washed with medium and cut into pieces; both ends were discarded. The pieces were placed, lumen side down, onto collagen-coated

flasks containing DMEM supplemented with 15% heat-inactivated FBS and penicillin/streptomycin and then incubated at 37°C in 5% CO₂. Once the cells had formed a monolayer, the explants were removed, and the cells were subcultured. The cells were identified by their endothelial cell-specific, cobblestone-like morphology, and the purity of the VECs was determined by immunocytochemical staining for Factor VIII. In all experiments, the cells used were passage 3–5.

MTT assay

VEC's were seeded in 96-well-plates (1×10⁵ cells/mL) and incubated in DMEM with 15% FBS at 37°C in 5% CO₂. After treatment with paeonol and ox-LDL, the cells were incubated for 4 h with fresh medium containing 20 μ L MTT (5 mg/mL). The medium was then removed, and 150 μ L DMSO was added to each well to dissolve the precipitate. The absorbance of MTT at 490 nm was detected using a microplate reader (Spectra MaxMze, VT, USA).

Transfection with a miR-21 mimic and inhibitor

VEC's were seeded in 6-well plates and incubated in DMEM with 15% FBS at 37°C and 5% CO₂. After dilution of 0.6 μ L miR-21 mimic (20 μ mol/L stock) or 6 μ L miR-21 inhibitor (20 μ mol/L stock) in 400 μ L culture medium without serum, 12 μ L HiPerFect transfection reagent was added to the diluted miR-21 mimic/inhibitor and mixed by vortexing to form transfection complexes. The complexes were added drop-wise to the VEC's medium, and the plate was gently agitated. The VEC's were incubated with the transfection complexes for 24 h, and the medium was then changed as required.

RNA extraction and qRT-PCR assay

Total RNA was extracted using QIAzol Lysis Reagent for both miR-21 and PTEN mRNA analyses. For detection of gene expression, qRT-PCR was performed using Quantitect SYBR Green PCR Kits according to the manufacturer's protocol. Relative expression was evaluated using the comparative CT method and normalized to the expression of U6 small RNA.

Dual-luciferase reporter assay

The sequence for miR-21 was obtained from PubMed, and the matched sites are shown below. Luciferase reporter plasmids containing a wild type or mutant 3'-UTR of PTEN were constructed to verify the putative binding site for miR-21. The wild type or mutant reporter plasmid was cotransfected into VEC's, along with the miR-21 mimic. After 36 h, luciferase activity was measured using the dual-luciferase assay system. The firefly luciferase activity of each sample was normalized to the *Renilla reniformis* luciferase activity.

3'AGUUGUAGUCAGACUAUUCGAU 5' rno-miR-21

— _ | _ | | : _ _ : _ : | | | | _

5'CAAUGAUUUACUCAGUAAGCUU 3'PTEN WT

5'CAAUGAUUUACUCACAUUCGAU 3'PTEN MUT

Western blotting

VECs were lysed in SDS sample buffer, and total protein was extracted from the lysates. Protein concentrations were determined using a BCA Protein Assay kit. The proteins were separated by 10% SDS-polyacrylamide gel electrophoresis and transferred onto nitrocellulose membranes, which were blocked for 3 h at room temperature with phosphate buffered saline-Tween containing 5% nonfat milk. Then, membranes were incubated with antibodies against β -actin and PTEN. Immunoreactive proteins were visualized by enhanced chemiluminescence detection of secondary antibodies against mouse or rabbit antigens conjugated to horseradish peroxidase.

Statistical analysis

All data are presented as means \pm SD and were analyzed using SPSS 17.0 (USA). Student's *t* test was used for statistical comparisons between two groups, and one-way ANOVA was used for multiple groups. A difference was considered statistically significant when $P < 0.05$.

Results

Establishment of injured VECs by exposure to ox-LDL

VECs were seeded in 96-well plates with DMEM containing 10% FBS at 1×10^5 cells/mL and cultured at 37°C in 5% CO₂. After achieving 80% confluence, the cells were treated with various concentrations of ox-LDL (5, 10, 20, 40, or 80 mg/L) for different periods (12, 24, or 48 h), and cells were then counted using the MTT assay (Figure S1, Figure 2). Ox-LDL treatment inhibited VEC growth in a dose- and time-dependent manner; the growth inhibition rate was close to 50% when the cells were stimulated with 20 mg/L ox-LDL for 24 h. These conditions were used as the optimal stimulation concentration and time in subsequent experiments.

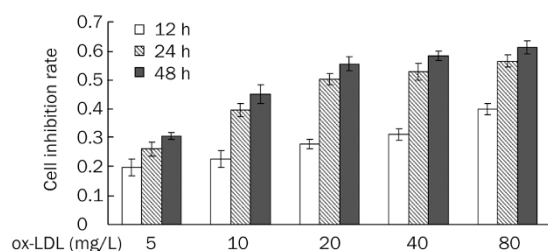


Figure 2. Inhibitory effects of ox-LDL on VECs viability rate were analyzed by MTT assay using a microplate reader at 490 nm. The data were expressed as mean \pm SD. $n=6$. Ox-LDL (5, 10, 20, 40, and 80 mg/L) stimulated VECs for 12, 24, and 48 h, respectively.

MiR-21 expression was stimulated in ox-LDL-injured VECs

The expression of miR-21 in ox-LDL-injured VECs was measured using qRT-PCR. We observed that miR-21 expression was significantly stimulated by treatment with ox-LDL (20 mg/L) compared with the control group ($P < 0.05$, Figure 3), suggesting that miR-21 is involved in ox-LDL-induced VEC damage. Additionally, transfection with a miR-21 mimic

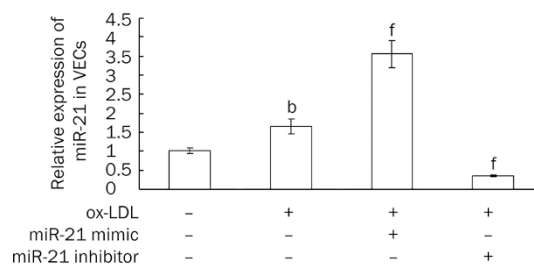


Figure 3. MiR-21 was specifically expressed in ox-LDL-injured VECs. The VECs were transfected with miR-21 mimic or inhibitor, then the cells were treated with ox-LDL (20 mg/L) for 24 h. All the data came from SYBR Green PCR assay and were expressed as mean \pm SD. $n=3$. ^b $P < 0.05$ vs control group. ^f $P < 0.01$ vs ox-LDL group.

significantly enhanced miR-21 expression in VECs, whereas transfection with a miR-21 inhibitor significantly suppressed miR-21 expression ($P < 0.01$) (Figure 3).

PTEN is a target gene of miR-21 in VECs

To determine whether miR-21 directly targets PTEN in VECs, we constructed luciferase reporter plasmids containing a wild type or mutant 3'-UTR for PTEN. Cotransfection of the wild type reporter plasmid with the miR-21 mimic significantly decreased the relative luciferase activity compared with a miR-21 mimic alone ($P < 0.01$), whereas cotransfection with mutant type 3'-UTR plasmid elicited significantly higher luciferase activity than that seen in the wild type group ($P < 0.01$) (Figure 4A). In addition, we found that ox-LDL treatment significant reduced the expression levels of PTEN mRNA and protein (Figure 4B and 4C) and that these changes were reversed by treatment with the miR-21 inhibitor. In contrast, a much more significant reduction in expression of PTEN mRNA and protein was observed in the miR-21 mimic-transfected group (Figure 4B and 4C). These findings suggest that PTEN is a downstream target gene of miR-21 in VECs and that miR-21 suppresses PTEN by directly binding to its 3'-UTR.

Paeonol suppresses miR-21 expression in ox-LDL treated VECs

First, we pretreated VECs with paeonol at various concentrations (15, 30, 60, 120, 240, or 480 μ mol/L) for various lengths of time (6, 12, 24, or 48 h). We next treated the cells with 20 mg/L ox-LDL for another 24 h. After treatment, cell survival was tested using the MTT assay. It was observed that the cells in paeonol-treated group had a higher survival rate than the ox-LDL group; this effect was dose and time dependent (Figure 5). An optimal condition for paeonol treatment in VECs (120 μ mol/L for 24 h) was established and used in subsequent experiments.

To test whether the protective effect of paeonol is associated with changes in miR-21 expression in VECs, the expression of miR-21 after paeonol and ox-LDL treatment was measured by qRT-PCR. Paeonol treatment significantly decreased ox-LDL-induced miR-21 expression in a dose-dependent manner ($P < 0.01$, Figure 6), suggesting that the protective effect of

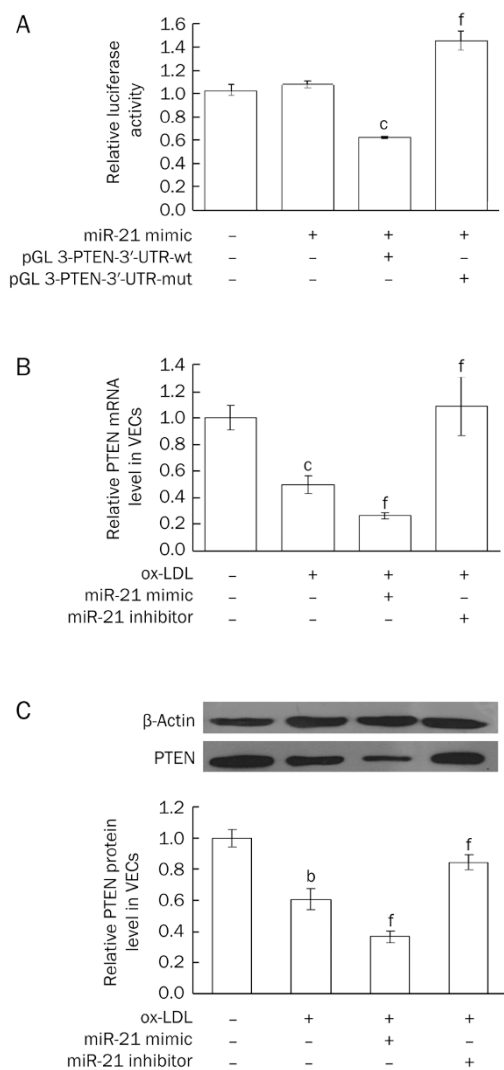


Figure 4. PTEN was a target gene of miR-21 and miR-21 expression was inversely correlated with PTEN expression in ox-LDL stimulated VECs. The data were expressed as mean \pm SD. $n=3$. (A) The wild type or mutant reporter plasmid was cotransfected into VECs with miR-21 mimic. $^{\circ}P<0.01$ vs mimic group. $^{\circ}P<0.01$ vs 3'UTR wild type group. (B) Effects of miR-21 mimic and inhibitor on PTEN mRNA expression were measured by SYBR Green PCR assay. $^{\circ}P<0.01$ vs control group. $^{\circ}P<0.01$ vs ox-LDL group. (C) Total protein was harvested from cultured VECs and analyzed by Western blotting for PTEN. $^{\circ}P<0.05$, $^{\circ}P<0.01$ vs control group. $^{\circ}P<0.01$ vs ox-LDL group.

paeonol on VECs was associated with modulation of miR-21 expression.

Paeonol prevents downregulation of PTEN in ox-LDL treated VECs

To test whether the protective effect of paeonol in ox-LDL-treated VECs was associated with regulation of PTEN, we treated VECs with 120 μ mol/L of paeonol for 24 h, followed by 24 h of ox-LDL treatment. Ox-LDL treatment significantly suppressed the expression of PTEN at both the mRNA and protein level, and these effects were prevented by pretreat-

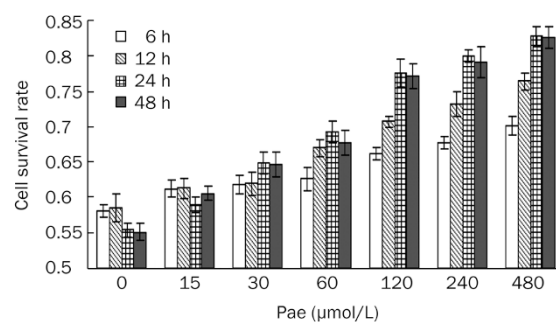


Figure 5. Effects of paeonol on ox-LDL-injured VECs survival rate were analyzed by MTT assay using a microplate reader at 490 nm. The data were expressed as mean \pm SD. $n=6$. VECs were pretreated with different concentrations of paeonol (15, 30, 60, 120, 240, and 480 μ mol/L) for 6, 12, 24, and 48 h, respectively, then treated with 20 mg/L ox-LDL for another 24 h.

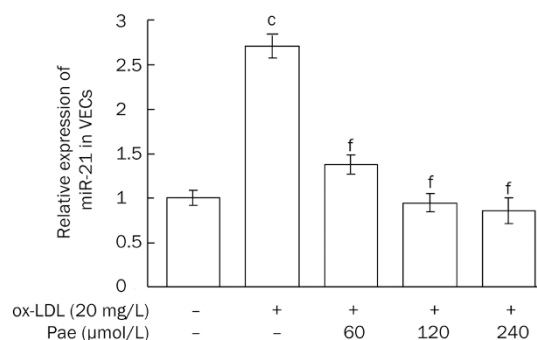


Figure 6. Downregulative effects of paeonol on miR-21 expression were analyzed by SYBR Green PCR assay. VECs were pretreated with varying concentrations of paeonol (60, 120, and 240 μ mol/L) for 24 h and stimulated by ox-LDL (20 mg/L) for another 24 h, then miR-21 expression was analyzed. Data were expressed as mean \pm SD. $n=3$. $^{\circ}P<0.01$ vs control group. $^{\circ}P<0.01$ vs ox-LDL group.

ment with paeonol (Figure 7A and 7B). In addition, to further characterize the relationship between miR-21 and PTEN in paeonol-treated VECs, the mRNA and protein expression levels of PTEN were assayed in ox-LDL-treated VECs pretreated with paeonol in the presence of the miR-21 mimic or miR-21 inhibitor. We found that the miR-21 mimic rescued the effect of paeonol on the protein expression of PTEN, whereas addition of a miR-21 inhibitor enhanced paeonol-induced increases in PTEN at both the mRNA and protein levels (Figure 7A and 7B). These results suggested that regulation of PTEN by miR-21 might be involved in the protective effect of paeonol in ox-LDL-induced VEC injury.

Paeonol treatment reduced TNF- α release by ox-LDL treated VECs

To test whether the effect of paeonol on ox-LDL induced injury in VECs is associated with suppression of TNF- α production, the levels of TNF- α were measured using an ELISA.

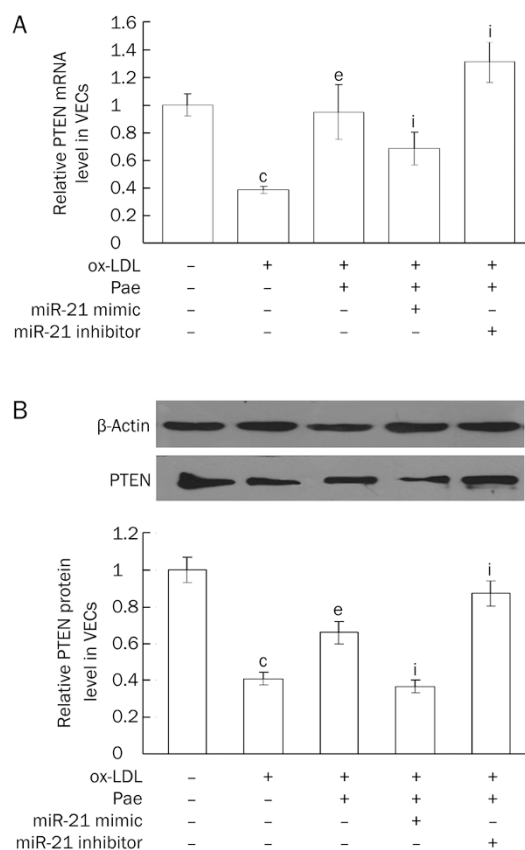


Figure 7. (A) Effects of paeonol on PTEN mRNA level was detected by SYBR Green PCR assay. (B) Effect of paeonol on PTEN protein level was analyzed by Western blotting. MiR-21 mimic or inhibitor was transfected into VECs, then VECs were pretreated with paeonol (120 μ mol/L) for 24 h and stimulated with ox-LDL (20 mg/L) for another 24 h. Data were expressed as mean \pm SD. $n=3$. $^{\circ}P<0.01$ vs control group. $^{\circ}P<0.01$ vs ox-LDL group. $^iP<0.01$ vs Pae group.

We observed that ox-LDL treatment significantly stimulated TNF- α release in VECs and that this effect was prevented by paeonol pretreatment. Moreover, the effect of paeonol on TNF- α release was partially reversed or significantly enhanced by transfection with miR-21 mimic or inhibitor, respectively (Figure 8). These results suggest that the protective effect of paeonol against ox-LDL induced injury in VECs might be associated with miR-21 mediated suppression of TNF- α release.

Discussion

Paeonol is a proven anti-atherosclerotic herbal extract with a potent anti-inflammatory activity^[11]. Our previous investigations have indicated that its anti-atherosclerotic effects are associated with protection of endothelial cells from injury^[5,11]. However, the underlying mechanisms are still unclear. miRNAs suppress target transcription and translation by binding to the 3'-untranslated regions (3'UTRs) of target mRNAs. These small endogenous silencers can regulate diverse biological processes, such as proliferation, apoptosis,

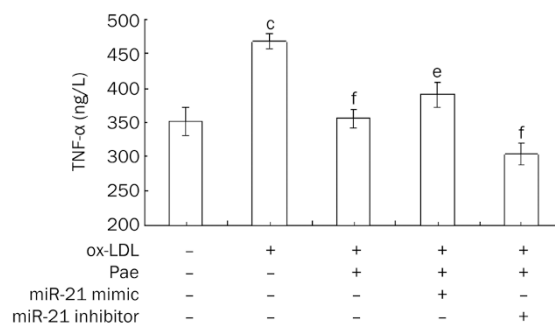


Figure 8. Paeonol inhibited TNF- α expression in ox-LDL induced VECs. MiR-21 mimic or inhibitor was transfected into VECs, then the VECs were pretreated with paeonol (120 μ mol/L) for 24 h and stimulated with ox-LDL (20 mg/L) for another 24 h. TNF- α expression was detected by ELISA assay kit. Data were expressed as mean \pm SD. $n=3$. $^{\circ}P<0.01$ vs control group. $^{\circ}P<0.05$, $^iP<0.01$ vs ox-LDL group.

angiogenesis, cholesterol metabolism and inflammation^[7]. Increasing evidence indicates that miRNAs play important roles in the development of AS^[12]. miR-21 is highly expressed in vascular endothelial cells^[13], cardiomyocytes^[14], cardiac fibroblasts^[15] and therefore may be critical in the development of cardiovascular disease. In this study, we found that the protective effect of paeonol against ox-LDL VEC injury was at least partially due to its effects on miR-21 expression. Moreover, we observed that pretreatment with paeonol also reversed the suppression of PTEN expression induced by ox-LDL. Interestingly, studies have demonstrated that PTEN is a downstream target gene of miR-21 in cancers^[16-19], because its 3'-UTR contains 16 base sites that are complementary to the miR-21 sequence^[1, 20]. However, it is unknown whether paeonol-mediated protection of VECs against ox-LDL injury involves miR-21-dependent modulation of PTEN.

In the current study, we found not only that paeonol pretreatment antagonized the suppression of PTEN induced by ox-LDL treatment but also that the expression of PTEN is regulated by miR-21. Additionally, our dual-luciferase reporter assay indicated that miR-21 binds directly to PTEN's 3'UTR, inhibiting PTEN expression. As a result, we suggest that miR-21-mediated regulation of PTEN contributes to the protective effects of paeonol.

TNF- α is an important inflammatory factor that is critical in mediating inflammatory diseases such as AS and arthritis^[21]. In the current study, we found that ox-LDL-induced TNF- α release was prevented by paeonol pretreatment and that transfection with a miR-21 mimic or inhibitor, when combined with paeonol, partially reversed or increased TNF- α release, respectively. Therefore, it is possible that the paeonol's effects on TNF- α release are miR-21-dependent and may contribute to the protection of VECs against ox-LDL-induced cell injury. Nevertheless, the relationship between PTEN and TNF- α release needs further investigation.

Taken these findings together, the current study indicates that the protective effect of paeonol against ox-LDL-induced injury of VECs is at least partially miR-21-dependent. This

miRNA modulates the downstream target gene, PTEN, as well as TNF- α release. Above all, this study validates miR-21 as a biomarker for vascular inflammation and clarifies the molecular mechanism by which paeonol aids in the treatment of vascular diseases.

Acknowledgements

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Author contribution

Ya-rong LIU wrote the paper; Ya-rong LIU and Jun-jun CHEN performed the experiments and analyzed the data; and Min DAI designed the study and supervised the project.

Supplementary materials

The supplementary figures are available on the web site of APS.

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