



Different Effects of Endothelial **Extracellular Vesicles and** LPS-Induced Endothelial Extracellular Vesicles on Vascular Smooth Muscle Cells: Role of **Curcumin and Its Derivatives**

Debiao Xiang^{1†}, Yamei Li^{2†}, Yuling Cao², Ying Huang³, Lili Zhou², Xiulian Lin², Yong Qiao¹, Xin Li¹ and Duanfang Liao^{2*}

¹ The Third Hospital of Changsha, Changsha, China, ² Key Laboratory for Quality Evaluation of Bulk Herbs of Hunan Province,

Hunan University of Chinese Medicine, Changsha, China, ³ The First Affiliated Hospital of Hunan University of Chinese

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*Correspondence:

Duanfang Liao dfliao@hnucm.edu.cn

[†]These authors have contributed equally to this work

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Background: During the progression of atherosclerosis (AS), the vascular endothelial and smooth muscle cells are reciprocally regulated by extracellular vesicles (EVs). EVs have different effects on pathological and physiological processes due to the different cargoes contained in EVs.

Purpose: To study the effects of endothelial cells-derived EVs on normal and inflammatory conditions. To investigate the effects of curcumin and curcumin derivatives (Nicotinic-curcumin) on endothelial EVs.

Methods: EVs were isolated from human umbilical vein endothelial cells (HUVECs) by ultracentrifugation. To examined the effect of normal and LPS-induced endothelial cells-derived EVs on the proliferation of human aortic smooth muscle cells (HASMCs), the CCK-8 assay was performed. Transwell and wound healing assays were conducted to assess cell migration. The effects of EVs on lipid accumulation following treatment with oxidized low-density lipoprotein (Ox-LDL) were evaluated with the oil red O staining assay and HPLC. The number of EVs was calculated using the nanoparticle tracking analysis (NTA) and BCA. The expression levels of Rab27a and Rab27b that regulate the EVs secretion were measured by Western blotting assay. The differential expression of miRNAs in endothelial EVs and LPS-induced endothelial EVs was analyzed using miRNA-Sequencing (miRNA-Seq) and RT-PCR.

Results: Treatment with endothelial EVs reduced the proliferation and migration of HASMCs as well as lipid accumulation in HASMCs. However, treatment with LPS-induced endothelial EVs did not inhibit the migration of HASMCs or lipid accumulation, instead it promoted the proliferation of HASMCs. Treatment with the two types of EVs induced differential expression of several miRNAs, including miR-92a-3p, miR-126-5p, miR-125a-3p, miR-143-3p, etc. Moreover, 1 µg/mL LPS induction greatly increased secretion of endothelial EVs. Treatment with curcumin and nicotinic-curcumin reduced endothelial EVs secretion, possibly by inhibiting inflammation.

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Conclusion: Endothelial EVs may confer beneficial effects on atherosclerosis by regulating vascular smooth muscle cell (VSMCs), whereas pro-inflammatory factors may disrupt this effect.

Keywords: atherosclerosis, endothelial EVs, vascular smooth muscle cell, inflammation, curcumin

INTRODUCTION

Atherosclerosis (AS) is a progressive disease characterized by abnormal accumulation of lipids and immune cells in the vascular wall. AS has been implicated in the development of several cardiovascular and cerebrovascular diseases such as myocardial infarction, angina pectoris, heart failure and stroke (1, 2). Meanwhile, AS is caused by diabetes, hypertension, dyslipidemia, and inflammation (3, 4). Atherosclerotic lesions are focal, asymmetrical thickening of the arterial intima, consisting of numerous cell types, connective tissue components and lipids (5). Vascular endothelial and smooth muscle cells as well as macrophages participate in the formation of atherosclerotic plaques (6). Several studies suggest that endothelial dysfunction, which occurs at the initial stage of atherosclerosis, is involved in the formation, progression and occurrence of complications related to atherosclerotic plaques (7, 8). Inflammation is an important marker for endothelial dysfunction. Specifically, several pro-inflammatory factors in serum such as oxidized lowdensity lipoprotein (ox-LDL), tumor necrosis factor- α (TNF- α), interleukin-1 (IL-1 β) etc, activates endothelial cells (9, 10) to release several cytokines, which recruit circulating monocytes and leukocytes and as well-induces abnormal proliferation and migration of vascular smooth muscle cells. Vascular smooth muscle cells are one of the main components of the fibrous cap of atherosclerotic plaques. Inflammation increases permeability of vascular intima, allowing entry of large amount of lipids in to the arterial lumen (11). Vascular smooth muscle cells proliferate and migrate to the intima, ingest large amounts of lipids and transform into foam subtypes that ultimately form the atherosclerotic plaque.

Extracellular vesicles (EVs) are small (30 to 1,000 nm) bilayered structures secreted by numerous cells (12, 13). Given the numerous biomolecules in EVs including miRNAs, DNA and proteins, the vesicles are used as specific diagnostic biomarkers for several diseases (14–16). Numerous studies also show that EVs are potential therapeutic targets and drug delivery systems (17, 18). Almost all types of cells secrete EVs, which are abundantly present in stable form in various tissues and body fluids. EVs mediate communication between cells, and through their loaded molecules such as miRNA, proteins, and cytokines, they regulate functions of the target cells (12, 19).

The interaction between endothelial and vascular smooth muscle cells is essential in the development of cardiovascular disease. Intriguingly, EVs enhance the efficiency of such interactions. Particularly, EVs signals can regulate vascular tone of Smooth muscle cells (SMCs) (20). Excessive secretion of EVs in VSMCs under endoplasmic reticulum stress induces apoptosis of ECs and infiltration of inflammatory cytokines (21). MiRNAs are one of main molecules in EVs. They regulate posttranscription gene expression by binding to the 3[']-untranslated region (UTR) of specific target mRNA sequences. This modulates protein expression by either promoting mRNA degradation or by blocking mRNA translation. miR-126-3p derived from endothelial EVs modulates proliferation of vascular smooth muscle cell (VSMCs) and disrupts neointima formation by inhibiting secretion of Low-density lipoprotein receptor-related protein 6 (LRP6) (22). Uptake of miR-143/145 containing EVs provides atheroprotection in smooth muscle cells. MiR-125a-3p inhibits proliferation and migration of VSMCs and the development of vascular stenosis by targeting MAPK1 (23). Over-expression of miR-155 promotes the formation of foam cells and aggravates atherosclerosis by targeting SOCS1 via STAT3 and NF-kB signaling pathway (24). It has been demonstrated that miRNA-10b regulates proliferation of VSMCs via the Akt pathway, specifically by targeting TIP30 (25). EV derived miRNAs are thought to participate in the development of cardiovascular disease. Functions RNAs have been extensively studied. Expression of miRNA in endothelial EVs under physiological and disease conditions suggests these structures play critical roles in such disease conditions. In this paper, we evaluated the effect of regulating the expression of miRNAs in endothelial derived EVs.

Curcumin isolated from the rhizome of *Curcuma longa* L. possess antioxidant, anti-inflammatory, antimicrobial, anticarcinogenic and cardiovascular protective properties (26). It has been evidenced that curcumin reduces the incidence of atherosclerosis due to its hypolipidemic effects, antioxidant and anti-inflammatory activity (27). Moreover, vast targets inevitably bring vast unnecessary side effects and low bioavailability of curcumin also restrict its clinical application. Nicotinic-curcumin (NC) is derivative of curcumin and niacin, synthesized through esterification (**Scheme 1**). We aimed to improve bioavailability of curcumin and combine the effects of curcumin with the lipid-lowering effects of niacin in atherosclerosis treatment. At present, we have found NC treatment remarkably increased efflux of cholesterol and reduced accumulation of ox-LDL in THP-1 cells (28).

MATERIALS AND METHODS

Materials

Niacin and curcumin were purchased from Sinopharm Chemical Reagents Co., Ltd (Beijing, China). ECM medium and human aortic smooth muscle cells (HASMCs) were purchased from ScienCell Co., Ltd (San Diego, USA). ELISA kits were purchased from Multi Sciences Co., Ltd (Shanghai, China). Other materials included antibodies from Abcam (London, England), Oil red O



stain from Solarbio Science & Technology Co., Ltd. (Beijing, China) and Cell culture flasks and transwell plates from Corning (New York, USA).

Cell Culture

Human umbilical vein endothelial cells (HUVECs) were isolated from the human fresh umbilical cord based on the CD31 marker protein, using an immunofluorescence microscope They were cultured at 37°C in ECM medium (ScienCell, USA) in a humid incubator under 5% CO₂ and 95% relative humidity (RH). Human aortic smooth muscle cells (HASMCs) were cultured in Dulbecco's modified Eagle medium (DMEM) (HyClone, Australia) supplemented with 10% fetal bovine serum (FBS), 1% of penicillin (100 U/mL) and streptomycin (100 μ g/mL). The antibiotics were first diluted in dimethyl sulfoxide (DMSO) and culture medium to obtain the desired concentrations.

Isolation and Identification of Endothelial EVs

HUVECs were cultured in Endothelial Cell Medium (ECM) (free of EVs). The supernatant was collected and centrifuged sequentially at 300 g for 10 min, 2,000 g for 10 min and 10,000 g for 30 min) under 4 °C. The supernatant was further centrifuged at 100,000 g for 70 min at 4 °C. The precipitate was resuspended in phosphate buffered solution (PBS) and centrifuged at 100,000 g for 70 min at 4 °C. Protein concentration in PBS ($100 \sim 200 \mu L$) was quantified using the bicinchoninic acid (BCA) method. The EVs were stored at -80 °C.

Uptake of Endothelial EVs by VSMCs

The cell culture medium was replaced with 2 mL of Exoperfect exo-CD63-GFP tracer virus medium, supplemented with 5 μ g/mL polybrene. After co-incubation for 8 h, the cells were observed under the microscope before replacing the EV-free serum-based media. Incubation was continued for another 48 h before adding 2 μ g/mL puromycin for screening CD63-GFP stably transfected cells. Normal cells served as controls. The culture medium was replaced every 2 days. Stable tranformants in the virus infected group served beyond death of all cells in the control group. The concentration of puromycin in the stable cells was halved in the subsequent culture. The transfected cells were

cultured for 48 h in EVs-free medium and centrifuged to obtain the supernatant. The EVs were labeled using CD63-GFP before isolation using ultra-high centrifugation. VSMCs were cultured in medium containing CD63-GFP-labeled EVs for 12 h and 24 h. Cells were then fixed for 15 min using 4% paraformaldehyde before being sealed with DAPI tablets. The cells were observed and photographed using a laser confocal microscope.

EVs Treatment and Co-culture of HUVECs and VSMCs

Purified endothelial EVs were diluted using DMEM medium and thereafter filtered using a bacterial membrane $(0.22 \,\mu\text{m})$ sieve. The VSMCs were then treated for 24 h using $20 \,\mu\text{g/mL}$ of endothelial EVs. A scratch was made in a monolayer of VSMCs co-cultured with HUVECs for 6 and 12 h to assess the migration ability of the VSMCs.

HUVECs were cultured for 8 h in the upper chamber of a 24-well-transwell plate ($0.4 \mu m$ aperture) in ECM medium supplemented with $1 \mu g/mL$ of LPS, whereas HASMCs were cultured separately on a 24-well-plate. After activation using $1 \mu g/mL$ lipopolysaccharide (LPS), the HUVECs and HASMCs were co-cultured for 24 h before adding 2.5 nmol/mL GW4869 (inhibits EVs release) in an upper chamber of a 24-well-transwell plate. The proliferation of VSMCs in a lower chamber was assessing using the CCK-8 assay after 24 h culture. VSMCs in a lower chamber were reseeded onto upper chamber of separate transwell plates ($8 \mu m$ aperture), incubated for 12 h before staining using 0.25% crystal violet. The stained cells were observed, photographed and counted under the inverted microscope.

Determination of Free Cholesterol in Cells

The HASMCs were treated for 24 h with 80 µg/mL ox-LDL to established lipid accumulation model. The HASMCs were then centrifuged at 1,000 r/min for 10 min under 4°C, resuspended in 500 µL 0.9% NaCl solution and repeatedly (3 ~ 4 times) freeze-thawed at between -80° C and room temperature. The cell membranes of the HASMCs were ruptured using an ultrasonic instrument. The supernatant was centrifuged at 12,000 r/min for 10 min under 4°C, before the protein concentration in xyz was determined using the BCA method. To determine



cholesterol concentration, equal volumes of the supernatant and KOH-ethanol were first mixed vigorously for 5 min to a uniform solution and saponified ultrasonically for 1 h in a water bath at 48°C. Thereafter, 6% trichloroacetic acid (45 μ L) was added to denature the protein before adding Iso-0volumetric cholesterol extractor (n-butane: isopropanol at the ratio of 60:40). After mixing and centrifugation at 4°C (12,500 r/min, 10 min), the upper organic phase was collected, dried in nitrogen and dissolved in 40 μ L acetonitrile. The concentration of free cholesterol was measured using HPLC (29). The aqueous phase was processed three times to ensure maximum cholesterol extraction.

RESULTS

Identification of Endothelial EVs

EVs are small bi-layer structures (30-1,000 nm) that carry numerous biomolecules such proteins, lipids, DNA and RNA.

Information between cells is transmitted through the constituent proteins and nucleic acids. Endothelial cells secrete numerous biomolecules, and given their wide distribution, they interact with innumerable cells. Endothelial cells secrete cytokines, which are carried via EVs to downstream target cells such as vascular smooth muscle cells, macrophages, etc. Averagely, Endothelial EVs are 150 nm wide (**Figure 1A**). They contain CD63, TSG101, and Calnexin surface markers (**Figure 1B**). Transmission electron microscopy revealed that the endothelial EVs are generally spherical (**Figure 1C**).

Effects of Endothelial EVs on the Proliferation, Migration, and Accumulation of Lipid in HASMCs

Uptake of Endothelial EVs by HASMCs

CD63-GFP labeled EVs in culture supernatant of transduced HUVECs (Figure 2B) were isolated using an ultra-high centrifugation. Laser scanning confocal microscope revealed



FIGURE 2 | Uptake of CD63-GFP-EVs by vascular smooth muscle cells. (A) Transmission electron microscopy showing successful uptake of EVs by HASMCs; (B) Uptake of CD63-GFP by HUVECs.



FIGURE 3 | (A) The effect of EVs derived from LPS activated HUVECs (EV.LPS) on proliferation of HASNCS. The proliferation of HASMCs was assessed using CCK-8 assay; **(B)** Effect of HUVECs culture on proliferation of HASMCs in the presence of LPS. The assessment was performed using the CCK-8 assay (Data represents mean \pm SD of 6 independent experiments, #p < 0.05 and ##p < 0.01, respectively. *p < 0.05, and **p < 0.01, respectively). #: Compared with Con, *: compared with Model.

successful uptake of CD63-GFP-EVs by HASMCs after 12 h and 24 h of incubation (**Figure 2A**).

The Effects of Endothelial EVs on the Proliferation and Migration of HASMCs

Endothelial EVs substantially inhibited proliferation of HASMCs (P < 0.01). In contrast, EVs derived from LPS stimulated HUVECs (EV.LPS) promoted proliferation of HASMCs (P < 0.05; **Figure 3A**). In addition, LPS activated HUVECs exerted a similar effect on HASMCs (P < 0.05). However, culture with LPS stimulated HUVECs promoted the proliferation of HASMCs (**Figure 3B**).

EVs secreted by non-activated HUVECs inhibited migration of HASMCs (P < 0.05). Wound scratch assay revealed that

LPS-activated HUVECs had no effect on the migration of HASMCs (**Figures 4A,B**). Further analyses revealed that LPS activated HUVECs disrupted migration of HASMCs. A culture of non-activated HUVECs substantially modulated the migration of HASMCs (P < 0.05). However, GW4869 treatment inhibited the effect of endothelial EVs on proliferation and migration of HASMCs (**Figures 4C,D**).

The Effects of Endothelial EVs on Accumulation of Lipids in HASMCs

Oil red O staining revealed that ox-LDL treatment induced uptake of large amount of lipids by HASMCs. However, rosuvastatin treatment reversed this phenomenon (**Figure 5A**). EVs secreted by inactivated HUVECs (EV.con) inhibited



endothelial cells on migration of HASMCs. (d) Effect of EVs from LPS-stimulated endothelial cells (EV.LPS) on the migration of HASMCs. (B) The effect of LPS-activated HUVECs on migration of HASMCs. (c) Transwell assay for the effect of HUVECs, LPS and GW4869, either alone or in combination, on the migration of HASMCs (a) HUVECs culture alone. (b) HUVECs and HASMCs co-culture (Co-culture.Con). (c) HUVECs and HASMCs co-culture in the presence of 1 μ g/mL LPS (Co-culture.LPS). (d) HUVECs and HASMCs co-cultured with 2.5 nmol/mL GW4869 (GW.Con). (e) HUVECs and HASMCs co-cultured in the presence of 2.5 nmol/mL GW4869 and 1 μ g/mL LPS (GW.LPS). (D) Transwell assay for effect of GW4869 treatment on the inhibition property of endothelial EVs on the migration of HASMCs (Data represents mean \pm SD of 6 independent experiments, #p < 0.05 and #p < 0.01, respectively. *p < 0.05 and **p < 0.01).

accumulation of lipids in HASMCs, whereas EVs derived from LPS activated HUVECs (EV.LPS) had no effect on the aforementioned phenomenon. HPLC revealed the average retention time of free cholesterol in HASMCS was 6.415 min. Compared with EV.con (43.97 \pm 9.02 ng/µg), ox-LDL treatment significantly enhanced retention of intracellular cholesterol (62.12 \pm 4.49 ng/µg) (P < 0.05). Interestingly, EV.LPS treatment exerted comparable effect as ox-LDL (59.21 \pm 4.55 ng/µg) (**Figures 5B,C**). These findings demonstrated that EVs from non-activated HUVECs (EV.con) reduces accumulation of lipids in HASMCs, in contrast to EV.LPS.

LPS Disrupted the Expression of miRNAs in Endothelial EVs

EVs derived from non-activated HUVECs disrupted the relative expression of several miRNAs. Significant change in the expression of miRNAs was based on (fold change) >0 for up-regulated expression and log2 (fold change)<0 for down-regulated expression, both at P < 0.05. Particularly, LPS treatment dysregulated the expression of 234 miRNAs. Of these 124 were over-expressed whereas 110 were under expressed. Further assessment revealed that 13 of the most differently expressed miRNAs participated in various aspects of atherosclerosis. Of these, four miRNAs including miR-27a-3p, miR-365b-3p,



miR-126-5p, and miR-155-5p were over-expressed. The rest, including let-7b-5p, miR-92a-3p, miR-30a-5p, miR-125a-3p, miR-181a-2-3p, miR-143-3p, miR-21-3p, miR-10a-5p, and miR-10b-5p (**Figures 6A,B**), were under-expressed.

Curcumin and Nicotinic-Curcumin Inhibited LPS-Induced Endothelial Inflammatory Responses

Certain inflammatory cytokines can cause inflammation and disrupt the normal functioning of endothelial cells. In the early AS stage, inflammatory cytokines stimulate endothelial cells to produce adhesion molecules which recruit mononuclear macrophages as well as promote proliferation and migration of vascular smooth muscle cells. Mononuclear macrophages and vascular smooth muscle cells engulfing large amounts of lipids and differentiate into foam cells, forming atherosclerotic plaques (30). ELISA and RT-PCR tests revealed that LPS simulation upregulated secretion of IL-1 β (P < 0.01; Figures 7A,B). However, NC treatment reversed this phenomenon. Comparatively, NC treatment (4 μ M) exerted better anti-inflammatory effect than rosuvastatin (1 μ M) (Figure 7). In addition, curcumin and NC significantly inhibited proliferation of HUVECs (P < 0.01; Figure 7C). Wound scratch assay further revealed that LPS remarkably promoted the migration of HUVECs (P < 0.01). However, NC (4 μ M) treatment inhibited the migration of HUVECs (P < 0.01; Figures 7D,E). Also, LPS and Cur exerted comparable apoptotic effect on HUVECS (Figures 8A,B).

LPS Promotes the Release of HUVECs-Derived EVs but Curcumin and Nicotinic-Curcumin Inhibits This Effect

The total exosomal proteins content in each group was measured using the BCA method. The size and the concentration



of the particles were also determined. Nanoparticle tracking analysis (NTA) and BCA analyses revealed that LPS treatment upregulated secretion of endothelial EVs. However, after curcumin (4 μ M) and NC (4 μ M) treatments significantly modulated production of EVs and inhibited inflammatory responses by HUVECs (**Figures 9A,B**). Western blot further revealed that LPS treatment up-regulated the expression of Rab27a. However, NC treatment downregulated expression of Rab27a but increased that of Rab27b (**Figures 9C,D**). Overall, NC reduced the secretion of EVs related to the inhibited expression of Rab27a proteins.

The Effects of Curcumin and Nicotinic-Curcumin on the Expression of miR-125a-3p, miR-92a-3p, miR-126-5p, and miR-143-3p in Endothelial EVs

RT-PCR revealed that LPS treatment reduced the relative expressions of miR-125a-3p, miR-92a-3p, and miR-143-3p, but increased that of miR-126-5p. However, NC treatment up-regulated the relative expressions of miR-125a-3p and miR-92a-3p but down-regulated that of miR-126-5p. Interestingly NC treatment had no effect on that of miR-143-3p (**Figure 10**). Curcumin treatment only decreased the expression of miR-126-5p, but increased the expression of miR-125a-3p, miR-92a-3p, and miR-143-3p.

DISCUSSION

Communication between endothelial cells and vascular smooth muscle cells are essential in the progression of cardiovascular disease (20). We focused on regulatory effects of endothelial EVs from LPS activated or non-activated HUVECS on VSMCs. In general, EVs of non-activated HUVECS modulate AS by regulating proliferation, migration and lipid storage in VSMCs. However, LPS-induced EVs disrupt the beneficial functions of endothelial EVs from non-activated HUVECS. Further analyses revealed that LPS treatment dysregulated the expression of numerous miRNAs in endothelial EVs. The miRNAs participates in regulating several processess and aspects of vascular smooth muscle cells. LPS treatment regulates production of microRNAs in endothelial EVs. The role of miR-27a-3p is controversial. For instance, miR-27a-3p is thought to suppress expression of ARHGEF26 and inhibits proliferation of SMCs by interacting with a conserved binding site in the 3' untranslated region of ARHGEF26 mRNA (31). The 3' region contains multiple conserved binding sites for miR-27a-3p, which regulates the expression TSP-1 gene. This suggests that miR-27a-3p accelerates arteriosclerosis development by inhibiting the expression of mRNA for TSP-1 gene (31). Meanwhile, miR-27a-3p regulates endothelial inflammatory response by targeting MAPK, TGFBR1 and TGFBR2 signaling pathways (32). In coronary atherosclerosis, miR-365b-3p inhibits proliferation and migration of human coronary artery smooth muscle cells by



directly targeting ADAMTS1 (33). Surprisingly, in our study, LPS treatment increased the expression of miR-365b-3p in endothelial EVs, which was contrary to the effect of LPS-induced EVs. This might be attributed to under-expression of miR-365b-3p, only a minor regulatory miRNA. MiR-126-5p modulates atherosclerosis, regulates endothelial stress response and promotes endothelial repair (34–36). At sites where disturbed laminar flow compromises endothelial cell

(EC) function, atherosclerosis plaques developed preferentially is correlated with low miR-126-5p levels in endothelial cells. Endothelial miR-126-5p promotes endothelial proliferation and limits atherosclerosis by suppressing expression of delta-like 1 homolog (Dlk1) (37). It also promotes contractile switching of aortic smooth muscle cells by targeting VEPH1 and alleviates Ang II-induced abdominal aortic aneurysm in mice. MicroRNA-155-5p promotes progression of atherosclerosis (38) mainly by



enhancing inflammatory response during the development of the disease by targeting SOCS1 via the STAT3 and NF- κ B signaling pathways. MiR-155-5p also promotes cholesterol uptake on macrophages, which may be related to the accumulation of cholesterol in VSMCs (24, 39).

In ECs, let-7b-5p promotes angiogenesis by inhibiting TGFBR1, its target gene. In addition, let-7b-5p modulated the secretion of TNF and IL-6 by monocytes as well as expression of SERPINE1 in LPS-activated macrophage (40). Let-7b-5p

also promotes the degradation of fatty acids in macrophages through mitochondrial respiration. The effect of let-7b-5p on the functioning of vascular smooth muscle cells is rarely reported. MiR-92a-3p promotes apoptosis of endothelial cells by targeting SIRT6 and activating MAPK signaling pathway (41). However, separate studies show that inhibition of miR-92a-3p regulates the development of atherosclerotic plaques (42, 43). Furthermore atheroprotective flow patterns decrease the level of miR-92a, which in turn increases KLF2 expression to



maintain endothelial homeostasis (44). Up regulated expression of miR-30a-5p in atherosclerotic plaques suggest the miRNA is a potential diagnostic marker for atherosclerotic plaques. However, the regulatory property of miR-30a-5p needs further investigation (45). Interventional therapy in restenotic arteries of lower extremity substantial inhibits the expression of miR-125a-3p, relative to normal arteries. A more in-depth study revealed that miR-125a-3p effectively inhibits the proliferation and migration of VSMCs and the occurrence of vascular stenosis by targeting MAPK1 (23). Moreover, miR-125a-3p was overexpressed in ECs. Meanwhile, a separate study revealed that by inhibits the expression of vascular endothelial growth factor (VEGF), miR-125a-3p repressed angiogenesis in renal cancer tissues (46). LPS treatment mediated reduction of miR-125a-3p expression disrupted the normal functions of EVs. We found MiR-181a-2-3p possibly regulates the TLR/NFkB signaling and activates inflammatory and inflammasome responses. MiR-143-3p modulates both cellular and EVs-mediated responses in pulmonary vascular cells (47). Extracellular microvesicles containing miR-143-3p secreted by l-flow or KLF2 overexpressed in endothelial cells inhibits atherosclerosis in adjacent VSMCs (48). In addition, bidirectional movement of extracellular miR-143-3p may occur between SMCs and ECs under atherosclerotic conditions (49). Exosomal miR-21-3p from nicotine-treated macrophages increases migration and proliferation VSMC by targeting PTEN. However, the expression of the miRNA in ECs is very low (50). LPS down-regulated the expression of homologous miR-10a-5p and miR-10b-5p. The expression of miR-10a-5p negatively correlated to atherosclerosis progression. In macrophages, miR-10a-5p suppressed the expression of ligand-dependent nuclear receptor co-repressor, thereby promoting fatty acid oxidation (51). However, in a separate study, atherosclerotic plaques induced the over-expression of miR-10b-5p in VSMCs, which promoted the proliferation of these cells by targeting TIP30 via the Akt pathway (25).

Inflammation-induced dysregulated expression of miRNAs in EVs regulates pathogenesis of atherogenesis. Accordingly, such miRNAs are potential diagnostic markers for atherogenesis. Based on LPS-induced changes in the expression of miRNAs in endothelial EVs alone, we believe over-expression of miR-155-5p participates in accumulation of lipids in VSMCs. Also, over-expression of miR-125a-3p and miR-143-3p mediates the proliferation and migration of VSMCs induced by endothelial EVs. We also believe the LPS-induced proteomic changes in endothelial EVs are the major factor influencing the functioning of VSMCs.

Meanwhile, LPS treatment up-regulated the secretion of endothelial EVs. However, curcumin and nicotinic-curcumin treatment suppressed LPS-mediated over-production of IL-1 β in endothelial cells during inflammation. In addition, curcumin and nicotinic-curcumin treatment reduced LPS-mediated secretion of EVs. Also, curcumin and nicotinic-curcumin treatment



upregulated the expression of miR-125a-3p, miR-92a-3p, and miR-143-3p, but modulated that of miR-126-5p, suggesting the two treatments suppresses inflammation.

Summarily, endothelial cells can regulate functioning of vascular smooth muscle cells via EVs, thereby influencing progression of atherosclerosis. Inflammation dysregulates the expression of endothelial EVs, which may be related to the expression levels of several miRNAs in the EVs. Therefore, inhibition of endothelial inflammation can potentially modulate atherosclerosis. Drugs that regulate secretion and expression of miRNAs in endothelial EVs are potential treatments for atherosclerosis.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by the Animal Ethics Committee of the Hunan

University of Chinese Medicine (HN-LL-KY-2016–004–01). Written informed consent for participation was not required for this study in accordance with the national legislation and the institutional requirements.

AUTHOR CONTRIBUTIONS

DX and YL: data curation and formal analysis. DL: funding acquisition, project administration, resources, and supervision. DX and YQ: investigation. LZ and XLi: methodology. YC and XLin: software. DX and YH: writing—original draft. DX and DL: writing—review and editing. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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