

Phenotypic transformation of intimal and adventitial lymphatics in atherosclerosis: a regulatory role for soluble VEGF receptor 2

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ABSTRACT: The role of lymphatics in atherosclerosis is not yet understood. Here, we investigate lymphatic growth dynamics and marker expression in atherosclerosis in apolipoprotein E-deficient (*apoE*^{-/-}) mice. The prolymphangiogenic growth factor, VEGF-C, was elevated in atherosclerotic aortic walls. Despite increased VEGF-C, we found that adventitial lymphatics regress during the course of formation of atherosclerosis (*P* < 0.01). Similar to lymphatic regression, the number of lymphatic vessel endothelial hyaluronan receptor 1 (LYVE-1⁺) macrophages decreased in the aortic adventitia of *apoE*^{-/-} mice with atherosclerosis (*P* < 0.01). Intimal lymphatics in the atherosclerotic lesions exhibited an atypical phenotype, with the expression of podoplanin and VEGF receptor 3 (VEGFR-3) but not of LYVE-1 and prospero homeobox protein 1. In the aortas of atherosclerotic animals, we found markedly increased soluble VEGFR-2. We hypothesized that the elevated soluble VEGFR-2 that was found in the aortas of *apoE*^{-/-} mice with atherosclerosis binds to and diminishes the activity of VEGF-C. This trapping mechanism explains, despite increased VEGF-C in the atherosclerotic aortas, how adventitial lymphatics regress. Lymphatic regression impedes the drainage of lipids, growth factors, inflammatory cytokines, and immune cells. Insufficient lymphatic drainage could thus exacerbate atherosclerosis formation. Our study contributes new insights to previously unknown dynamic changes of adventitial lymphatics. Targeting soluble VEGFR-2 in atherosclerosis may provide a new strategy for the liberation of endogenous VEGF-C and the prevention of lymphatic regression.—Taher, M., Nakao, S., Zandi, S., Melhorn, M. I., Hayes, K. C., Hafezi-Moghadam, A. Phenotypic transformation of intimal and adventitial lymphatics in atherosclerosis: a regulatory role for soluble VEGF receptor 2. *FASEB J.* 30, 2490–2499 (2016). www.fasebj.org

KEY WORDS: lymphangiogenesis · LYVE-1 · podoplanin · macrophages · VEGF-C

INTRODUCTION

Atherosclerosis is a chronic inflammatory disease that involves both innate and adaptive immunity (1). As part of the inflammatory process, activated macrophages accumulate in the aortic wall and stimulate angiogenesis—the growth of new blood vessels—and thereby contribute to plaque formation (2). Angiogenesis in plaques is associated with plaque

vulnerability (3). Accumulation of macrophages in atherosclerotic plaques represents an imbalance of macrophage recruitment from blood into the aortic wall and macrophage clearance from the plaques *via* lymphatics (4). In atherosclerosis, macrophage recruitment is increased as a result of angiogenesis and the presence of proinflammatory cytokines (5, 6). Lymphatics are expressed in human atherosclerosis (7, 8); however, little is known about the dynamics of lymphatic growth in the course of the formation of atherosclerosis.

Lymphatics are essential to immune function and key in such pathologies as cancer (9, 10). The role of lymphatics in atherosclerosis is just now undergoing study (11). The lymphatic endothelium heterogeneously expresses lymphatic vessel endothelial hyaluronan receptor 1 (LYVE-1), VEGF receptor 3 (VEGFR-3), podoplanin, and prospero homeobox protein 1

ABBREVIATIONS: *apoE*^{-/-}, apolipoprotein E-deficient; LYVE-1, lymphatic vessel endothelial hyaluronan receptor 1; M:F, male-to-female ratio; PROX1, prospero homeobox protein 1; VEGFR, VEGF receptor; WD, Western diet; WT, wild-type

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(PROX1) (9). Lymphangiogenesis, the growth of new lymphatics, shares several characteristics with angiogenesis, the growth of new blood vessels (12), and several factors that induce angiogenesis also promote lymphangiogenesis (13). These factors, members of the VEGF family and cytokines such as TNF- α , are elevated in chronic inflammatory conditions (14). Lymphangiogenesis and angiogenesis often accompany each other in chronic inflammatory conditions (15). Despite their commonalities and joined occurrence in pathologies, lymphangiogenesis and angiogenesis have distinct dissimilarities (12).

Lymphatic growth lags behind angiogenic growth (12). We found that actively growing blood vessels suppress lymphatic growth (12). The suppression occurs because the angiogenic endothelium competes for and binds with the growth factor that would otherwise induce the growth of the lymphatic endothelium (12). The endothelium of actively growing blood vessels expresses higher levels of VEGFR-2 (12). These endothelial VEGFR-2 molecules bind with and deplete the growth factor, VEGF-C, from the extracellular matrix (12). VEGFR-2 is proteolytically cleaved from the endothelial membrane (16). Similar to the membrane-bound form, soluble VEGFR-2 binds with VEGF-C and impedes its biologic activity.

VEGF-C has an affinity to endothelial VEGFR-2 but also to VEGFR-3 on the lymphatic endothelium. The binding of VEGF-C to VEGFR-3 induces lymphangiogenesis (14). The removal of VEGF-C by the angiogenic endothelium impedes lymphatic growth, whereas blood vessel endothelium thrives (12). This mechanism, which is termed receptor-mediated growth factor internalization, explains why lymphatics are not found together with actively growing blood vessels (12). Whether angiogenesis in atherosclerosis regulates lymphatic growth remains to be studied.

When angiogenic vessels mature, they down-regulate their VEGFR-2 to the baseline levels that are found in normal quiescent endothelium (12). With VEGFR-2 expressed at low levels, the endothelium of the matured blood vessel no longer depletes extracellular VEGF-C. Consequently, VEGF-C signals *via* the lymphatic endothelial VEGFR-3 and induces lymphatic growth. This explains the temporal lag of lymphangiogenesis relative to angiogenesis, presumably until the newly formed blood vessels mature (12).

Apolipoprotein E-deficient (*apoE*^{-/-}) mice spontaneously develop atherosclerosis, and feeding them a Western diet (WD) further accelerates the process (17). Angiogenesis has been established in the context of atherosclerosis in these mice (18); however, lymphatics in the atherosclerotic lesions of these mice have not been studied. To explore lymphangiogenesis in the dynamic context of atherosclerosis formation, we examined aortas of young and aged *apoE*^{-/-} mice. We further studied the burden of WD on their lymphatics in the aorta. In the aortas of these mice, we quantified immune cells, lymphatic markers, and soluble VEGFR-2.

TABLE 1. *Composition of the semipurified WD*

WD composition	
Ingredient	Amount
CHO/fat/protein (% calories)	60:20:20
Cholesterol (g/kg)	10
Casein (g/kg)	105
Lactalbumin (g/kg)	105
Dextrose (g/kg)	186
Sucrose (g/kg)	186
Cornstarch (g/kg)	194
Margarine B (80% fat) (g/kg)	116 (93 fat)
Mineral mix (g/kg)	46
Vitamin mix (g/kg)	12
Choline chloride (g/kg)	3

Carbohydrate, fat, and protein ratio (60:20:20% calories) represents the percentage of dietary calories provided by each nutrient category. Diet cholesterol was added at 1% by weight, and margarine B was a blend of fats common in the U.S. diet. It was based on saturated animal fat as 24% milk fat and 40% beef tallow, with sources of monounsaturated and polyunsaturated fats from 20% chicken fat and 16% soybean oil. The ratio of saturated to monounsaturated to polyunsaturated fatty acids was approximately 44:38:18. The diet was adjusted for vitamin and mineral concentrations as a percentage of calories (4.2 kcal/g). CHO, carbohydrate; prot, protein.

MATERIALS AND METHODS

Animals

Young (8- to 12-wk-old mice) and aged (54- to 58-wk-old retired breeders) C57BL/6J mice (000664; The Jackson Laboratory, Bar Harbor, ME, USA) and young (8- to 12-wk-old mice) and aged *apoE*^{-/-} mice (54- to 58-wk-old retired breeders from our own colony) were housed in a temperature-controlled animal facility with a 12-h light/dark cycle and were fed standard laboratory chow and water *ad libitum*. To accelerate atherosclerosis formation, *apoE*^{-/-} animals, starting at 6 wk of age, were fed a WD for 26 wk. For all experiments, we used both male and female mice and indicated the male-to-female ratio (M:F) for each experiment. All animal experiments were approved by the Institutional Animal Care and Use Committee of Brigham and Women's Hospital.

Diet

WD was made from semipurified components. The carbohydrate/fat/protein ratio was calculated as percentages of total calories and adjusted for vitamins and minerals (Table 1). The regular diet was in accordance to the standard composition, Formulab Diet 5008C33.

Plasma cholesterol and triglyceride measurements

Mouse blood (1 ml) was obtained through cardiac puncture under anesthesia. Heparinized blood was centrifuged at 12,500 rpm for 10 min at 4°C, and plasma was collected. Plasma total cholesterol levels and triglycerides were measured by using Thermo Infinity kits (Thermo Electron, Pittsburgh, PA, USA) and a serum triglyceride determination kit (Sigma-Aldrich, St. Louis, MO, USA), respectively.

Immunohistochemistry

Aortic trees were perfused with PBS, and aortic arches were harvested and snap-frozen in optimal cutting temperature

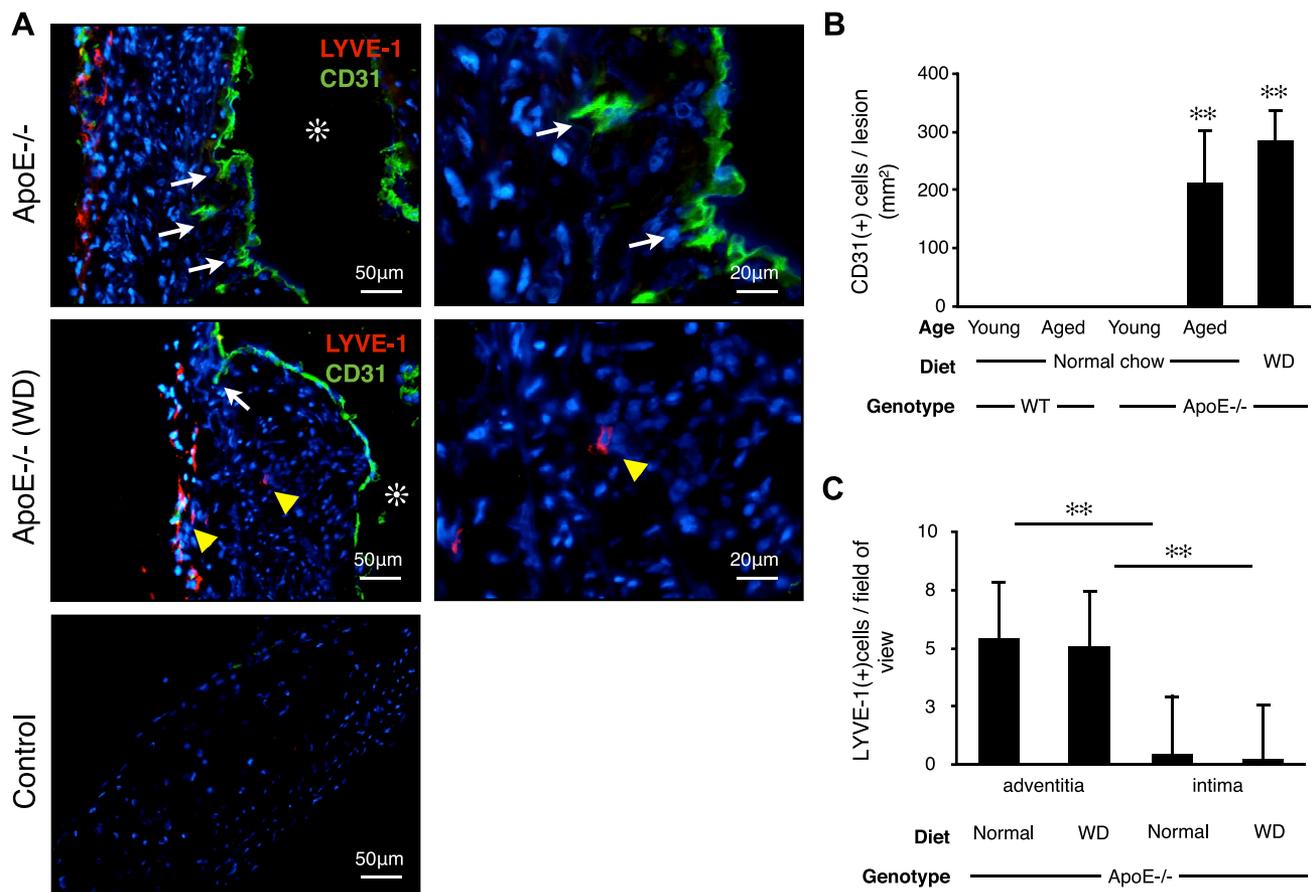


Figure 1. Scarcity of LYVE-1⁺ cells in atherosclerotic plaques despite abundant angiogenesis. Vascular endothelium staining (CD31, green) and lymphatic endothelium staining (LYVE-1, red) in atherosclerotic lesions of *apoE*^{-/-} mice. **A**) In aged *apoE*^{-/-} mice fed normal diet ($n = 3$; 54–56 wk; M:F, 0.6) and in WD-fed *apoE*^{-/-} mice ($n = 3$; 32 wk; M:F, 0.4), CD31⁺ cells were found in the intimal space, which indicated angiogenesis (white arrows). Asterisks indicate aortic lumen; yellow arrowheads indicate LYVE-1⁺ macrophages. **B**) Quantification of CD31⁺ cells in the intima (aged *apoE*^{-/-}: $n = 3$; 54–56 wk; M:F, 0.6; WD: $n = 3$; 32 wk; M:F, 0.4), means \pm SEM. **C**) Quantification of intimal vs. adventitial LYVE-1⁺ cells (aged *apoE*^{-/-}: $n = 5$; 54–58 wk; M:F, 0.6; WD: $n = 3$; 32 wk; M:F, 0.4), means \pm SEM. ****** $P < 0.01$.

compound (Sakura Finetek, Torrance, CA, USA). Sections (10 μ m) were prepared, air dried, and fixed in ice-cold acetone for 10 min. Sections were blocked with 3% nonfat dried milk bovine working solution (Sigma-Aldrich) and stained with anti-mouse CD31 mAb (1:50, 0.31 μ g/ml; BD Pharmingen, Brea, CA, USA) and anti-mouse LYVE-1 Ab (1 μ g/ml; ReliaTech, Wolfenbüttel, Germany), anti-mouse VEGFR-2 (5 μ g/ml; R&D Systems, Minneapolis, MN, USA), or anti-mouse podoplanin (1 μ g/ml; ReliaTech), anti-mouse VEGFR-3 (5 μ g/ml; R&D Systems), anti-human PROX1 (5 μ g/ml; R&D Systems), and CD11b (1:200, 0.63 μ g/ml; BD Pharmingen). After an overnight incubation at 4°C, sections were washed and stained for 60 min at room temperature with Alexa Fluor 647 goat anti-rabbit IgG (2 μ g/ml; Thermo Fisher Scientific Life Sciences, Waltham, MA, USA), Alexa Fluor 546 goat anti-rabbit IgG (2 μ g/ml; Thermo Fisher Scientific Life Sciences), Alexa Fluor 488 goat anti-rat IgG (2 μ g/ml; Thermo Fisher Scientific Life Sciences), and Alexa Fluor 488 goat anti-hamster IgG (2 μ g/ml; Thermo Fisher Scientific Life Sciences).

Whole-mount immunofluorescence

Abdominal aortas were excised and fixed in 4% paraformaldehyde for 1 h at 4°C, washed in PBS and methanol, blocked in 10% goat serum and 1% Triton X-100, and stained

with anti-mouse LYVE-1 Ab (4 μ g/ml; ReliaTech) and anti-mouse CD31 mAb (1:12.5; BD Pharmingen) overnight at 4°C. Samples were washed and stained with Alexa Fluor 647 goat anti-rabbit IgG (20 μ g/ml; Thermo Fisher Scientific Life Sciences) and Alexa Fluor 488 goat anti-rat IgG (20 μ g/ml; Thermo Fisher Scientific Life Sciences) overnight at 4°C. Photomicrographs were obtained across the length of the entire aortic whole mount. Single images were subsequently merged with a mosaic whole mount by using ImageJ software with the MosaicJ plugin (National Institutes of Health, Bethesda, MD, USA). Lymphatic vasculature was measured as the sum of areas > 100 -pixel threshold and expressed as the ratio of lymphatics to the entire area of the whole mount.

Western blot

Aortic trees were perfused *in situ* with PBS. Whole aortas were harvested, minced, and snap-frozen. Of RIPA lysis buffer (Sigma-Aldrich), 200 μ l was added to each minced aorta. RIPA lysis buffer was prepared to contain protease and phosphatase inhibitors (P2850, P5726, P8340; Sigma-Aldrich). Tissues were homogenized with a tissue homogenizer and centrifuged (12,500 rpm for 10 min at 4°C). Supernatants were collected, and protein concentrations were measured by using protein assay (Bio-Rad, Hercules, CA, USA). Equal amounts of total protein were loaded onto SDS-PAGE (Bio-Rad). Gels were run at 170 V for ~ 40 min.

Elevated soluble VEGFR-2 expression in atherosclerotic aortas

We next investigated the expression of VEGFR-2 in immunohistochemistry. In all examined aortas, VEGFR-2 colocalized with luminal CD31⁺ endothelial cells (Fig. 5A). In atherosclerotic lesions in *apoE*^{-/-} mice, VEGFR-2 was expressed in the intimal angiogenic vessels that costained for CD31 (Fig. 5A).

To explore potential mechanisms for the adventitial LYVE-1 regression, and despite the elevated VEGF-C in atherosclerotic aortas, we quantified the expression of soluble VEGFR-2 in the aortas. We performed Western blot analysis for soluble VEGFR-2. The expression of the 75 kDa soluble VEGFR-2 splice variant was significantly higher in aged *apoE*^{-/-} mice fed regular diet as well as in WD-fed *apoE*^{-/-} mice compared with controls.

DISCUSSION

In atherosclerosis, intimal and adventitial angiogenesis are correlated with disease progression (2, 20);

however, the expression and function of lymphatics in atherosclerosis are less well understood (21). We discovered a surprising regression of adventitial lymphatics and a reduction in LYVE-1⁺ macrophages with atherosclerosis. LYVE-1⁺ macrophages are key contributors to lymphangiogenesis (19). In contrast to the adventitia, in which these cells abound, few LYVE-1⁺ cells were found in the intima.

A lack of classic lymphatics, in which angiogenic vessels proliferate, is intriguing as lymphangiogenesis and angiogenesis are closely related processes that are often caused by the same factors (9). Yet, despite the prolymphangiogenic milieu of the atherosclerotic aortas, which is evidenced by increased VEGF-C, there is a paucity of classic lymphangiogenesis in the atherosclerotic lesions and even lymphatic regression in the adventitia.

To investigate the mechanism that underlies the adventitial lymphatic regression, we considered whether VEGF-C is potentially trapped by VEGFR-2, as we previously showed (12). Angiogenic endothelial cells express higher levels of VEGFR-2. We showed angiogenesis and VEGFR-2 expression in atherosclerotic aortas both in the intimal and adventitia,

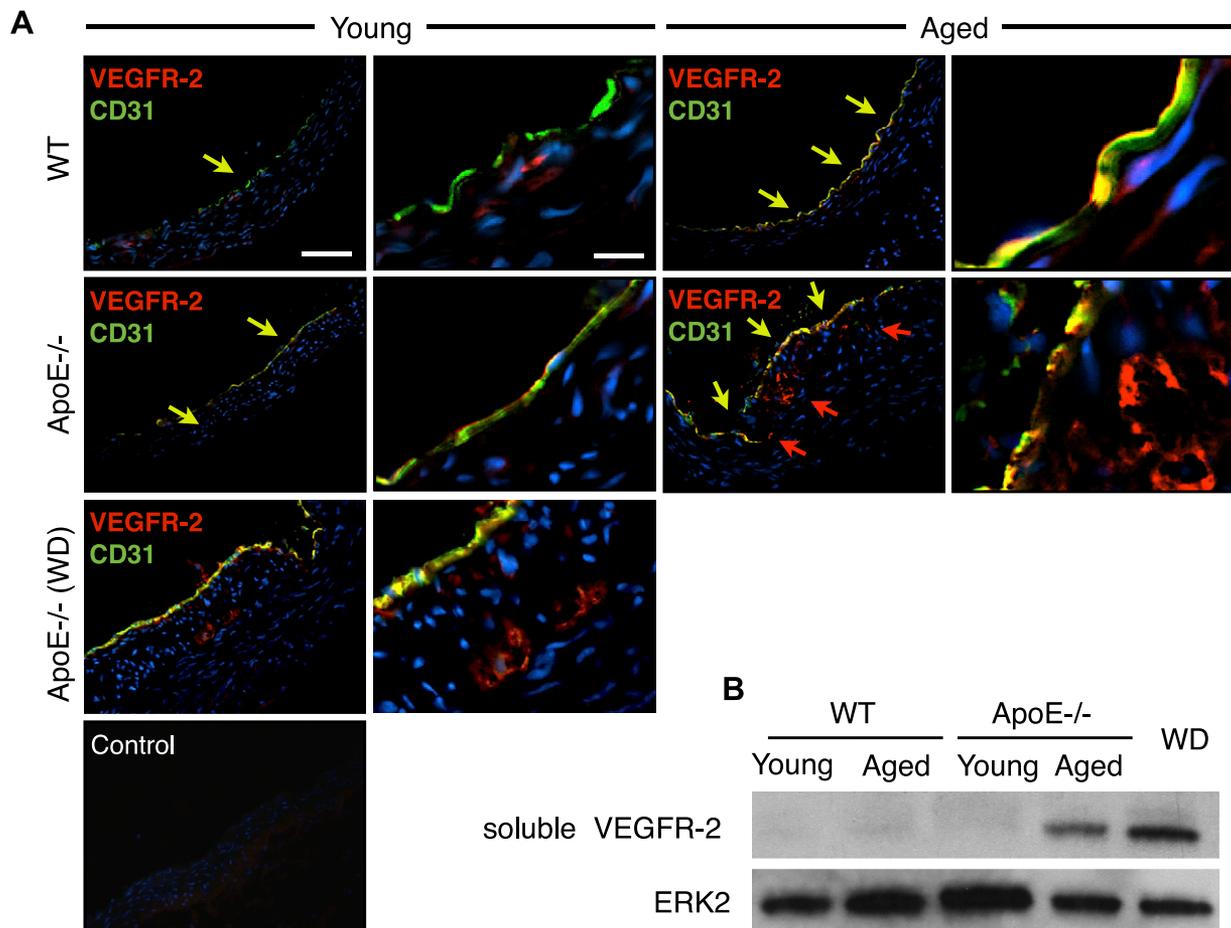


Figure 5. VEGFR-2 and soluble VEGFR-2 in aortas of aged and atherosclerotic mice. **A**) Immunofluorescence staining for VEGFR-2 (red) and the vascular endothelial cell marker, CD31 (green). Coimmunostaining for CD31 and VEGFR-2 (yellow arrows) and VEGFR-2 staining (red arrows). **B**) Western blot analysis for soluble VEGFR-2 (75 kDa) protein in aortic walls, ERK-2 loading control (young WT mice: *n* = 4; 8–12 wk; M:F, 0.5; aged WT mice: *n* = 4; 54–58 wk; M:F, 0.5; young *apoE*^{-/-} mice: *n* = 4; 8–12 wk; M:F, 0.5; aged *apoE*^{-/-} mice: *n* = 4; 54–58 wk; M:F, 0.5; WD: *n* = 4; 32 wk; M:F, 0.5).

