Although the response of endothelial cells to the disturbed blood flow in the vicinity of atherosclerotic lesions is known to be distinct from that elicited by nonatherogenic laminar flow, the mechanisms involved are poorly understood. Our initial studies confirmed that expression of the endothelial receptor tyrosine kinase Tie1 was evident at regions of atherogenic flow in mature animals. We therefore hypothesized that Tie1 plays a role in the endothelial response to atherogenic shear stress. Consistent with this, we found that Tie1−/− mice bred to the apoE-deficient background displayed a 35% reduction in atherosclerosis relative to Tie1+/−;ApoE−/− mice. Since deletion of Tie1 results in embryonic lethality secondary to vascular dysfunction, we used conditional and inducible mutagenesis to study the effect of endothelial-specific Tie1 attenuation on atherogenesis in Apoe−/− mice and found a dose-dependent decrease in atherosclerotic lesions. Analysis of primary aortic endothelial cells indicated that atheroprotective laminar flow decreased Tie1 expression in vitro. Attenuation of Tie1 was associated with an increase in eNOS expression and Tie2 phosphorylation. In addition, Tie1 attenuation increased IkBα expression while decreasing ICAM levels. In summary, we have found that shear stress conditions that modulate atherogenic events also regulate Tie1 expression. Therefore, Tie1 may play a novel proinflammatory role in atherosclerosis.

Introduction
Atherosclerotic lesions have a predilection for locales exposed to disturbed blood flow that can trigger inflammation and apoptosis (1–5). Disturbed flow is seen at the aortic sinus and at branch points such as bifurcations of the carotid, coronary, and renal arteries (2). In contrast, lesions rarely form along nonbranching segments of the descending aorta, where laminar flow is predominant and endothelial cells are protected against inflammatory activation and apoptosis (6, 7), and show increased expression of SOD and eNOS (8).

The force exerted by blood flow on the endothelium is termed shear stress. Mechanosensors of shear stress in endothelial cells include integrins, G protein-coupled receptors, PECAM-1, and the membrane lipid layer (5). The most recent additions to the list of shear stress mediators are the members of the Tie family of receptors. Tyrosine kinase with immunoglobulin-like and EGF-like domains 1, Tie1 is expressed almost exclusively in endothelial cells (11). Ablation of Tie1 expression results in early embryonic lethality after day 13.5 as a result of severe edema, hemorrhage, and loss of microvessel integrity (12, 13). Puri et al. (12) and Sato et al. (13) have independently shown that Tie1 is not essential for vasculogenesis but is required during embryonic development for the integrity and survival of vascular endothelial cells. Although Tie1 is important for embryonic vessel development, its expression is also increased during pathological conditions in the adult (14–16). Recent studies have indicated expression of Tie1 in inflammatory tissue of rheumatoid arthritis and osteoarthritis patients (17, 18). Additionally, overexpression of Tie1 augmented the inflammatory markers VCAM-1, ICAM-1, and E-selectin (19). Conversely, analyses of Tie1 siRNA-transfected HUVECs showed a decrease in the levels of inflammatory markers, including TLR2 and IL-1β. These data suggest that Tie1 expression mediates proinflammatory responses in endothelial cells.

Tie1 expression in the adult correlates distinctively to vascular regions exposed to disturbed flow in both physiological and pathological conditions. In optic microvessels, Tie1 promoter activity is increased at bifurcations and downstream of branches (9). The effect of fluid shear on Tie1 has also been demonstrated in vitro, whereby expression is decreased with brief application of laminar flow and altered by acute changes in shear stress magnitude (10). Conversely, disturbed flow in vitro upregulates Tie1 promoter activity (9). Tie1 is commonly found associated with Tie2 (20–22). Overexpression of Tie1 with Tie2 results in a cooperative recruitment of both receptors to the cell border (23). Tie1 can be activated by an Ang1 chimeric protein (COMP-Ang1) as well as by native Ang1 and Ang4, but not by Ang2 or Ang3 (21). Notably, coexpression with Tie2 was required for Tie1 activation (21), and association of Tie1 with Tie2 has been reported to modulate Tie2 activation (22, 24, 25). In bovine aortic endothelial cells subjected to shear stress in vitro (10), Tie1 was cleaved, and the intracellular product was found to coimmunoprecipitate with the protein tyrosine phosphatase Shp2 (26), which indicates that Tie1 might transduce signals in the absence of ligand activation. To our knowledge, the role of Tie1 in shear stress–induced vascular diseases such as atherosclerosis has not previously been studied.

We hypothesized that Tie1 plays an important role in shear stress–induced atherosclerosis and that attenuation of Tie1 would ameliorate atherosclerotic burden. In the present study, we show that Tie1...
expression was induced in atherosclerosis-prone regions of the vasculature, characterized by disturbed flow. Moreover, we show that increasing shear stress under laminar flow conditions downregulated Tie1 promoter activity in vivo. We also report a dose-dependent reduction in atherosclerosis in Tie1 attenuated ApoE−/− mice. Furthermore, we devised what we believe to be a novel method of isolating murine aortic endothelial cells (MAECs), facilitating in vitro experiments in parallel with our in vivo model. In vitro laminar flow with high shear stress for 24 hours downregulated Tie1 levels. Additionally, these in vitro studies document that Tie1 deletion increased shear stress–mediated eNOS and Tie2 phosphorylation, while increasing IκBα and decreasing ICAM levels. Our findings suggest that Tie1 may be a key modulator of endothelial response to pathological shear stress and a novel target of therapy for atherosclerosis.

Results
Tie1 is expressed at regions of atherogenic disturbed flow. Previous studies have shown that Tie1 is ubiquitously expressed in various endothelia at birth, including those of heart, lungs, kidney, liver, and brain (27). To map the expression pattern of Tie1 in adult macrovasculature, we used a transgenic mouse with Tie1 promoter–driven LacZ expression (12). In this model, the LacZ gene is knocked into the Tie1 locus (Tie1βc−/−) and placed under the control of endogenous Tie1 regulatory elements. As expected, Tie1-LacZ activity was ubiquitously expressed throughout the aorta and its branches in 4-week-old animals (Figure 1A), consistent with the role of an activated endothelium. Interestingly, in the 12-week-old adult mouse aorta, this expression was attenuated in the descending aorta (thoracic and abdominal segments). Whereas expression was in the thoracic and abdominal aorta was diminished, X-gal staining persisted in the aortic arch and proximal portion of the descending aorta, aortic arch branch arteries, and branches of the descending aorta, including vertebral and renal arteries (Figure 1, B–E). Distinct staining at the outer walls of these vascular bifurcations was observed where conditions of disturbed flow and low shear stress prevail (Figure 1E, inset).

LacZ expression was also detected in the endothelium of the aortic sinus and the aortic valves, areas characterized by disturbed flow (Supplemental Figure 1, A and B; supplemental material available online with this article; doi:10.1172/JCI42040DS1). We also noted more robust LacZ expression on the fibrosa surface, which experiences disturbed flow compared with the lamina flow of the ventricularis (28, 29). Additionally, X-gal staining was minimal in the ascending aorta, where laminar flow prevails (Supplemental Figure 1G). Since the aortic valve region is prone to atherosclerosis, we studied the effect of hypercholesterolemia caused by ApoE deletion on the endothelial Tie1 expression pattern. In young 4-week-old Tie1βc−/−;ApoE−/− mice, prior to the appearance of quantifiable atherosclerotic lesions, we noted a similar Tie1-LacZ expression pattern at the aortic sinus (Supplemental Figure 1, C and D) as seen in nonatherosclerotic littermates. This expression pattern persisted in 24-week-old Tie1βc−/−;ApoE−/− mice with advanced atherosclerotic plaque burden (Supplemental Figure 1, E and F). These results confirmed the persistence of accentuated Tie1 expression in regions characterized by disturbed flow.

In vivo Tie1 expression is reduced by laminar flow with increased shear stress. To study whether shear stress modification in vivo would affect Tie1 expression, we used shear stress–modifying casts (Figure 2, A and B). We implanted casts around the right common carotid artery of Tie1-LacZ mice for 7 days. These casts modulate endogenous shear stress by inducing lowered shear stress immediately upstream, increased shear stress inside the cast, and oscillatory shear stress with vortex flow immediately downstream (30, 31). Cheng et al. previously described increased eNOS activity in the tapered region of the cast, where laminar flow with increased shear stress is found (30), and documented formation of atherosclerotic lesions in the adjacent upstream region experiencing laminar flow with low shear stress and in the immediate downstream area undergoing oscillatory flow (31). Right common carotid arteries implanted with control nontapered casts (n = 4) expressed LacZ uniformly along the vessel (Figure 2D), similar to nonimplanted left subclavian and left carotid arteries. In contrast, X-gal staining was attenuated in the tapered region of the cast, where laminar flow with increased shear stress is induced (n = 4; Figure 2C). Comparable to controls, LacZ activity persisted in the aforementioned regions adjacent to the cast. Persistent endothelial expression of VE-cadherin in the areas of increased shear stress confirmed that the attenuation of Tie1-LacZ activity was not the result of endothelial denudation (Supplemental Figure 2), as also previously documented by others.

Figure 1
Tie1 expression at shear stress–specific sites in the adult aorta. Representative en face X-gal–stained, whole-mount distal aorta of 4-week-old (A) and 12-week-old (B–E) Tie1-LacZ mice. (A) Pervasive endothelial Tie1-LacZ expression in the aortic arch, branch arteries, and descending aorta wall of immature 4-week-old mouse. (B) Low-magnification view of aorta (aortic arch and thoracic and abdominal aorta) and (C) higher-magnification view depicting endothelial Tie1-LacZ expression in the aortic arch diminishing in the thoracic aorta as laminar flow is restored. (D) High-magnification image of thoracic aorta wall, illustrating endothelial Tie1-LacZ expression at ostia of vertebral artery branches (arrowheads). (E) High-magnification image of abdominal aorta wall, showing distinct X-gal staining at the outer walls of renal artery bifurcations (arrowheads). Inset: illustration of proatherogenic, disturbed flow as experienced at aortic branch points. White arrows indicate direction of blood flow. Original magnification, ×10 (B); ×20 (A and C); ×40 (D and E).
Tie1 suppression reduces distal aorta atherosclerosis progression in Apoe-null mice. To determine the role of Tie1 in atherosclerosis, Tie1−/−;Apoe−/− female mice (n = 49) fed a regular chow diet were evaluated for atherosclerotic plaque burden and compared with control Tie1+/+;Apoe−/− female littermates (n = 41). Female mice were chosen because of a clear gender-dependent difference in the progression of atherosclerosis (32). Mice were assessed at early (12 weeks old), intermediate (18 weeks old), and advanced (24 and 49 weeks old) stages of atherosclerosis. There were no statistically significant differences in serum cholesterol and triglyceride levels between experimental and control mice (Table 1). Semiquantitative RT-PCR analysis of descending aorta confirmed a 49% decrease in Tie1 mRNA expression of Tie1−/− mice (Figure 3A). Western blot assay of pulmonary endothelia showed a 40% reduction in Tie1 protein expression throughout the vascular endothelium (Figure 3B).

In the atherosclerosis-prone, disturbed-flow regions of the aorta, en face analysis showed 37.9% suppression of atherosclerosis. There were no statistically significant differences in lesion area between Tie1−/−;Apoe−/− and Tie1+/+;Apoe−/− mice at any stage (12 weeks, 40,030 ± 15,311 vs. 51,719 ± 6,094 μm²; 18 weeks, 156,155 ± 40,275 vs. 152,210 ± 25,157 μm²; 24 weeks, 286,433 ± 16,160 vs. 334,760 ± 41,861 μm²; 49 weeks, 426,680 ± 40,877 vs. 529,025 ± 52,711 μm²; Supplemental Figure 3A). The trend toward increased lesions in Tie1−/−; Apoe−/− mice, the opposite of the effects seen in the distal aorta, is interesting but does not warrant further comment, given the lack of statistical significance.

Endothelial-specific Tie1 deficiency alleviates atherosclerosis progression in a dose-dependent manner. We next assessed the effect of endothelial-specific Tie1 deletion on development of atherosclerotic lesions in Apoe−/− mice. Since Tie1 deletion leads to early embryonic lethality (12, 13), we used Cre-Lox technology to delete Tie1 in a spatially and temporally defined manner. We used SCL-ER²-Cre mice to achieve endothelial-specific, tamoxifen-mediated induction of Cre expression (33). To determine an efficient tamoxifen induction dosage, SCL-ER²-Cre mice were bred to Rosa-LacZ reporter background. Using a protocol of 2 mg intraperitoneal tamoxifen injection every 48 hours, for 14 days, we were able to obtain robust, endothelial-specific, Cre-mediated deletion (evidenced by X-gal staining), as documented in the aortic valve endothelium and in the microvasculature of the endocardium (Figure 4A). Tie1−/− mice were then bred with SCL-ER²-Cre mice, and generalized endothelial-specific deletion was quantified by Western assay of pulmonary endothelium, which showed a 81% reduction in Tie1 protein levels (Supplemental Figure 4). In the resulting mice with a null Tie1 allele, a floxed Tie1 allele, and endothelial-specific Cre (Tie1−/−;SCL-ER²-Cre mice), tamoxifen treatment induced a 65% reduction in aortic endothelial Tie1 mRNA level (Figure 4B).
Table 1
Plasma cholesterol and triglyceride levels for Apoe<sup>−/−</sup> mice

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<td>185.8</td>
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<td>18 weeks old</td>
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<td></td>
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<tr>
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<td>233.5</td>
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<td>14</td>
<td>275.3</td>
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<tr>
<td>24 weeks old</td>
<td></td>
<td></td>
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<tr>
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<td>49 weeks old</td>
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<td>9</td>
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Tie1 reduction caused no discernible effect; no significant differences were found between groups.

To elucidate the effect of Tie1 deletion on atherosclerosis progression, Tie1<sup>−/−</sup>;SCL-ER<sup>Cre</sup> mice were bred onto the Apoe<sup>−/−</sup> background. Tie1<sup>−/−</sup>;SCL-ER<sup>Cre</sup>;Apoe<sup>−/−</sup> female mice and control Tie1<sup>fl/fl</sup>;Apoe<sup>−/−</sup> female littermates were injected with tamoxifen as described above. Because tamoxifen administration has previously been associated with attenuation of atherosclerosis after chronic and unfiltered oral intake (34), we first determined whether acute and limited exposure to tamoxifen had any effect in our experimental model. We noted a protective effect (46% decrease) 4 weeks after treatment (i.e., 12 weeks old; 1.41% ± 0.22% vs. 0.76% ± 0.11%, P < 0.01; Supplemental Figure 5), which was not maintained at the 16-week time point (i.e., 24 weeks old; 4.32% ± 0.54% vs. 3.64% ± 0.86%, P = NS). Tamoxifen treatment temporarily decreased serum cholesterol in Tie1<sup>fl/fl</sup>;Apoe<sup>−/−</sup> mice compared with untreated controls at 12 weeks of age (456.44 ± 40.58 vs. 291.43 ± 19.36 mg/dl, P < 0.001), but this effect was lost by 24 weeks (313.54 ± 28.54 vs. 345.20 ± 24.61 mg/dl, P = NS; Table 2). Tamoxifen treatment also reduced serum triglycerides of 12- and 24-week-old animals (12 weeks old, 186.8 ± 8.00 vs. 147.98 ± 7.38 mg/dl, P < 0.01; 24 weeks old, 200.64 ± 9.81 vs. 141.95 ± 9.44 mg/dl, P < 0.01; Table 2). However, the tamoxifen-induced reduction in serum triglycerides was not associated with an atheroprotective effect at 24 weeks.

Consistent with our previous studies, en face analysis of the aorta from Tie1-deleted animals showed a statistically significant reduction in atherosclerotic lesions. In 12-week-old mice with early disease progression, we found a 68% decrease in lesions of Tie1-deleted mice compared with controls (0.76% ± 0.114% vs. 2.44% ± 0.46%, P < 0.0005; Figure 4D). The decrease in atherosclerosis persisted in mice up to 24 weeks of age, when Tie1 deletion resulted in a 70% decrease in plaque progression compared with controls (0.33 ± 0.03 vs. 0.07 ± 0.44 U, n = 3, P < 0.01; Supplemental Figure 6A), consistent with a decrease in inflammation associated with the attenuation in atherosclerosis. Similarly, analyses of tamoxifen-treated Tie1<sup>−/−</sup>;SCL-ER<sup>Cre</sup>;Apoe<sup>−/−</sup> and Tie1<sup>fl/fl</sup>;Apoe<sup>−/−</sup> control animals revealed a 70% reduction in macrophage infiltration (0.46 ± 0.07 vs. 0.14 ± 0.03 U, n = 3, P < 0.05; Supplemental Figure 6B). To determine whether Tie1 deletion within macrophage populations contributes to the attenuation in atherosclerosis, we evaluated the expression of Tie1 in primary macrophages. RT-PCR analysis demonstrated that there was no detectable Tie1 mRNA in macrophages (Supplemental Figure 6C); thus, a cell-autonomous effect of Tie1 in the macrophage population is unlikely to contribute to the decreased inflammation observed.

**Tie1 attenuation in vivo reduces expression of Rho-associated, coiled-coil containing protein kinases (ROCK1 and ROCK2) and cell adhesion molecules and increases eNOS levels.** To elucidate the effect of Tie1 deletion on inflammatory biomarker expression in vivo, we assayed RNA from the descending thoracic and abdominal aorta of tamoxifen-treated Tie1<sup>−/−</sup>;SCL-ER<sup>Cre</sup>;Apoe<sup>−/−</sup> mice and control Tie1<sup>fl/fl</sup>;Apoe<sup>−/−</sup> mice. mRNA levels of ROCK1 and ROCK2 were attenuated by 70% and 52%, respectively, whereas eNOS mRNA was increased (Figure 5, B–D). ICAM and VCAM mRNA expression were also decreased by 55% and 87%, respectively (Figure 5, E and F).

**Tie1 deletion in vitro augments shear stress-induced eNOS expression and reduces promflammatory signaling.** To determine the effect of Tie1 deletion on endothelial cell response to altered shear stress in vitro, we developed an immortalized MAEC line from both wild-type and Tie1<sup>fl/fl</sup>;SCL-ER<sup>Cre</sup> mice. Conditionally immortalized cells were derived from the H-2K<sup>b</sup>-tsAS8 transgenic mouse. Expression of the inducible tsAS8 tumor Ag (TAg) allows for the extended propagation of MAECs under permissive conditions. At passage 10, fluorescence-activated cell sorting analysis showed 89.7% positive immunostaining for CD31 (Supplemental Figure 7A). PECAM immunostaining showed uniform cytoplasmic distribution and increased localization to the cell border in apposition with adjacent cells (Supplemental Figure 7B). We also documented that MAECs took up 1,1'-dioctadecyl-3,3,3',3'-tetramethyldiocarbocyanine perchlorate–labeled acetylated LDL (DiI-AcLDL; Supplemental Figure 7C) and showed robust net work formation within 24 hours in Matrigel (Supplemental Figure 7D). Our MAECs displayed cobblestone morphology (Supplemental Figure 7E) and aligned to the direction of laminar flow with high
shear stress after 24 hours (Supplemental Figure 7F). These results demonstrated that MAECs isolated from immorto mice display the same characteristics of many primary endothelial cell lines.

Next, we isolated endothelial cells from homozygous \textit{Tie1}^{fl/fl};SCL-\textit{ER}^{T} \textit{Cre} immorto mice, allowing for the use of 4-hydroxytamoxifen (4OHT), the active metabolite of tamoxifen, to initiate Cre transcription and hence induce Tie1 deletion. To ascertain activation of Cre in vitro, we immunostained 4OHT-treated \textit{Tie1}^{fl/fl};SCL-\textit{ER}^{T} \textit{Cre} MAECs using Cre antibody. Increased nuclear staining indicates successful translocation of Cre (Figure 6, A and B). RT-PCR and Western blot analyses of full-length Tie1 indicate 4OHT administration resulted in 41% and 45% reductions, respectively, of Tie1 expression (Figure 6, C and D). Additionally, the cleaved Tie1 intracellular fragment (26, 35, 36) was reduced by 72% (Figure 6E).

We then exposed our MAECs to high-shear stress laminar flow for 24 hours. In our MAECs, laminar flow at 20 dynes/cm² decreased Tie1 receptor expression by approximately 40% (Figure 6F), and oscillatory flow at 5 dynes/cm² increased Tie1 expression by 30% (Supplemental Figure 8). Laminar flow also increased eNOS protein levels and potentiated eNOS phosphorylation in \textit{Tie1}^{fl/fl};SCL-\textit{ER}^{T} \textit{Cre} MAECs, as expected (37). Interestingly, eNOS expression and phosphorylation were further augmented by laminar flow after tamoxifen-induced deletion of Tie1 (Figure 6, G and H). In addition, Tie1 deletion further decreased shear stress–induced ICAM expression and increased \textit{IkB}α expression in response to laminar shear (Figure 6, I and J). We next observed that laminar flow increased Tie2 phosphorylation (38), and this shear stress–mediated increase in phosphorylation was accentuated by deletion of Tie1 (Figure 6K).

Moreover, in vitro oscillatory flow resulted in an increase in Tie1 expression (Supplemental Figure 8A), and attenuation of Tie1 under similar flow conditions increased eNOS expression and decreased p50 nuclear translocation (Supplemental Figure 8, B and C). In conclusion, these results suggest that Tie1 expression is regulated by shear stress and that Tie1 may have a proinflammatory role in promoting the progression of atherosclerosis.

Discussion

Our present findings provide major insights into the role of Tie1 in atherosclerosis. First, Tie1 expression was accentuated in regions of atherogenic shear stress, such as those with disturbed flow in the adult aorta. Second, there was dynamic downregulation of Tie1 expression in vitro and in vivo under conditions of laminar flow with high shear stress. Third, in conjunction with the expression of Tie1 at atherosclerosis-prone areas, we found that Tie1 reduction alleviated atherosclerosis progression in a dose-dependent manner. Finally, our in vitro and in vivo data support a proinflammatory role for Tie1.

Previous in vivo studies have focused on the role of Tie1 in early stages of embryonic development. Here, we showed that the expression pattern of Tie1 in the adult macrovasculature was different from that of the immature mouse. In the aorta of a young...
4-week-old mouse, we observed pervasive endothelial Tie1 expression consistent with aortic immaturity. However, in the 12-week-old adult mouse, Tie1 expression decreased specifically in the descending aorta. We noted strong X-gal staining at locales subjected to atherogenic disturbed flow (39), such as the aortic arch and its branch vessels, and bifurcations of the vertebral and renal arteries. LacZ expression was also observed at the aortic sinus and the aortic valves, areas characterized by disturbed flow.

Figure 4
Endothelial-specific Tie1 deletion reduces atherosclerosis burden. (A) Representative H&E-stained aortic valve showing endothelial-specific LacZ expression (arrowheads) from tamoxifen-treated SCL-ERT-Cre;Rosa26R-LacZ mouse. Original magnification, ×100. (B) RT-PCR analysis of aortic Tie1 levels from tamoxifen-treated Tie1fl/fl;SCL-ERT-Cre mice (65% reduction, P < 0.05). (C) Representative Sudan IV–stained distal aorta from Tie1fl/fl;SCL-ERT-Cre;ApoE–/– and Tie1fl/fl;ApoE–/– mice demonstrated lesion reductions in Tie1-deleted mice. (D and E) Mean aortic lesion area in tamoxifen-treated Tie1fl/fl;SCL-ERT-Cre;ApoE–/– and Tie1fl/fl;ApoE–/– mice analyzed at (D) 12 weeks (68% reduction, P < 0.0005) and (E) 24 weeks (70% reduction, P < 0.006). (F–H) Atherosclerotic lesion areas in Tie1-deleted mice, as assessed in 3 regions of the aorta, showed reduction of atherosclerosis in all regions of disturbed flow. Insets show respective regions of lesion analysis. Shown are mean atherosclerotic lesion areas at (F) lesser curvature of the aortic arch (12 weeks, 0.066% ± 0.023% vs. 0.038% ± 0.014%, P = NS; 24 weeks, 1.20% ± 0.379% vs. 0.206% ± 0.048%, 82% reduction, P < 0.01); (G) aortic arch branch arteries, including brachiocephalic, left common carotid, and left subclavian arteries (12 weeks, 0.211% ± 0.055% vs. 0.086% ± 0.026%, 56% reduction, P < 0.05; 24 weeks, 1.29% ± 0.283% vs. 0.568% ± 0.119%, 59% reduction, P < 0.05); and (H) descending aorta (12 weeks, 0.487% ± 0.074% vs. 0.118% ± 0.032%, 70% reduction, P < 0.01; 24 weeks, 1.13% ± 0.291% vs. 0.333% ± 0.097%, 75% reduction, P < 0.01). Data points denote individual animals, and horizontal bars indicate group average. *P < 0.05; **P < 0.01; ***P < 0.001.
The effect of altered shear stress on Tie1 expression was demonstrated in vivo by the use of vascular casts. Cheng et al. previously demonstrated that these in vivo shear-stress-modifying casts altered the shear forces in the carotid arteries of Apoe-null mice (31). Lowered shear stress upstream of the cast was found to induce formation of unstable plaques, and the immediate downstream region of the cast formed stable plaques in the presence of oscillatory flow. In our Tie1-LacZ reporter mice, we found persistence of Tie1 promoter–driven LacZ expression also in these immediate vicinities around the cast. Furthermore, Cheng and colleagues reported increased eNOS expression and detected no atherosclerotic lesions within the boundaries of the tapered cast (30). Interestingly, we found a dramatic reduction in Tie1 promoter activity after only 7 days of implanting tapered casts in Tie1-LacZ mice. Attenuation of LacZ expression was distinctly confined to the boundaries of the tapered cast, where shear stress increases from 10 to 25 N/m². Hence, the attenuation of Tie1 expression may be caused, at least in part, by both the higher shear stress magnitude and laminar flow conditions.

Interestingly, the in vivo response of endothelial cells to the same magnitude of shear stress may be context dependent, with endothelial cells from different vascular beds having different set points for response to shear stress. There is significant variability in the absolute wall shear stress of vascular regions within a given species, as well as between similar regions in the same species (reviewed in ref. 40). The shear stress of the mouse abdominal aorta is approximately 9 N/m² (41), and in our studies, little Tie1 expression was detected in the regions of laminar shear, but Tie1 expression persisted in regions of nonlaminar flow. By comparison, the average shear stress of the carotid arteries in mice has been calculated to be 15 N/m² (31), and Tie1 was expressed in carotid endothelial cells under normal conditions. However, when shear stress in the carotid arteries was increased by placement of vascular casts to 25 N/m², a magnitude comparable to the level of shear stress produced in most in vitro studies, there was a marked attenuation in Tie1 expression. Clearly, shear stress is not the only factor regulating Tie1 expression, but our data suggest that expression of Tie1 is downregulated in vitro and in vivo by shear stress profiles that are thought to be atheroprotective.

Since we found expression of Tie1 at regions of atherogenic shear stress, and a previous study reported Tie1 expression on both the surface and the subendothelium of atherosclerotic lesions (9), we postulated that attenuation of Tie1 may modulate atherosclerosis progression. We found that moderate reductions in Tie1 (Tie1−/−;Apoe−/− mice) elicited 35% and 38% reductions in atherosclerotic lesions of 24- and 49-week-old mice, respectively. Remarkably, further reductions of Tie1 by up to 80% in Tie1−/−;SCL-ER−/Cre;Apoe−/− mice resulted in 68% and 70% reduction in lesions of 12- and 24-week-old mice, respectively. Hence, using 2 different mouse models, our results demonstrated that Tie1 attenuation conferred amelioration of atherosclerosis progression.

To our knowledge, this is the first report of an in vivo biological function for Tie1 in the adult mouse. Whereas Tie1 plays a requisite role in prenatal vascular integrity (12, 42), our results suggest that loss of Tie1 in late adult stages confers an atheroprotective effect. This divergence in function may be due in part to the complex shear stress magnitudes and flow profiles experienced pre- and postnatally. Kontos et al. previously showed that Tie1 is capable of activating Akt to promote cell survival (43); Tie1 is also required for the maintenance of various adult vascular endothelial cell populations (12). Hence, Tie receptors may possess context-dependent roles (44) that are not only defined by endothelial heterogeneity, but also influenced by the physiologic environment that changes during vascular maturity.

We also noted that attenuation of Tie1 did not significantly alter the extent of atherosclerotic lesions, as assessed by quantification of lipid accumulation in the aortic valve region. However, there was a clear decrease in macrophage infiltration in this region. Interestingly, Tie1 attenuation elicited a trend of increased lesions at the aortic sinus, an effect opposite to that observed in the distal aorta. Previous studies have demonstrated a predilection of lesions for specific locales of the distal aorta in PECAM-knockout mice on either the Apoe-deficient or the LDL receptor–deficient background (45, 46). The aortic valve also experiences a unique and complex shear stress profile not evidenced elsewhere in the vasculature. Peacock previously described the flow profile in the aortic sinus as turbulent, composed of spinning vortices superimposed with random motion of fluid particles (28). In contrast, the flow profile at aortic bifurcations distal to the aortic sinus has been described as recirculatory (4), characterized by bidirectional blood flow with the resultant effect of a low time-averaged shear stress (2). Additionally, Butcher et al. demonstrated that the shear stress–induced transcriptional response of valvular endothelial cells is vastly different from aortic endothelial cells. More than 400 genes were differentially expressed, and overall, the inflammatory response of valvular endothelial cells were found to be less pronounced than that of aortic endothelial cells (47). Hence, Tie1 may regulate atherosclerosis progression in a shear stress–specific manner, discriminating between different flow profiles characteristic of unique anatomical and physiological locations within the vascular system and further highlighting the context-dependent response of heterogeneous endothelial populations.

To further investigate the effect of shear stress on Tie1 expression in vitro, we devised a method of isolating genetically modified MAECs that maintain a high percentage of cells displaying

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**Table 2**

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<td>14</td>
<td>291.4d</td>
<td>147.9d</td>
</tr>
<tr>
<td>Tie1−/−;SCL-ER−/Cre;Apoe−/−</td>
<td>Yes</td>
<td>12</td>
<td>293.9</td>
<td>172.9c</td>
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<td><strong>24 weeks old</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
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<tr>
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<td>10</td>
<td>352.1</td>
<td>200.6</td>
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<tr>
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<td>Yes</td>
<td>13</td>
<td>331.5</td>
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Tamoxifen temporarily decreased cholesterol levels at 12 weeks (i.e., 4 weeks after treatment), but the effect was lost by 24 weeks. Tamoxifen treatment decreased triglycerides at 12 and 24 weeks. Tie1 deletion decreased triglycerides. P < 0.05. dP < 0.01 vs. untreated Tie1−/−;SCL-ER−/Cre;Apoe−/−. *P < 0.05 versus tamoxifen-treated Tie1−/−;Apoe−/−.
endothelial characteristics. The culture of MAECs is a technically challenging process, often resulting in low yields with poor propagation properties. We circumnavigated these problems through the generation of conditionally immortalized aortic endothelial cells from Tie1<sup>fl/fl</sup>;SCL-ER<sup>T</sup>-Cre immortal mice and selecting with PECAM by flow-assisted cell sorting. Jat et al. first reported the derivation of conditionally immortalized cell lines from the H-2K<sup>s</sup>-tsA58 transgenic mouse (48); subsequently, several groups have successfully isolated a variety of cell lines from this mouse (49–51).

In this model, the large TAg of a temperature-sensitive strain (tsA58) of the simian virus 40 (SV40) is fused with the major histocompatibility complex promoter H-2K<sup>s</sup>, which is active in many primary endothelial cell lines.

We achieved 50% deletion of Tie1 in vitro, concordant with in vivo tamoxifen-induced deletion in Tie1<sup>fl/fl</sup>;SCL-ER<sup>T</sup>-Cre mice. We surmise that Cre expression under the SCL promoter may not be sufficient for extensive deletion of biallelic floxed Tie1 and anticipate that enhanced deletion could be achieved by development of Tie1<sup>–/fl</sup>;SCL-ER<sup>T</sup>-Cre cells. Marron et al. previously reported cleavage of Tie1 with phorbol ester or VEGF (35), while Chen-Konak and colleagues reported changes in levels of Tie1 intracellular fragment after brief alterations in shear stress (10). Upon administration of 4OHT, we also noted a concomitant decrease in levels of cleaved Tie1 intracellular fragment. Whereas deletion of Tie1 resulted in a 50% reduction in full-length Tie1 receptor, the cleaved Tie1 fragment was decreased by 74%. The significance of this observation is currently being explored.

In vitro experiments in MAECs showed that 24 hours of laminar flow at 20 dynes/cm<sup>2</sup> suppressed Tie1 expression and increased eNOS activity, while oscillatory flow at 5 dynes/cm<sup>2</sup> for a similar duration augmented expression of Tie1. Using bovine aortic endothelial cells, Chen-Konak et al. found that brief in vitro application of shear stress at 10 dynes/cm<sup>2</sup> temporarily decreased Tie1, but expression levels returned to baseline after 2 hours (10). Therefore, in our in vitro model recapitulated our in vivo data showing increased Tie1 expression at regions of disturbed flow. Stability of eNOS may be regulated by ROCK1 and ROCK2 (53, 54). We showed that genetic deletion of Tie1 augmented eNOS activity in vivo and in vitro while downregulating the transcription of ROCK1 and ROCK2 in vivo. Hence, our data suggest that Tie1 may signal upstream of the ROCK-eNOS signaling pathway.

An octameric negative shear stress response element that downregulates Tie1 expression has also been found (10). Shear stress response elements are targets of the NF-κB transcription factor, which in turn can also be regulated by shear stress (55). Recent studies have shown that siRNA silencing of Tie1 in vitro reduced expression of inflammatory markers (56), and overexpression studies demonstrated complementary findings (19), suggestive of a proinflammatory role for Tie1. We found that Tie1 deletion augmented shear stress–induced IkBα expression in vitro while decreasing expression of inflammatory biomarkers both in vivo and in vitro.

Finally, the effect of Tie1 deletion on the progression of atherosclerosis is unlikely to be exclusively related to decreases in Tie1-mediated signaling. As mentioned above, Tie1 and Tie2 have been shown to interact physically (21, 23, 24), but the in vivo biologi-
The relevance of these interactions has not been demonstrated. Attenuation of Tie1 has been shown to upregulate Tie2 phosphorylation (22), as has laminar flow (38). Our results showed that the combination of shear stress and Tie1 deletion augmented the increased phosphorylation of Tie2. Interestingly, increased Tie2 signaling has recently been associated with a decrease in atherosclerosis (57). A recent fluorescent resonance energy transfer-based proximity assay has documented Tie1-Tie2 complex formation at the cell surface and demonstrated that these interactions are inhibitory for Tie2 signaling (25). We did not observe a difference in the expression of either Ang1 or Ang2 in response to Tie1 attenuation following 24 hours of in vitro laminar flow (data not shown). Therefore, it is likely that the increased phosphorylation of Tie2 under these conditions (Figure 6K) may be explained, at least in part, by the relief of inhibitory interactions normally imposed by association with Tie1.

Figure 6
In vitro deletion of Tie1 and endothelial cell response to shear stress. (A–E) Treatment of Tie1<sup>fl/fl</sup>;SCL-ER<sup>T</sup>-Cre MAECs with 4OHT activated Cre and reduced Tie1 expression. Representative immunofluorescent images showing Cre nuclear localization following 4OHT treatment (B) compared with untreated controls (A). Scale bars: 100 μm. (C) RT-PCR analysis of Tie1 mRNA levels. Western blot analyses of full-length protein levels (D) and cleaved Tie1 (Cl Tie1) 45-kDa endodomain levels (E). (F) Western blot analysis of full-length Tie1 receptor after 24 hours of shear stress showed that laminar flow at 20 dynes/cm<sup>2</sup> (Lss20) decreased Tie1 expression. (G–K) Western blot analyses of Tie1-deleted (i.e., 4OHT-treated) versus untreated control cells, comparing laminar shear-induced eNOS expression (G), eNOS phosphorylation (H), ICAM expression (I), IkBα expression (J), and Tie2 phosphorylation (normalized to Tie2) (K). *P < 0.05.
In summary, we found that Tie1 was expressed in specific regions experiencing atherogenic shear stress, and reduction of Tie1 expression attenuated atherosclerosis progression in a shear stress–specific and dose-dependent manner. Our in vitro shear stress studies further suggest a proinflammatory role for Tie1, which we believe to be novel. This prophylactic alleviation of atherosclerosis burden by the reduction of Tie1 expression and signaling points to Tie1 as an attractive candidate for targeted therapeutic approaches.

Methods

Animal breeding and tamoxifen injections. Tie1fl/fl mice were a gift from M.C. Puri (University of Toronto, Toronto, Ontario, Canada; ref. 12), and SCL-ERCre mice were a gift from J.R. Gothert (University of Duisburg-Essen, Essen, Germany; ref. 33). Tie1fl/fl;Apoe−/−, Tie1<−/−;Apoe−/−, Tie1fl/fl;SCL-ERCre, and Tie1<−/−;SCL-ERCre;Apoe−/− mice were bred in our laboratory. All mice used in this study were bred onto a pure C57BL/6 background and were maintained in microisolator cages on a rodent chow diet (Purina Mills Inc.) and autoclaved water ad libitum. Animal care and experimental procedures were approved by the Institutional Animal Care and Use Committee of Vanderbilt University. Tamoxifen injections were performed on 6-week-old mice. Intraperitoneal injections of 2 mg tamoxifen (catalog no. T5648; Sigma-Aldrich) dissolved in 10% ethanol and 0.1 ml sunflower oil (catalog no. S0007; Sigma-Aldrich) were administered once every 2 days (7 injections total).

Generation of Tie1 mutant alleles. To generate Tie1fl/fl mice, a loxP site and a neomycin resistance cassette were introduced within the first intron of Tie1, and another loxP site was introduced just upstream of the minimal promoter region. This strategic placement of loxP sites would allow for excision of the Tie1 minimal promoter and exon 1 upon Cre activation (58).

Detection of β-galactosidase. Whole tissue was collected in HBSS; fixed in 0.2% glutaraldehyde, 2 mM MgCl2, and 5 mM EGTA solution; and stained in 1 mg/ml X-gal, 2 mM MgCl2, 5 mM EGTA, 0.01% deoxycholate, 0.02% NP-40, and 0.1 M phosphate buffer (pH 7.3) solution overnight at 30°C. The stained tissues were postfixed in 4% paraformaldehyde, cleared in a glycerol gradient, and photographed in 100% glycerol with a dissecting photomicroscope. Serial cryosections of aortic sinus adjacent to oil red O–stained sections were stained as described above, postfixed with 4% paraformaldehyde, dehydrated in ethanol, cleared in xylene, counterstained with eosin, and mounted in permount.

Analysis of aortic lesions. The extent of atherosclerotic lesions was evaluated by 2 alternate methods: cross-sections of the aortic sinus and the whole aorta pinned out en face (59). Aortas were flushed through the left ventricle, and the entire aorta was dissected and pinned out for en face analysis as described previously (59). In addition, to quantify atherosclerotic lesions in the aortic sinus, the heart was embedded in OCT medium and frozen. 5-μm cryosections of the aortic valve region were collected and processed for oil red O staining. Lesions in the aorta were analyzed using Imaging System KS 300 (Kontron Elektronik GmbH). The mean atherosclerotic lesion area for each animal was calculated as total lesion area relative to total surface area. Second, lesions were assessed according to 3 regions: (a) lesser curvature of the aortic arch; (b) greater curvature of the aortic arch, including the brachiocephalic trunk, left common carotid artery, and left subclavian artery; and (c) descending aorta, including the thoracic, abdominal, and lumbar regions. For each region, the lesion area was calculated as a percentage of total aorta surface area.

Serum lipids and lipoprotein distribution analysis. Serum total cholesterol and triglyceride levels were determined on samples obtained from mice fasted for 4 hours as described previously (32). Fast performance liquid chromatography was performed on an HPLC system model 600 (Waters) using a Superox 6 column (Pharmacia).

Immunocytochemistry. The localization of mouse macrophages in the arterial lesions was examined by immunocytochemistry using 5-μm cryosections of the aortic sinus fixed in acetone at 4°C. The sections were immersed in PBS (pH 7.2), blocked in goat serum dilution (containing 4% BSA and 2% goat serum), and incubated overnight at 4°C with monoclonal rat antibody MOMA-2 (Accurate Chemical & Scientific Corp.). The sections were treated with goat biotinylated antibody rabbit IgG (BD Biosciences—Pharmingen) for 45 minutes at 37°C, then incubated with avidin-biotin complex labeled with alkaline phosphatase (Vector Laboratories). Enzyme was visualized with Fast Red TR/Naphthol AS-NX substrate (Sigma-Aldrich) and counterstained with hematoxylin. Nonimmune rabbit or rat serum was used in the place of primary antibody as a negative control. Photomicroscopy was performed on a Zeiss Axioskop microscope with Plan-FLUAR objectives. Area of staining was normalized to total plaque area previously calculated for each mouse.

Shear stress–modifying cast. Casts were manufactured at the Vanderbilt Physics Machine Shop with medical grade polyetherketone (Invibio Inc.) using single lip micro cutters (provided by Alignment Tools Pte Ltd.). Casts with nonconstricting bores were used as controls. As described by Cheng et al., the tapered casts induce lowered shear stress (10 N/m2) immediately upstream, increased shear stress inside the cast and oscillatory shear stress with vortex flow immediately downstream (30, 31). The tapered region increases shear stress from 10 to 25 N/m2.

8 mice 12 weeks of age were assigned randomly to 2 groups. Shear stress in the right common carotid artery was altered by cast placement. Each animal was anesthetized with isoflurane, and the anterior cervical triangle was accessed by a sagittal anterior neck incision. Both halves of the cast were placed around the right common carotid artery and fixed with a suture. Wounds were closed, and the animals were allowed to recover. Animals with control and tapered casts were sacrificed 7 days after surgery.

Immunohistochemistry. Mouse carotid arteries were processed for whole-mount immunostaining with VE-cadherin after β-gal staining. Tissue samples were blocked in 10% goat serum at room temperature for 30 minutes and incubated with rabbit antibody to VE-cadherin (catalog no. 160840; Cayman Chemical) diluted in blocking solution at 4°C overnight.

Sections were treated with goat biotinylated antibody to rabbit IgG (BD Biosciences—Pharmingen) for 45 minutes at 37°C. Then, sections were incubated with avidin-biotin complex labeled with peroxidase (Vector Laboratories). Enzyme was visualized with 3,3′-diaminobenzidine (DAB) substrate (Sigma-Aldrich).

Protein isolation and Western blotting. Tissue samples or cells were lysed on ice using a lysis buffer (50 mM Tris, 50 mM NaCl, 1% Triton X-100, 1 mM EGTA, 1 mM Na2VO4, 5 mM NaF, and 1X β-glycerophosphate) supplemented with complete protease inhibitor cocktail (Roche). After centrifugation, the supernatant was collected. Protein concentrations were determined with Protein Assay Reagent (BioRad Laboratories). The nuclear pellet was rinsed 3 times in lysis buffer and incubated at 100°C with 2X sample buffer to extract nuclear proteins. Lysates were resolved by SDS-PAGE electrophoresis and transferred onto Hybond ECL nitrocellulose membranes (GE Healthcare). Blots were blocked with 5% nonfat milk and probed with rabbit antibodies to Tie1 (catalog no. sc342; Santa Cruz), NOS3 (catalog no. sc654; Santa Cruz), phospho-NOS3 (catalog no. sc12972; Santa Cruz), Tie2 (catalog no. sc342; Santa Cruz), ICAM (catalog no. sc1511; Santa Cruz), IkBα (catalog no. sc1643; Santa Cruz), phospho-Tie2 (catalog no. AF2720; R&D Systems), and mouse antibody to
β-actin (catalog no. A5316; Sigma-Aldrich) overnight at 4°C. After washing with 0.1% Tween-20–supplemented PBS (PBST), blots were incubated with secondary antibodies IRDye 800CW goat anti-rabbit IgG and IRDye 680CW goat anti-mouse IgG for 1 hour at room temperature with gentle agitation. Blots were washed and scanned on the Odyssey Infrared Imaging System (LI-COR Biosciences) and densitometry analyses were performed using Odyssey Software.

**MAEC isolation.** The immorto mouse is a transgenic mouse generated by the introduction of thermolabile SV40 TAg, tsAS8 (48). In this model, the SV40 large TAg of temperature-sensitive strain tsAS8 is fused with the major histocompatibility complex promoter H-2Kb. Expression of the promoter and the large TAg protein is only evident when immorto mouse-derived cells are cultured at a permissive temperature (33°C). The addition of IFN-γ was also used to further enhance promoter activity (48). A 4- to 8-week-old immorto mouse was anesthetized with isoflurane and sacrificed by cervical dislocation. The lumbar aorta was punctured, and 5 ml of HBSS was perfused through the left ventricle to flush out blood. The heart, lung, and aorta block was excised and transferred to a dish containing HBSS and antibiotics. The heart and lungs were removed. The rest of the aorta was incubated at 37°C for 15 minutes with filtered 10 mg/ml collagenase type II (Worthington Chemicals) and 1x antibiotics dissolved in dispase (Roche). The adventitia was gently removed, and the remaining intima layer comprising the endothelium and internal elastic lamina was cut out longitudinally and incubated at 37°C for 30 minutes in filtered 20 mg/ml collagenase type II solution and 1x antibiotics dissolved in dispase. The aorta was dissociated by pipetting several times in the dissolving solution and filtered through a 100 μm sterile cell strainer (Fisher Scientific). Culture media (MCDB 131 [Gibco, Invitrogen], 10% FBS [Hyclone], 10 U/ml heparin [Sigma-Aldrich], 2.75 mM hydrocortisone [Sigma-Aldrich], 0.2% bovine brain extract [Hammond Celltech], 10 U/ml recombinant murine IFN-γ [Peprotech], and 1x antibiotics [Gibco, Invitrogen]) was added to the cell suspension and centrifuged at 375 g for 10 minutes. The cell mixture was then plated onto a fibronectin coated culture dish and propagated at 33°C in a mixture of 5% carbon dioxide and 95% oxygen. MAECs were isolated from wild-type and Tie2ΔEC-SCL-ERCre immorto mice.

**Flow-assisted cell sorting.** MAECs were released from the plate by incubating with Accutase (Invitrogen) for 10 minutes at 37°C. Cells were collected by centrifugation at 375 g for 10 minutes at 4°C and resuspended in HBSS (supplemented with 1% FBS and 1x antibiotics). The cells were then incubated with APC-conjugated CD31 (catalog no. 551262; BD Biosciences) at 1 mg/ml x 10⁶ cells for 30 minutes on ice with occasional agitation. MAECs were washed 3 times in HBSS and filtered through a 40-μm cell strainer (Fisher Scientific). Propidium iodide (Invitrogen) was added at 1:1,000 dilution prior to sorting. CD31+ cells were collected in culture media, centrifuged, and plated onto collagen-coated (Rat tail collagen type I; BD Biosciences) dishes.

**MAEC characterization.** MAECs were rinsed with PBS, briefly fixed with methanol at –20°C, and blocked with 10% normal goat serum (Jackson ImmunoResearch) for 1 hour at room temperature. Immunostaining was performed at 4°C overnight with either CD31 (catalog no. 553370; BD Biosciences) or VE-cadherin (catalog no. 160840; Cayman Chemical) and counterstaining with goat anti-rat Alexa Fluor 488 (BD Biosciences) or goat anti-rabbit Alexa Fluor 488. MAECs were also cultured on Matrigel-coated (BD Biosciences) dishes overnight and observed for network formation. Dil-AcLDL (Biomedical Technologies) was added to the media, centrifuged, and plated onto collagen-coated (Rat tail collagen type I; BD Biosciences) dishes. MAECs were released from the plate by incubating with Accutase (Invitrogen) for 10 minutes at 37°C. Cells were collected in culture media (MCDB 131 [Gibco, Invitrogen], 10% FBS [Hyclone], 10 U/ml heparin [Sigma-Aldrich], 2.75 nM hydrocortisone [Sigma-Aldrich], 0.2% bovine brain extract [Hammond Celltech], 10 U/ml recombinant murine IFN-γ [Peprotech], and 1x antibiotics [Gibco, Invitrogen]) and treated with 5 μM 4OHTh (Sigma-Aldrich) every 2 days for 6 days. To ascertain ER²Cre nuclear localization, 4OHTh-treated cells were rinsed once with HBSS, fixed with cold 100% methanol for 3 minutes, and rinsed twice with HBSS. Cells were blocked with 5% goat serum, incubated overnight at 4°C with Cre antibody (Abcam, AB24608), rinsed, and immunostained with Alexa Fluor 488–conjugated goat anti-rabbit secondary antibody (BD Biosciences).

**Shear stress experiments.** Shear stress experiments were performed using a custom cone-and-plate shear stress viscometer design. An inverted servo motor (ElectroCraft) was attached to a plexiglass cone with 0.5° angle. A Motomatic II motor controller (Reliance Electric, ElectroCraft) regulated velocity in laminar flow, and a digital function generator (Instek) was used to produce a sinusoidal waveform modulating oscillatory flow. A digital oscilloscope (Tektronix) was used in parallel to monitor the output magnitude and waveform. The servomotor and cone were lowered onto the culture dish by a step-controlled base stand. Laminar shear was attained with unidirectional motion of the cone, while oscillatory shear was achieved by bidirectional motion at 5 dynes/cm². Shear stress experiments were performed for 24 hours in complete media in a sterile, 5% CO₂ incubator at 37°C.

**RNA extraction and real-time RT-PCR.** RNA from tissue or cells was isolated using TriReagent (Molecular Research Center), and SuperScript III First-Strand System (Invitrogen) was used to synthesize first-strand cDNA. Quantitative real-time RT-PCR was performed using iQ SYBR Green Supermix (BioRad) on the iQ5 Real Time PCR Detection System (BioRad). Experiments were performed in triplicate using glyceraldehyde-3-phosphate dehydrogenase as the housekeeping gene to normalize the data. The relative change in gene expression was determined using the critical threshold (Ct) and the calculation 2ΔΔCt. Primers were used as follows: Tie1, 5′-GTGCCACACTTTTGACACTG-3′ and 5′-CAGGCAAGCAGGTTGTAGA-3′; glyceraldehyde-3-phosphate dehydrogenase, 5′-CAGTGGCATGCCCCTCGGT-3′ and 5′-AGGAAATGACCTTGACAAAG-3′; VCAM, 5′-AAAGAGACACCCAGTGAGG-3′ and 5′-GGGGCAAGTGTGCAATAAG-3′; ICAM, 5′-GAAGGTGTTCTTCTGACGC-3′ and 5′-GGGCGTGGCAGAGTTGAACTATA-3′; Nos3, 5′-TGAGATCTCTGCTTCACATC-3′ and 5′-AGTTCTAGAGGACCTACACAAAGTGGT-3′.

**Statistics.** All statistical differences in this study were determined by a 2-tailed, unpaired Student’s t test with 95% confidence intervals, using Prism 4.0 software (Graphpad Inc.). Statistical significance was accepted for P values less than 0.05. Graphical data representations are presented as mean ± SEM.

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Address correspondence to: H. Scott Baldwin, 9435-A MRB IV, Vanderbilt University Medical Center, Nashville, Tennessee 37232-0493, USA. Phone: 615.322.2703; Fax: 615.322.6541; E-mail: scott.baldwin@vanderbilt.edu.


