CD163⁺ macrophages promote angiogenesis and vascular permeability accompanied by inflammation in atherosclerosis

Liang Guo, … , Renu Virmani, Aloke V. Finn

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Intake of hemoglobin by the hemoglobin-haptoglobin receptor CD163 leads to a distinct alternative non–foam cell antiinflammatory macrophage phenotype that was previously considered atheroprotective. Here, we reveal an unexpected but important pathogenic role for these macrophages in atherosclerosis. Using human atherosclerotic samples, cultured cells, and a mouse model of advanced atherosclerosis, we investigated the role of intraplaque hemorrhage on macrophage function with respect to angiogenesis, vascular permeability, inflammation, and plaque progression. In human atherosclerotic lesions, CD163⁺ macrophages were associated with plaque progression, microvascularity, and a high level of HIF1α and VEGF-A expression. We observed irregular vascular endothelial cadherin in intraplaque microvessels surrounded by CD163⁺ macrophages. Within these cells, activation of HIF1α via inhibition of prolyl hydroxylases promoted VEGF-mediated increases in intraplaque angiogenesis, vascular permeability, and inflammatory cell recruitment. CD163⁺ macrophages increased intraplaque endothelial VCAM expression and plaque inflammation. Subjects with homozygous minor alleles of the SNP rs7136716 had elevated microvessel density, increased expression of CD163 in ruptured coronary plaques, and a higher risk of myocardial infarction and coronary heart disease in population cohorts. Thus, our findings highlight a nonlipid-driven mechanism by which alternative macrophages promote plaque angiogenesis, leakiness, inflammation, and progression via the CD163/HIF1α/VEGF-A pathway.

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CD163+ macrophages promote angiogenesis and vascular permeability accompanied by inflammation in atherosclerosis

Liang Guo,1 Hirokuni Akahori,2 Emanuel Harari,1 Samantha L. Smith,1 Rohini Polavarapu,2 Vinit Karmali,2 Fumiyuki Otsuka,1 Rachel L. Gannon,1 Ryan E. Braumann,1 Megan H. Dickinson,1 Anuj Gupta,3 Audrey L. Jenkins,4 Michael J. Lipinski,4 Jinhoon Kim,5 Peter Chhour,5 Paul S. de Vries,6 Hiroyuki Jinnouchi,1 Robert Kutys,1 Hiroyoshi Mori,1 Matthew D. Kutyna,1 Sho Torii,1 Atsushi Sakamoto,1 Cheol Ung Choi,2 Qi Cheng,1 Megan L. Grove,4 Mariem A. Sawan,3 Yin Zhang,2 Yihai Cao,7 Frank D. Kolodgie,1 David P. Cormode,3 Dan E. Arking,6 Eric Boerwinkle,6,9 Alanna C. Morrison,6 Jeanette Erdmann,10 Nona Sotoodehnia,11 Renu Virmani,1 and Alok V. Finn1,3

Inflammation in atherosclerosis and vascular permeability accompanied by

School of Medicine, University of Pennsylvania, Philadelphia, Pennsylvania, USA. 6Human Genetics Center, Department of Epidemiology, Human Genetics, and Environmental Sciences, School of Public Health, The University of Texas Health Science Center at Houston, Houston, Texas, USA. 7Department of Microbiology, Tumor and Cell Biology, Karolinska Institute, Stockholm, Sweden. 8McKusick-Nathans Institute of Genetic Medicine, Johns Hopkins University School of Medicine, Baltimore, Maryland, USA. 9Human Genome Sequencing Center, Baylor College of Medicine, Houston, Texas, USA. 10Institute for Cardiogenetics, University of Lübeck, and German Centre for Cardiovascular Research (DZHK), Partner Site Hamburg/Kiel/Lübeck, Germany. 11Division of Cardiology, Department of Medicine and Epidemiology, Cardiovascular Health Research Unit, University of Washington, Seattle, Washington, USA.

Intake of hemoglobin by the hemoglobin-haptoglobin receptor CD163 leads to a distinct alternative non–foam cell antiinflammatory macrophage phenotype that was previously considered atheroprotective. Here, we reveal an unexpected but important pathogenic role for these macrophages in atherosclerosis. Using human atherosclerotic samples, cultured cells, and a mouse model of advanced atherosclerosis, we investigated the role of intraplaque hemorrhage on macrophage function with respect to angiogenesis, vascular permeability, inflammation, and plaque progression. In human atherosclerotic lesions, CD163+ macrophages were associated with plaque progression, microvascularity, and a high level of HIF1α and VEGF-A expression. We observed irregular vascular endothelial cadherin in intraplaque microvessels surrounded by CD163+ macrophages. Within these cells, activation of HIF1α via inhibition of prolyl hydroxylases promoted VEGF-mediated increases in intraplaque angiogenesis, vascular permeability, and inflammatory cell recruitment. CD163+ macrophages increased intraplaque endothelial VCAM expression and plaque inflammation. Subjects with homozygous minor alleles of the SNP rs7136716 had elevated microvessel density, increased expression of CD163 in ruptured coronary plaques, and a higher risk of myocardial infarction and coronary heart disease in population cohorts. Thus, our findings highlight a nonlipid-driven mechanism by which alternative macrophages promote plaque angiogenesis, leakiness, inflammation, and progression via the CD163/HIF1α/VEGF-A pathway.

Introduction

Macrophages play a critical role in atherogenesis but are a heterogeneous group of immune cells (1). The atherosclerotic plaque microenvironment is an important stimulus driving their differentiation into distinct cell types that undoubtedly influence plaque evolution. Our understanding of how these stimuli affect macrophage behavior and in turn influence atherosclerosis remains incomplete (2).

Related Commentary: p.910

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Conflict of interest: The authors have declared that no conflict of interest exists.
which is thought to result from plaque neovascularization and increased microvessel permeability (6, 7). These features are considered critical elements of advanced atherosclerotic plaques and are linked to plaque progression through the accumulation of erythrocyte membranes, which are rich in free cholesterol and promote necrotic core enlargement (8).

Within areas of IPH, oxidative stress leads to erythrocyte lysis, which releases free hemoglobin (Hb) (9). Hb is quickly bound by the plasma protein haptoglobin, and hemoglobin:haptooglobin (HH) complexes are internalized into the macrophage via the CD163 receptor exclusively expressed on cells of this lineage (10, 11). We and others have previously shown that intake of HH by CD163 leads to a distinct phenotype of macrophage termed M(Hb) (6) (also referred to as Mcem) found in areas of neoangiogenesis and hemorrhage characterized by high surface expression of CD163, reduced proinflammatory cytokine production, and a lack of lipid retention, all characteristics which distinguish them from foamy macrophages (6, 12). On the basis of these findings, the M(Hb) macrophage has been termed atheroprotective (12, 13). However, the association of M(Hb) macrophages with areas of intraplaque angiogenesis and permeability raises the important question of whether these cells are merely bystanders or play an active role in these processes.

An abundance of extracellular ferric iron and lowered ROS formation are important features defining areas in which M(Hb) macrophages are found (6). We have previously shown that increased export of iron through upregulation of the iron exporter ferroportin (FPN) lowers intracellular free iron within M(Hb) macrophages (6, 14). A relationship between iron and the proangiogenic transcription factor HIF1α can be found through its interactions with the prolyl hydroxylase proteins (PHDs). Under conditions of normoxia or hypoxia, HIF1α becomes hydroxylated by PHDs on proline residues 402 and 564, within its oxygen-dependent domain. This allows it to be recognized by the von Hippel–Lindau tumor-suppressor protein (pVHL), which targets it for ubiquitin-mediated degradation (15). While hypoxia is a well-known activator of HIF1α, iron itself also indirectly controls the activation of HIF1α by serving as an essential cofactor for the activity of PHDs (16). Although 3 HIF prolyl hydroxylases (PHD1, PHD2, PHD3) mediate the hydroxylation of proline residues on HIF1α, PHD2 is responsible for more than 95% of the activity in cultured cells (17). PHD2 hydroxylase activity is stimulated by the addition of iron in vitro, and its activity is inhibited by iron chelators (16). Thus, changes in iron metabolism within M(Hb) macrophages might affect their angiogenic potential via inhibition of PHDs.

Here, we reveal an important molecular pathway within alternatively polarized M(Hb) macrophages that promotes intraplaque angiogenesis, permeability, inflammatory cell recruitment, and plaque progression in human atherosclerosis. Intracellular iron deprivation within M(Hb) macrophages inhibits PHD2 activity, which leads to the accumulation of HIF1α and VEGF-A. Using human atherosclerotic specimens and in vitro assays with human cells, we demonstrate that M(Hb) macrophages are associated with intraplaque angiogenesis, defective endothelial junctional protein expression, and increased vascular inflammation mediated through VEGF-A-VEGFR2 interactions. Genetic deletion of CD163 in a mouse model of IPH reduced plaque permeability, progression, and inflammatory cell recruitment. These findings demonstrate an unexpected but important proatherogenic role for M(Hb) macrophages in the response to IPH by promoting plaque angiogenesis and permeability. Moreover, these studies point toward the HIF1α/VEGF-A system as a potential target for antitherosclerotic therapies.

**Results**

**Human carotid plaque pathological characterization, plaque progression, and association with CD163+ macrophages.** A total of 60 atherosclerotic samples from patients undergoing surgical carotid endarterectomy (CEA) were available in the CVPath Registry (18). A detailed history was not included for 22 of these samples, and they were therefore excluded. Thirty-eight specimens containing culprit segments along with proximal and distal (flanking) segments were available for analysis and assessed for plaque morphology. We classified 114 plaques (38 culprit, 38 proximal flanking, and 38 distal flanking plaques) and performed IHC to assess each one for the area occupied by CD163, a marker of M(Hb) macrophages. We found a strong positive correlation between CD163+ macrophage area and plaque progression (i.e., pathological intimal thickening [PIT] or intimal xanthoma, fibroatheroma or thin-cap fibroatheroma (TCFA), and plaque rupture or healed rupture) (Figure 1, A and B). The correlation with plaque progression could also be demonstrated by stratifications of stenosis and necrotic area percentage (Figure 1, C and D). These results suggest a strong association between CD163 expression in alternative macrophages and atherosclerotic disease progression.

**Human carotid plaque assessment of angiogenesis in high and low CD163/CD68 plaque areas.** We also used IHC to assess each plaque for the area occupied by the general macrophage marker CD68. Areas were separated into low and high CD163 expression on the basis of their CD163/CD68 ratio (low CD163/CD68 <0.20 vs. high CD163/CD68 <0.70). (Foam cells predominated in the low CD163 areas.) We determined the extent of microvesselarity in each of these areas using antibodies against CD31 and vWF. Table 1 shows that the total content of macrophages in each area (as assessed by CD68 expression) was not significantly different between groups. The microvessel number per unit area, however, was significantly greater within high CD163 areas (Table 1, see also representative images in Figure 1, E and F). Because most microvessels were seen around CD163+ M(Hb) cells both within low and high CD163 plaque areas, we further subdivided each area into CD163+ and CD163- regions for a local effect assessment in the plaque. Regardless of a high or low CD163/CD68 ratio, we detected greater numbers of microvessels in regions that had CD163+ macrophages (CD163+ area in Table 1). This suggests that CD163+ macrophages are associated with plaque angiogenesis.

**Alternative CD163+ macrophages express HIF1α and VEGF-A and are associated with intraplaque angiogenesis and vascular permeability.** We next evaluated the expression of the proangiogenic transcription factor HIF1α and its transcriptional target VEGF-A using similar methodology. High CD163 areas demonstrated heavy expression of HIF1α and VEGF-A compared with CD163 low/foam cell-rich areas (Figure 2A). HIF1α was predominantly located within CD163+ macrophages. To confirm our findings, protein was extracted from high CD163 plaques and compared with protein...
Inhibition of PHDs by relative iron deprivation within M(Hb) cells increases HIF1α/VEGF-A and promotes angiogenesis. We previously demonstrated that human plaque areas rich in M(Hb) cells had reduced ROS, signs of oxidative damage, and an abundance of ferric iron, consistent with iron release from macrophages (6). CD163+ M(Hb) cells can be reproduced in vitro by differentiation of human monocytes over 5 to 7 days in HH-enriched media (6). Hb/iron intake into macrophages causes an increase in the iron-binding protein ferritin and upregulation of the iron exporter FPN, which results in significant lowering of intracellular free iron (6, 22). By laser capture microdissection, lesion areas with high levels of CD163+ macrophages (high CD163) in human carotid plaques showed elevated FPN expression compared with areas containing low levels of CD163+ macrophages (low CD163) (Figure 2G). Consistent with this, HH-differentiated macrophages had lower levels of intracellular free iron (Figure 3A). Because PHD2 function is dependent on iron as a cofactor, we examined PHD2 activity using cellular lysates from human macrophages differentiated in HH-enriched media. Lysates were exposed to a peptide derived from the HIF1α oxygen-dependent degradation domain, and hydroxylation was measured using an antibody specific for hydroxylated HIF1α (Pro564). We found that PHD2 activity was significantly lower in HH-differentiated cells (Figure 3B). To confirm our results, we measured hydroxylated HIF1α (HIF1α-Oh) levels in control and HH-differentiated macrophages in the presence of 10 μM MG132, a proteasome inhibitor, using an antibody against HIF1α-Oh. We found that HIF1α-Oh levels were significantly lower in HH-differentiated cells (Figure 3C). There were no changes in the total amount of PHD2 levels by Western blotting in control or HH-differentiated cells (Figure 3D). Next, to confirm relative iron deprivation as the cause of elevated HIF1α levels, we used 2 different approaches. First, we modulated intracellular iron by treating HH-differentiated cells with hepcidin, a peptide whose major function is to degrade FPN (23). Heparin given 24 hours before harvesting was able to increase intracellular iron within HH-differentiated macrophages but had no effect on control cells (Supplemental Figure 1A; supplemental material available online with this article; https://doi.org/10.1172/JCI93025DS1). HIF1α levels were significantly increased in HH-differentiated cells and dramatically decreased to control levels after hepcidin treatment, while there was no effect on control-treated cells (Figure 3E). We observed similar results for VEGF-A levels, which were determined by ELISA analysis of supernatants of these cells (Figure 3F). HIF1α levels were also decreased in HH-differentiated cells when iron (10, 50, or 100 μM FeCl3) was added directly to them 24 hours before harvesting (Supplemental Figure 1B). Last, to confirm the angiogenic ability of M(Hb) cells, supernatants from control and HH-treated macrophages, with or without added hepcidin, were evaluated in an in vitro Matrigel angiogenesis assay using human aortic endothelial cells (HAECs). Tube formation was increased by more than 2-fold with HH supernatants, but was not significantly changed in the presence of hepcidin (Figure 3G). These data suggest that relative iron deprivation within M(Hb) cells inhibits PHD2 activity to increase HIF1α-mediated angiogenesis.

CD163+ M(Hb) cells promote vascular permeability via VEGF-A/VEGFR2 signaling. HIF1α accumulation within M(Hb) cells drives the transcription of VEGF-A, a key regulator of endothelial per-
HAECs using transendothelial electrical resistance (TEER) and control or HH-differentiated macrophages on the permeability of VE-cadherin. To determine whether M(Hb) cells promote vascular permeability (24). This results in rapid internalization and degradation of and promotes specific tyrosine phosphorylation of VE-cadherin by its major ligand VEGF-A leads to the formation of a complex between endothelial cells, which contrasted with HH-differentiated macrophage supernatants. There was distinct expression of VE-cadherin in membrane protein extractions (Figure 4, C and D). We confirmed this result by immunoblotting for VE-cadherin in membrane protein extractions (Figure 4, E and F). Collectively, these results suggest that M(Hb) macrophages promote endothelial permeability via VEGF-A/VEGFR2 signaling.

Deletion of CD163 in mice reduces intraplaque neovascularization and plaque progression. To explore the role of Hb in phenotypic modulation of macrophage behavior with respect to angiogenesis, we isolated peritoneal macrophages from CD163-knockout mice we had previously generated (25). Our published data as well as those of others suggest that mice can support a M(Hb) phenotype similar to that of humans (14, 26, 27). Exposure of peritoneal macrophages to mouse Hb (0.1 mg/ml) overnight resulted in significant upregulation of HIFα by Western blotting and release of VEGF-A (Figure 5, A and B) in WT but not CD163–/– mice. (Ingestion of Hb by mouse CD163–/– macrophages does not seem to be as dependent on haptoglobin binding as human CD163 is [ref. 27].) Moreover, treatment of mouse dermal endothelial cells (MDECs) with supernatants from WT mouse peritoneal macrophages exposed to Hb resulted in significant increases in tube formation in an in vitro Matrigel angiogenesis assay (Supplemental Figure 2), a decrease in membrane VE-cadherin expression (Supplemental Figure 3), and an increase in membrane permeability (Supplemental Figure 4). There was no effect of supernatants from Hb-exposed, CD163-deficient macrophages in any of these assays with respect to these endpoints (Supplemental Figures 2–4). Collectively, these results suggest that CD163 mediates the proangiogenic and pro-permeability effects of Hb on macrophages.

To determine whether Hb is an important modulator of macrophage-mediated angiogenesis and permeability in the setting of atherosclerosis, we crossed CD163–/– mice with atherosclerosis-prone apolipoprotein E-deficient (ApoE–/–) mice (both on a C57BL6 background). Because IPH is not typically observed in the aortic sinus after high-fat diet feeding in ApoE–/– mice, we examined the brachiocephalic arteries (BCAs) of aged, Chow-fed mice, which reproducibly develop IPH (28). While we detected no differences in the aortic root plaque lesion areas in ApoE–/– CD163–/– versus Apo E–/– mice (Supplemental Figure 5), in 1-year-old ApoE–/– CD163–/– double-knockout mice, we observed a significant reduction in BCA plaque lesion area and intraplaque neovascularization and hemorrhage (by immunostaining using antibodies against VE-cadherin and Ter-119, respectively) (Figure 5, C–E, H, and I, by confocal microscopy, was reduced by more than 50% in ApoE–/– CD163–/– mice, had significantly less BCA plaque lesion area and intraplaque neovascularization and hemorrhage (by immunostaining using antibodies against VE-cadherin and Ter-119, respectively) (Figure 5, C, E, H, and I, and Supplemental Figure 9). One-year-old ApoE–/– and ApoE–/– CD163–/– mice had similar body weights, plasma cholesterol, glucose, and insulin levels (Supplemental Figure 6), and total numbers and types of WBCs (Supplemental Figure 7), as well as similarities in various measures of iron status (Supplemental Figure 8). CD163–/– macrophages were routinely seen in areas of angiogenesis within BCA plaques from ApoE–/– mice (Figure 5H). In comparison with ApoE–/– mice, CD163–/–ApoE–/– mice had significantly less BCA plaque complexity (Figure 5F), smaller necrotic areas (Figure 5G), and greater fibrous cap thickness (Supplemental Figure 10). To measure intraplaque neovascular permeability, FITC-dextran was administered i.v. We found that intraplaque FITC-dextran, quantified by confocal microscopy, was reduced by more than 50% in ApoE–/– CD163–/– mice versus ApoE–/– controls, consistent with the lower microvessel density observed in ApoE–/– CD163–/– mice (Figure 5, I and J). These results collectively indicate that Hb and CD163 in particular are critical mediators of macrophage-mediated angiogenesis, vascular permeability, and plaque progression.

<table>
<thead>
<tr>
<th>Table 1. Macrophage content and angiogenesis in human carotid plaques</th>
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<tr>
<td><strong>Macrophages</strong></td>
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<tr>
<td>-----------------</td>
</tr>
<tr>
<td>CD163/CD68 (ratio)</td>
</tr>
<tr>
<td>Total CD163 area (mm²)</td>
</tr>
<tr>
<td>% Area positive for CD163</td>
</tr>
<tr>
<td>Total CD68 area (mm²)</td>
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<tr>
<td>% Area positive for CD68</td>
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<tr>
<td><strong>Angiogenesis</strong></td>
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<tr>
<td>Microvessels/plaque area (n/mm²)</td>
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Results are presented as the mean ± SD (n = 10 plaques per group). All P values were determined by 2-sided Student’s t test. *P < 0.01.
**Microvascular endothelial cells surrounding CD163+ macrophages have elevated VCAM expression.** VCAM has been shown to be critical in the development of atherosclerosis in experimental models, because it promotes inflammatory cell entry into the vessel wall (29). Intraplaque neovessels have been previously shown to express VCAM, which was associated with increased macrophage density (30). We hypothesized that CD163+ macrophages might also promote leukocyte entry into the plaque via VEGF-mediated increases in VCAM expression on intraplaque microvessels. HAECs treated with supernatant collected from macrophages differentiated with HH showed a greater phosphorylated NF-κB (p−NF-κB) (p65) to total NF-κB ratio (Figure 6, A and C) compared with those treated with supernatant from control macrophages. When cells were pretreated with the NF-κB inhibitor BAY 11-7082, VCAM upregulation by CD163+ M(Hb) macrophage supernatants was blocked (Supplemental Figure 11). Since VEGF is known to be able to upregulate VCAM in endothelial cells via activation of NF-κB (31), we examined whether VEGFR2 knockdown in the endothelial cells treated with control supernatants versus CD163+ M(Hb) macrophage supernatants could attenuate this inflammatory response. Indeed, knockdown of VEGFR2 by siRNA in cultured human endothelial cells blocked the upregulation of VCAM and NF-κB in endothelial cells treated with M(Hb) supernatants (Figure 6, G–I). Moreover, treatment of MDECs with supernatants from WT mouse peritoneal macrophages exposed to Hb resulted in a significant increase in VCAM expression; however, there was no effect of supernatants from Hb-exposed, CD163-deficient (CD163−/−) macrophages (Supplemental Figure 12).

To confirm that VEGF mediated the increase in intraplaque endothelial VCAM expression and macrophages recruitment to the plaque, we performed studies in mice using a VEGF-blocking antibody and immunohistochemical analyses for VE-cadherin and VCAM colocalization and macrophage tracing. Six- to eight-month-old ApoE−/− mice were given a VEGF-blocking antibody or a control antibody via i.p. administration for 4 weeks. At the end of the antibody treatment, we injected gold-nanoparticle–labeled (Au-nanoparticle–labeled) monocytes i.v., measured monocyte infiltration into BCA plaques in serial histological sections, and performed immunohistochemical analyses using antibodies against VE-cadherin and VCAM. The results showed less VE-cadherin and VCAM colocalization in plaques from ApoE−/− mice treated with a VEGF-blocking antibody than was seen in plaques treated with the control antibody (Supplemental Figure 13) and a decrease in the number of labeled monocytes in BCA plaques after VEGF-blocking antibody treatment (Figure 6J). A third group of ApoE−/− CD163−/− mice were also treated with a control antibody, and few or no labeled monocytes could be found in the BCA plaques (Figure 6).

We next evaluated the severity and distribution of inflammation in the human plaques by histological scoring. Macrophage areas with high CD63 expression were associated with significantly higher inflammation scores (Figure 6, K and L). Confocal immunofluorescence showed that the microvessels surrounding high CD163+ macrophages were associated with more CD3+ T cells (Figure 6, M and N). These mouse and human data suggest a critical role for CD163+ macrophages in promoting microvascular inflammation and leukocyte recruitment.

**CD163+ macrophages are associated with increased angiogenesis, microvessel permeability, and intraplaque endothelial inflammation in human coronary artery plaques.** To further explore the proangiogenic and pro-permeability roles of CD163+ macrophages in human coronary atherosclerotic disease, Evans blue dye (EBD) (complexed with albumin) staining, a commonly used method to determine microvascular permeability, was used to perfuse intact and freshly collected human coronary artery samples with evidence of advanced atherosclerosis. Cross-sectional examination of EBD-perfused coronary arteries revealed distinct areas of tissue staining within plaques (Figure 7A). Heavily EBD-stained areas were colocalized with intraplaque microvessels surrounded by CD163+ macrophages, as identified by immunofluorescence and confocal imaging (Figure 7A, positive area 1). VCAM was also shown to be colocalized with microvascular endothelial cells by VE-cadherin staining (Figure 7A, positive area 2). Quantitation of microvessels, CD163+ macrophages, and VCAM in EBD-positive versus EBD-negative areas showed significantly higher microves sel density, more CD163+ macrophages, and upregulated VCAM expression in EBD-positive versus EBD-negative areas in the plaque (Figure 7, B–D). These results suggest that areas of high microvascular permeability in human coronary artery lesions are associated with the presence of CD163+ macrophages and increased expression of endothelial VCAM.

Last, SNPs in the CD163 gene were examined in a cohort of 346 African American patients from the CVPath Registry. Among the SNPs examined, homozygous G-allele carriers of rs7136716, an upstream intergenic variant that is reported to be associated with serum creatine kinase levels (32), were found to be significantly associated with coronary plaque rupture in a recessive model (odds ratio [OR] = 2.04, 95% CI, 1.01–4.09, P < 0.05) (Figure 7E, Table 2, and see Supplemental Table 1 for patients’ characteristics). (Individuals of European descent have a lower prevalence of this SNP, with a minor allele frequency [MAF] of 11%–12% vs. approximately 50% for individuals of African American descent.) To further examine the effect of the homozygous minor alleles in ruptured coronary plaques, 47 individuals with plaque rupture carrying homozygous major versus minor allele genotypes were selected for microvasculature analysis. Measurement of microvessel density at the rupture site of the coronary artery revealed that subjects with a homozygous minor allele genotype (GG genotype carriers) had significantly more angiogenesis than did those with a homozygous major allele genotype (AA genotype carriers) (Figure 7F). We found that the traditional cardiovascular risk factors were not different between these 2 groups (see patients’ characteristics).
Figure 7. Alternative CD163+ macrophages are associated with intraplaque angiogenesis and vascular permeability and express HIF1α and VEGF-A.

Human plaques from CEA specimens were analyzed by histology and immunofluorescence. Images were acquired by confocal microscopy using ×40 (A and E) or ×60 (C, RNAscope) objectives, with optical slicing in the z axis. (A) Representative microvessels (MV) from low and high CD163 areas (green) dual immunostained for HIF1α and VEGF-A (red channels). Scale bars: 20 μm. (B) immunoblotting for CD163, HIF1α, VEGF-A, and GAPDH of protein extracted from human atheroma expressing high or low levels of CD163 (n = 4 samples each). Bar graphs show quantitation of densitometry for the indicated proteins. (C) RNAscope ISH analysis using CD163 (red) and VEGF (green) probes on human carotid plaques. H&E and CD163 immunohistochemically stained images of human carotid plaques are shown on the left to indicate the areas (red boxes) of RNAscope ISH images of CD163 and VEGF, shown on the right. Scale bars: 1 mm (H&E- and immunostained images) and 2 μm (ISH). (D) Quantitative graph of VEGF RNA punctate counts in macrophages with low (<1-4 punctates) or high (≥5 punctates) CD163 expression (total of 3 plaques used for data collection, with 5 to 6 areas per plaque). (E) representative microvessels from low and high CD163 areas (green) dual immunostained for VE-cadherin and vWF (red channels). Note that VE-cadherin expression is located exclusively along the interendothelial contacts. The fluorescence signal for VE-cadherin appeared to be attenuated in a microvessels surrounded by CD163+ macrophages, with diffuse expression of vWF antigen, suggestive of leaky endothelial junctions. Scale bars: 20 μm. (F) Bar graphs show quantification of fluorescence signals for VE-cadherin and vWF in low and high CD163 areas (multiple areas from a total of 10 plaques per group were examined; see Methods). (G) Laser capture microdissection and qPCR analysis of FPN expression in low CD163+ and high CD163+ macrophage areas (n = 15 plaques sampled). Results are presented as the mean ± SEM (B and F) or the mean ± SD (D and G). (B and F) *P < 0.05, by 2-sided Student’s t test. (D and G) P < 0.01 and P < 0.05, by 2-sided Student’s t test.

Discussion

IPH and macrophages play important roles in the development of advanced atherosclerotic plaques. However, our understanding of the mechanisms by which they do so remains incomplete. Within this context, it seems natural to assume that within areas of IPH, the release of free Hb and its uptake by macrophages would protect against its deleterious effects by suppressing iron-derived toxicity. Indeed, previous work by our group and others has demonstrated that Hb is an important stimulus for alternative macrophage differentiation into a non–foam cell phenotype with antioxidative properties and reduced inflammatory cytokine expression that some have termed atheroprotective (6, 12, 13). Here, we show the unexpected but important finding that, within M(Hb) macrophages, relative iron deprivation inhibited the function of PHD2, leading to increased HIF1α-mediated angiogenesis, vascular permeability, and microvascular inflammation — features seen in advanced atherosclerotic plaques. These results suggest that the macrophage inflammatory response to hemorrhage actually promotes plaque destabilization by initiating a vicious cycle that likely leads to further intraplaque bleeding, inflammatory cell recruitment, and necrotic core expansion (Figure 8). Consistent with this, mice with combined deletion of CD163 and ApoE had significantly reduced plaque progression as compared with ApoE-deficient mice. Our findings highlight a nonlipid-driven mechanism by which alternative macrophages promote human plaque progression and indicate that HIF1α/VEGF-A is an important target for antiatherosclerotic therapies.

Wartman proposed more than 75 years ago that IPH might be a major contributor to the progression of coronary artery lesions (33). We and others have previously shown that IPH is a critical event in atherogenesis that fundamentally alters the plaque microenvironment by increasing the content of red cell membrane–derived free cholesterol causing necrotic core expansion and plaque progression (8). Studies involving the injection of silicon polymer into ath- erosclerotic human coronary arteries have revealed an elaborate microvascular network extending from the adventitia through the media and into the thickened intima (34). IPH is believed to arise from the disruption of thin-walled microvessels that are lined by discontinuous endothelium without supporting smooth muscle cells (19). However, the means by which such extensive angiogenesis occurs within intraplaque vessels has remained uncertain.

Table 2. Homozygous carriers of the minor allele of the CD163 SNP rs7136716 versus noncarriers

<table>
<thead>
<tr>
<th>SNP number</th>
<th>SNP type</th>
<th>Total minor allele frequency (n = 346 pts)</th>
<th>Homozygous minor allele in ruptures (n = 51 pts)</th>
<th>Homozygous minor allele in nonruptures (n = 295 pts)</th>
<th>P value</th>
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<tr>
<td>rs7136716</td>
<td>Upstream intron</td>
<td>43% (G allele)</td>
<td>14 (27.5%)</td>
<td>46 (15.6%)</td>
<td>0.046</td>
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Homozygous carriers of the minor allele of the CD163 SNP rs7136716 versus noncarriers were evaluated in plaque rupture versus nonrupture cases in the CVPath cohort (n = 346), pts, patients. The P value was determined by Fisher’s exact test to examine the association of the genetic variant with the plaque rupture.

in Supplemental Table 2. Significantly higher levels of CD163 mRNA were found in the ruptured coronary arteries of the homozygous G allele carriers (Figure 7G). To confirm the significance of rs7136716 in a large, independent cohort, 3,364 participants of African American ancestry from the Atherosclerosis Risk in Communities (ARIC) study were examined for genotypic and phenotypic data. A total of 3,225 subjects free of prevalent coronary heart disease (CHD) at baseline were included in the analysis. Baseline characteristics for the 3,225 participants with genotypic data and free of prevalent CHD at baseline are shown in Supplemental Table 3. A total of 1,311 incident CHD events occurred during follow-up, of which 338 were myocardial infarctions (MI). The mean follow-up period was 18.4 years for MI patients and 12.8 years for CHD patients. The rs7136716 variant indicated significant associations with both incident MI and incident CHD (hazard ratio = 1.21, 95% CI, 1.05–1.40, P = 0.0054, and hazard ratio = 1.20, 95% CI, 1.05–1.35, P = 0.0026, respectively) (Figure 7H). Together, these human data suggest that the rs7136716 variant in CD163 is a possible risk factor of CHD in patients of African American ancestry.
changes in macrophage iron metabolism that lead to relative intracellular iron deprivation and inhibition of iron-dependent PHDs. This causes an accumulation of HIF1α, which mediates angiogenesis through increased VEGF. While hypoxia may be an additional and important factor, our data clearly demonstrate that Hb intake by human macrophages under conditions of normoxia drives their differentiation into proangiogenic cells. Further supporting our contention is the fact that genetic deletion of CD163, a macrophage-specific protein, resulted in a significant reduction in angiogenesis (P < 0.05, Figure 5, H and I) and plaque progression (P < 0.01, Figure 5, C–G) in our mouse model. Interestingly, a recent study by Marsch and colleagues showed that PHD1-knockout mice have a metabolic phenotype that is generally deemed protective for cardiovascular disease (36). Additional changes in macrophage iron metabolism that lead to relative intracellular iron deprivation and inhibition of iron-dependent PHDs. This causes an accumulation of HIF1α, which mediates angiogenesis through increased VEGF. While hypoxia may be an additional and important factor, our data clearly demonstrate that Hb intake by human macrophages under conditions of normoxia drives their differentiation into proangiogenic cells. Further supporting our contention is the fact that genetic deletion of CD163, a macrophage-specific protein, resulted in a significant reduction in angiogenesis (P < 0.05, Figure 5, H and I) and plaque progression (P < 0.01, Figure 5, C–G) in our mouse model. Interestingly, a recent study by Marsch and colleagues showed that PHD1-knockout mice have a metabolic phenotype that is generally deemed protective for cardiovascular disease (36). Additional changes in macrophage iron metabolism that lead to relative intracellular iron deprivation and inhibition of iron-dependent PHDs. This causes an accumulation of HIF1α, which mediates angiogenesis through increased VEGF. While hypoxia may be an additional and important factor, our data clearly demonstrate that Hb intake by human macrophages under conditions of normoxia drives their differentiation into proangiogenic cells. Further supporting our contention is the fact that genetic deletion of CD163, a macrophage-specific protein, resulted in a significant reduction in angiogenesis (P < 0.05, Figure 5, H and I) and plaque progression (P < 0.01, Figure 5, C–G) in our mouse model. Interestingly, a recent study by Marsch and colleagues showed that PHD1-knockout mice have a metabolic phenotype that is generally deemed protective for cardiovascular disease (36). Additional
studies are needed to understand the differential function of each PHD isoform in the context of atherosclerosis.

Earlier work by our group and that of others has shown that microvessels within advanced atherosclerotic plaques exhibit compromised endothelial cell morphology and integrity characterized by loss of VE-cadherin (19, 20). However, the causes for increased vascular permeability with human atherosclerosis have remained unclear. Bobryshev and colleagues previously reported that while VE-cadherin was readily detectable in areas of human intraplaque vessels lacking inflammatory cells, it was irregularly distributed in vessels surrounded by large numbers of inflammatory cells (20). Our data indicate that VEGF-A secreted from CD163+ macrophages is the cause of these abnormalities. Supernatants from HH-differentiated M(Hb) macrophages caused significant increases in vascular permeability ($P < 0.05$, Figure 4, A and B) and VE-cadherin internalization ($P < 0.05$, Figure 4, C–F) in cultured endothelial cells but not in the presence of endothelial cells treated with VEGFR2 siRNA. Moreover, similar experiments demonstrated increases in endothelial VCAM, which was inhibited in the presence of VEGFR2 siRNA. Additionally, in ApoE-/-
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**A**

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**B**

VEGF-A concentration (pg/ml)

**C**

ApoE<sup>−/−</sup>  ApoE<sup>−/−</sup> CD163<sup>−/−</sup>

**D**

Plaque area

P < 0.01

**E**

Stenosis

P < 0.01

**F**

Lesion classification

P < 0.01

**G**

Necrosis

P < 0.05

**H**

BCA  CD163 DAPI  VE-cadherin DAPI  CD163 VE-cadherin DAPI

**I**

Microvessel density

P < 0.05

**J**

Intraplaque FITC-dextran fluorescence (AU)

P < 0.05
mouse, a VEGF-blocking antibody not only decreased intraplaque endothelial VCAM expression but also inhibited monocyte recruitment. Similarly, human plaque microvessels surrounded by CD163 cells showed abnormalities in VE-cadherin and extravasation of vWF, unlike those located away from CD163 cells. Deletion of CD163 in mice reduced plaque permeability, microvessel quantity, microvascular inflammation, inflammatory cell recruitment, and plaque progression. These data strongly suggest that macrophage-secreted VEGF-A is an important factor increasing endothelial permeability within human intraplaque vessels.

While the macrophages inflammatory response in areas of IPH is essential for controlling the toxic effect of free iron derived from erythrocytes, the resultant proangiogenic changes in macrophage function are almost certainly pathogenic within the context of atherosclerosis. Small amounts of intimal angiogenesis and bleeding might be made worse by the resultant inflammatory response. Clinical studies using serial carotid MRI to detect IPH in humans with carotid artery disease have shown that IPH has long-standing deleterious effects on plaque growth that remain detectable for years after the initial bleeding event (37). Although not specifically explored by us, it is likely that the macrophage response to hemorrhage shown here is similar to that for other diseases in which hemorrhage (i.e., tumors, diabetic retinopathy, and macular degeneration) plays an important role.

We previously hypothesized that antiangiogenic therapies using VEGF-A inhibitors might be important for the normalization of plaque vasculature and thus limit further risk of IPH and necrotic core expansion (7). By reducing vessel growth, permeability, and leukocyte recruitment, such therapies might be able to limit the initial effects of IPH. Our data suggest that in patients with IPH identified via MRI or perhaps by molecular imaging of M(Hb) macrophages, local anti-VEGF therapies might be useful in halting plaque progression. Given the known side effects of proteinuria, hypertension, and thromboembolism, the risks of systemic anti-VEGF therapies might not be justified (38).

We would be remiss if we did not briefly discuss the limitations of this study. Surgical CEA specimens involve a certain selection bias of surgical manipulation, in that these findings may not be representative of humans with IPH who do not undergo endarterectomy. The anti-CD163 antibodies used in this study could also potentially detect soluble CD163 in the specimens, but we believe this is unlikely, given that the specimens were perfusion fixed and that the vast majority of CD163 expression was associated with nucleated cells. Moreover, given the observational nature of human pathology studies, it remains uncertain as to what extent the intimal angiogenesis and vascular permeability seen in the samples examined can be attributed to M(Hb) cells as opposed to other causes. Last, we focused on local iron and Hb release at sites of IPH; however, there may be other important potential contributors to plaque progression not explored here, including lesional cytokine production by CD163 macrophages and local thrombin generation with consequent changes in cellular functions induced by protease-activated receptor action. Nonetheless, the in vitro demonstrations of these effects in cultured endothelial cells under normoxic conditions and in a CD163-deficient mouse model of IPH lend further support to our conclusions. In addition, while we identified a relationship between a human CD163 polymorphism (i.e. rs7136716) and its relationship to increased risk of MI and CHD in African Americans, this SNP has not been previously related to the risk of CHD in larger GWAS studies, which have mostly been conducted in individuals of European and Asian descent (39). This might be due to the low frequency of homozygous minor allele carriers in European populations (<1%) in contrast to its expression in other ethnic groups (20%–30%). Moreover, because the location of this variant is 30 kbp upstream of the CD163 promoter, its function in relationship to CD163 remains unknown and difficult to determine. Nonetheless, our data showing greater plaque angiogenesis and CD163 expression in subjects with 2 copies of the G-allele suggest a functional role for this SNP with regard to CD163.

In conclusion, our study demonstrates for the first time to our knowledge that a specific alternative macrophage subset driven by uptake of Hb promotes intimal angiogenesis, vascular permeability, and plaque inflammation, features that probably promote plaque progression. Our data also reveal the molecular mechanisms underlying these effects and pinpoint the HIF1a/VEGF-A system as a potential target for antiatherosclerotic therapies.

**Methods**

*Human carotid specimens.* Human carotid plaques removed from patients eligible for surgical CEA and pathological specimens of coronary artery samples were selected from the CVPath Registry (18). All plaques were identified according to a modified American Heart Association classification (40). Thirty-eight human carotid plaques removed from patients eligible for surgical CEA, in which the carotid plaque (common and internal carotid artery plaque) was removed in an undisturbed manner, were selected from the CVPath Registry of sixty cases. Indications for surgery were ipsilateral symptoms involving transient ischemic attack or stroke in the previous 6 months or signifi-
Figure 6. CD163+ macrophages increase intraplaque endothelial VCAM and leukocyte recruitment via VEGF. Activation of VCAM and NF-κB in HAECs (A–J) and association of high CD163+ macrophages and inflammation in human atherosclerotic lesions (K–N). In A–C, HAECs were incubated with culture media supernatants collected from human macrophages differentiated in HH [M(Hb) Sup] or control media [M(con) Sup] overnight. The expression of VCAM, p-p65 (p–NF-κB), and total NF-κB was measured by immunoblotting. LPS (100 ng/ml) was used as a positive control (n = 4 per group). (D) Immunostaining for VEGF in carotid (green) and VCAM (red) in BCA plaques from 1-year-old ApoE−/− and ApoE−/− CD163+/− mice. Scale bars: 100 μm. (E) The colocalization of VE-cadherin and VCAM in confocal microscopic images was analyzed for correlation coefficient (n = 6–7 per group). (F) Total macrophage content in BCA plaques as described in H was analyzed using macrophage marker Mac3 immunohistochemical staining (n = 5 per group). (G–I) Cells were transfected with VEGFR2 siRNA for 48 hours before incubation with culture media supernatants differentiated with HH or control media. The expression of VCAM, ICAM, p-p65 (p–NF-κB), and total NF-κB was measured by immunoblotting (n = 4 per group). Scr siRNA, scrambled siRNA. (J) Monocyte infiltration into BCA plaques from 6- to 8-month-old mice after a 4-week treatment with a control or VEGF-blocking antibody. Monocyte tracing was performed by injecting Au-nanoparticle–labeled monocytes, and labeled monocytes were counted on histological images (n = 6 per group). (K) Representative images of H&E and CD163 immunohistochemical staining showing high CD163 and low CD163 areas. Scale bars: 500 μm and 100 μm. (L) Areas of high and low expression of CD163+ macrophages in human atherosclerotic plaques were scored for inflammation (n = 7–8 per group). (M) CD163+ macrophages and CD3+ T cells were detected surrounding microvessels in high CD163 areas (white dashed lines show microvessels). Scale bars: 20 μm. (N) Quantitation of CD3+ T cells in high and low CD163+ macrophages surrounding intraplaque microvessels (n = 8 per group). Data represent the mean ± SEM (B, C, and H–J), the mean ± SD (E, F, J, and N), or the median for the box and whisker plot (L). (H–J) *P < 0.05, by 1-way ANOVA, and, if the variance ratio test (F test) was significant, a more detailed post-hoc analysis of differences between groups was performed using a Tukey-Kramer honest significant difference test. (B and C) *P < 0.05, by 2-sided Student’s t test. (E, F, and N) P < 0.01 and P 0.05, by 2-sided Student’s t test. (L) P < 0.01, by Mann-Whitney-Wilcoxon test.

cant carotid stenosis (>70%) measured by duplex ultrasound. In total, there were 18 symptomatic cases and 20 asymptomatic cases. At surgery, the carotid tissue was chilled immediately after excision and then promptly frozen at –80°C and shipped to the CVPath Institute. The CECA specimen was partially thawed and frozen in OCT (Miles Inc.). Cryosections were prepared using a Hacker Bright OTF/HS microtome cryostat equipped with a tungsten carbide knife with an installed CryoJane (Leica Microsystems). The CryoJane Tape-Transfer System improves cryosectioning outcomes on glass slides, particularly for the heavily calcified tissues common to carotid plaques (41). The frozen tissue was step sectioned at 10-μm thickness, 15 to 20 serial sections for staining were collected on glass slides, and adjacent frozen sections of 1.0 to 1.2 mm were collected in plastic tubes and frozen for protein extraction (see below). These steps were repeated until the frozen block was exhausted. This method allowed matching of histological findings with protein analysis. The entire carotid plaque removed at surgery was sampled in this manner.

Lesion classification and selection. Lesion morphology was assessed according to a simplified scheme as previously reported by our laboratory (40). Three sections from each carotid plaque were selected for morphometric analysis, the lesions with the plaque rupture site or ulceration or the most severely narrowed segment were used as the culprit site, and lesions flanking plaques) located 1.2 mm above or below the presenting lesion were also assessed by IHC. Thirty-eight specimens containing culprit segments along with proximal and distal flanking segments were available for analysis and assessed for plaque morphology. A total of 114 plaques (38 culprit, 38 proximal flanking, and 38 distal flanking plaques) were classified, of which 38 were fibroatheromas, 20 were healed plaque ruptures, 14 were fibrocalcific lesions, 11 were plaque ruptures, 9 were TCFAs, 7 were calcified nodules, 5 were PITS, 5 were intimal xanthomas, 2 were chronic total occlusions, 2 were fissures, and 1 was a fibrous plaque.

Inflammation scoring. Pathological scoring of plaque inflammation was performed to assess its severity and distribution. Plaque inflammation was regionally assessed using a scoring system accounting for the severity according to the number of inflammatory cells and the relative distribution based on quadrants. Severity was scored on a scale of 0 to 4: 0–20 inflammatory cells = 0; 21–100 cells = 1; 101–200 cells = 2; 201–400 = 3; and more than 400 cells = 4. Distribution was also scored on a scale of 0 to 4: 0–45 degrees = 0; 45–90 degrees = 1; 90–180 degrees = 2; 180–270 degrees = 3; and 270–360 degrees = 4. The overall inflammation score was drawn from the scoring matrix (Supplemental Table 4). Inflammatory cells (macrophages and T cells) were identified on H&E staining on the basis of their characteristic appearance. Scoring was confirmed by a pathologist (R. Virmani).

Immunohistochemical and immunofluorescence staining. Cryosections were stained using both H&E and Movat pentachrome stains. Areas of foam cell macrophages and M(Hb) macrophage were identified with CD68 and CD163 immunohistochemical staining, respectively, as previously reported (6). The ratio of CD163 to CD68 staining was analyzed, and ratios below 0.20 were considered low CD163 plaque areas, while those above 0.70 were considered high CD163 plaque areas. Ten plaque samples contained high CD163/CD68 areas, while ten plaques contained low CD163/CD68 areas and were collectively chosen for further analysis. Five to six samples each from the plaques with low and high CD163 areas were selected for immunoblotting to detect CD163, HIF1α, and VEGF-A expression in the plaque. Negative control plaques that were low in both CD68 and CD163 showed no HIF1α and VEGF-A expression by immunoblotting (data not shown). Ten plaques were selected from the low and high CD163 area samples for quantitation of angiogenesis. Microvessel numbers were assessed following immunohistochemical staining for anti-CD31 and anti-vWF.

Immunofluorescence staining for CD163/vWF, CD68/vWF, CD163/HIF1α, CD163/VEGFA, CD163/VE-cadherin, VE-cadherin/VCAM, and CD163/Cd3 was carried out on cryosections or formalin-fixed, paraffin-embedded sections. The frozen sections were fixed in cold acetone for 10 minutes, with further exposure to 0.15% H2O2 for 20 minutes. The sections were treated with Dako protein block (catalog X0909) for 10 minutes before incubation with primary antibodies against CD163 (Santa Cruz Biotechnology, catalog sc-20066, clone GH/61, dilution 1:200, overnight at 4°C); CD68 (Dako, clone KP1, dilution 1:800, 1 hour); vWF (Strategic BioSolutions, S4003GND1, dilution 1:2,000, 1 hour); HIF1α (Novus Biologicals, NB100-105, dilution 1:40, 1 hour); VEGF-A (BioGenex, PU483-UP, dilution 1:40, overnight); VE-cadherin (R&D Systems, AF1002, dilution 1:100, and BD Biosciences, 555661, dilution 1:400, overnight); CD3 (Roche, 790-4341, prediluted); and VCAM (Abcam, ab134047, dilution 1:100). Antibody detection included biotinylated goat anti-rabbit, horse anti-mouse, and rabbit anti-goat (Vector Laboratories, Reveal ABC kit).

RESEARCH ARTICLE

The Journal of Clinical Investigation

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The Journal of Clinical Investigation
Human coronary artery microvessel permeability was assessed by EBD perfusion. (A) Representative images of EBD-perfused human coronary arteries, H&E-stained images, and confocal immunofluorescence images of CD163 (red) and VE-cadherin (yellow) or VE-cadherin (red) and VCAM (green) in an EBD-negative area (top row), EBD-positive area 1 (middle row), and EBD-positive area 2 (bottom row). Positive areas 1 and 2 are shown in progressively higher-magnification H&E-stained images from left to right in the second and third rows. Red and white arrowheads point to microvessels. Confocal images of the EBD-negative areas for CD163/VE-cadherin and VE-cadherin/VCAM are shown in the top row of columns 3 and 4, respectively, while the positive area 1 is shown for CD163/VE-cadherin in the middle rows of columns 3 and 4 (higher-magnification image on the right), and positive area 2 is shown for VE-cadherin/VCAM in the bottom row of columns 3 and 4 (higher-magnification image on the right). (B–D) Quantification of microvessel density, CD163+ macrophages, and VCAM in an EBD-positive area versus an EBD-negative area (n = 6–8 per group). (E) The SNP rs7136716 was associated with human coronary artery atherosclerotic plaque rupture and risk of coronary artery disease. See Table 2 for SNP analysis of the CVPath cohort. Representative images showed ruptured plaques in human coronary arteries from subjects with the rs7136717 AA genotype versus those with the GG genotype. Scale bars: 500 μm. Arrowheads point to microvessels. (F) Microvessel density per plaque area at the ruptured coronary artery plaque site from individuals with 0 copies (AA genotype, n = 22) versus 2 copies of the minor allele (GG genotype, n = 25) who died of plaque rupture. (G) Relative CD163 mRNA expression in ruptured coronary artery plaques from AA versus GG genotype groups as measured by qPCR (n = 13 per group). (H) Cox proportional hazards ratio assessment for the association of the genetic variant rs7136717 with incident MI and incident CHD in the ARIC cohort (n = 3,225). Data represent the mean ± SD (B–D, F, and G) or ORs and 95% CI (H). P < 0.01 and P < 0.05, by 2-sided Student’s t test (B–D, F, and G). In the ARIC cohort, Cox proportional hazards models were used to examine the association of the genetic variant with incident MI and incident CHD, and the analyses were adjusted for age, sex, ancestry-informative principal components, and study center (H).

Human coronary artery samples and EBD permeability assay. Human coronary artery samples were selected from freshly collected autopsy specimens from the CVPath Registry, with a post-modern interval (PMI) cut-off less than 12 hours and evidence of advanced atherosclerosis based on x-ray images of the heart. EBD (0.5%) solution was prepared and filtered in saline solution containing 5% BSA and then perfused into both the right and left coronary arteries through a mean pressure of approximately 60 mmHg for 15 minutes at 37°C. After dye perfusion, samples were further washed with PBS perfusion and fixed by neutral buffered formalin perfusion. Fixed tissue specimens were harvested, cut at 2- to 3-mm intervals, and embedded in OCT for frozen sectioning, staining, and histology as well as immunofluorescence analysis. A description of the demographic characteristics of the subjects whose hearts were studied in this experiment is provided in Supplemental Table 5.

RNAseq ISH and laser capture microdissection. In situ detection of CD163 and VEGF transcripts in human atherosclerotic plaques was performed using an RNAseq assay with RNAscope Probe-Hs-CD163-C2 (Advanced Cell Diagnostics, catalog 417061-C2), an RNAscope Probe-Hs-VEGFA (Advanced Cell Diagnostics, catalog 423161), and an RNAscope Fluorescent Multiplex Reagent Kit (Advanced Cell Diagnostics, catalog 320850) following the manufacturer’s protocols. Both positive (POLR2A and PPIB) and negative controls were included in the procedure according to the manufacturer’s instruction. All slides were digitally scanned with an LSM 800 laser-scanning confocal microscope (Carl Zeiss) using a ×60 objective with Airyscan (Carl Zeiss). Laser capture microdissection was performed as previously described (6).

Human coronary plaque and CD163 SNPs. SNPs from the CD163 gene were examined in 346 subjects (51 with plaque rupture and 295 with nonruptured plaques) from the CVPath Registry on an Illumina Exome BeadChip array. For further analysis of rs7136716, there were a total 111 subjects with homoygous major AA alleles, 175 with heterozygous AG alleles, and 60 with homoygous minor GG alleles. Evidence of plaque rupture was detected by serial sectioning of culprit plaques as identified by H&E and Movat pentachrome staining. Among the subjects with plaque ruptures, 15 had homoygous major AA alleles, 22 had heterozygous AG alleles, and 14 had homoygous minor GG alleles. The ruptured culprit lesions in coronary arteries were analyzed for intraplaque microvessel density in age- and sex-matched subjects with homoygous major alleles in subjects with homoygous minor alleles. RNA was extracted from ruptured artery specimens, and CD163 expression levels were measured by quantitative PCR (qPCR) using SYBR Green Master Mix (Life Technologies, Thermo Fisher Scientific).

CHD risk of SNPs in the human cohort. The Atherosclerosis Risk in Communities (ARIC) study is a prospective, population-based cohort study designed to investigate the etiology and predictors of cardiovascular disease and has been described in detail previously (42). Briefly, participants aged 45 to 64 years at baseline were recruited from 4 communities: Forsyth County, North Carolina; Jackson, Mississippi; Minneapolis, Minnesota; and Washington County, Maryland. A total of 15,792 individuals, predominantly of European and African American ancestry, participated in the baseline examination between 1987 and 1989, with 3 additional triennial follow-up examinations and a fifth examination between 2011 and 2013. In the present study, we included only participants of African American ancestry. Human Exome BeadChip, version 1.0 (Illumina), was used to genotype 247,870 exonic variants across the genome, which were called within the context of the Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) consortium as described previously (43, 44). Only genotypes for the genetic variant rs7136716 were used in the present study. The genotype was coded using the additive model. We studied the genetic variant in relation to 2 cardiovascular disease endpoints: (a) MI: fatal or nonfatal MI, and (b) CHD: fatal or nonfatal MI, fatal CHD, sudden death within 1 hour of onset of symptoms, or revascularization (percutaneous coronary artery intervention such as stenting or balloon angioplasty or coronary artery bypass grafting).
Statistics. Analysis of normality was performed using a Shapiro-Wilk test. Normally and non-normally distributed data were analyzed for significance by Student’s t test or Mann-Whitney-Wilcoxon tests, respectively. For multiple group comparisons, 1-way ANOVA was applied. If the variance ratio test (F test) was significant, a more detailed post-hoc analysis of differences between groups was made using a Tukey-Kramer honest significant difference test. A P value of less than 0.05 was considered statistically significant. See Supplemental Methods for more detailed information.

Author contributions
LG, HA, EH, SLS, FO, RLG, REB, MHD, HJ, RK, HM, QC, FDK, RV, and AVF contributed to the analysis of human atherosclerotic specimens. VK, RP, and HA performed in vitro experiments involving human and mouse macrophages. LG, EH, SLS, REB, CUC, ALJ, and MJL performed mouse experiments and analysis. PSDV, MLG, DEA, EB, ACM, JE, and NS performed SNP analysis. JK, PC, DPC, YZ, and YC provided experimental materials. LG, HA, AG, MAS, RV, and AVF drafted and/or edited the manuscript, which was critically reviewed by all the authors.

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The Journal of Clinical Investigation

RESEARCH ARTICLE

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Address correspondence to: Aloe V. Finn, CVPath Institute, 19 Firstfield Road, Gaithersburg, Maryland, 20878, University of Maryland School of Medicine, Baltimore, Maryland, 21201, USA. Phone: 301.208.3570; Email: afinn@cvpath.org.

Hirokuni Akahori’s present address is: Department of Internal Medicine, Cardiovascular Division, Hyogo College of Medicine, Nishinomiya, Hyogo, Japan.

Emanuel Harari’s present address is: Division of Cardiology, Rabin Medical Center, Petah-Tikva, and Sackler Faculty of Medicine, Tel-Aviv University, Tel Aviv, Israel.

Fumiyuki Otsuka’s present address is: National Cerebral and Cardiovascular Center, Suita, Osaka, Japan.

Cheol Ung Choi’s present address is: Cardiovascular Center, Division of Cardiology, Departments of Internal Medicine, Korea University Guro Hospital, Guro-gu, Seoul, Korea.

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