Homocysteine-mediated thrombosis and angiostasis in vascular pathobiology

Joseph Loscalzo

Department of Medicine, Brigham and Women’s Hospital, Harvard Medical School, Boston, Massachusetts, USA.

The mechanisms by which homocysteine contributes to atherothrombosis are complex and their in vivo relevance uncertain. In this issue of the JCI, Jacovina and colleagues report a unique in vivo mechanism by which homocysteine may contribute to vascular disease (see the related article beginning on page 3384). This group had previously reported that homocysteine impairs endothelial cell surface plasminogen activation by posttranslationally modifying annexin A2, the coreceptor for plasminogen and tissue plasminogen activator. They now show that an annexin A2–deficient mouse rendered hyperhomocysteinemiac by dietary means has impaired fibrinolysis, perivascular fibrin persistence, and attenuated angiogenesis (angiostasis). Potential mechanisms by which homocysteine-dependent changes in endothelial phenotype link thrombosis to angiostasis are reviewed and their relationship to homocysteine-dependent vascular disease considered.

Over the past 30 years, evidence from many in vitro and in vivo studies supports an association between mild to moderate hyperhomocysteinemia and vascular dysfunction and disease (1). A key mechanism that predisposes to vascular disease in hyperhomocysteinemia is endothelial dysfunction (2). Homocysteine induces endothelial dysfunction in part by promoting oxidant stress, as illustrated in cellular studies (3, 4) and in genetic animal models (2, 5, 6). The mechanisms for homocysteine-induced oxidant stress are complex and include inhibiting the translation of glutathione peroxidase-1 (7), a major antioxidant enzyme in vascular cells that regulates mitochondrial reactive oxygen species flux (8) and whose overexpression rescues the normal vascular phenotype of hyperhomocysteinemic mice (9); upregulation of NADPH oxidase expression (10); enhanced expression of inducible nitric oxide synthase (10) and uncoupling of nitric oxide synthases (11); and decreased glutathione levels owing to decreased cysteine synthesis (12), thereby shifting the redox balance of endothelial cells toward oxidant stress.

Among the specific phenotypic properties of the endothelial cell that go awry in hyperhomocysteinemia are its antithrombotic properties. Hyperhomocysteinemia produces a prothrombotic state (13), some mechanisms for which include enhanced platelet activation, enhanced coagulation (likely as a consequence of increased tissue factor expression; ref. 14), and attenuated fibrinolysis. Two recognized mechanisms for reduced fibrinolysis include posttranslational modification of fibrinogen by homocysteinylation, rendering the fibrin derived from fibrinogen relatively resistant to plasmin (15), and increased activity of thrombin-activatable fibrinolysis inhibitor (TAFI) (16).

In this issue of the JCI, Jacovina and colleagues (17) report the existence of another antifibrinolytic mechanism caused by homocysteine. In the group’s earlier work (18, 19), they demonstrated that the calcium-regulated, phospholipid-binding protein annexin A2 is an endothelial coreceptor for TPA and plasminogen that facilitates plasminogen activation by enhancing the catalytic efficiency of TPA. They next demonstrated that homocysteine forms a mixed disulfide bond with the Cys9 residue of annexin A2, which decreases the binding affinity of the receptor for TPA. As a result of this posttranslational modification, endothelial cell surface plasminogen activation is markedly attenuated (18). These observations were next confirmed in vivo using Anxa2−/− mice, which showed increased perivascular fibrin deposition and impaired fibrin clearance. Fibrin formation and plasmin-dependent remodeling are not only important in the hemostatic response to vascular injury, but also for luminal recanalization and angiogenesis during the healing phase. Consistent with this role for fibrin and plasmin, the Anxa2−/− mice showed significantly impaired angiogenesis in a range of vascular injury models (neoangiogenesis) (19). In their latest study, reported in this issue of the JCI (17), the investigators found that moderate hyperhomocysteinemia induced by feeding mice methionine also leads to impaired fibrinolysis with perivascular fibrin persistence as well as attenuated angiogenesis (angiostasis). In this animal model, they observed significantly reduced TPA–annexin A2 binding and plasminogen activation consistent with the importance of endothelial cell surface plasmin activity for normal fibrin clearance and angiogenesis. The pathophenotype was readily reversed by infusion of wild-type recombinant annexin A2 into the hyperhomocysteinemic mice (17).

Homocysteine and angiostasis

These interesting data (17) support yet another novel mechanism by which hyperhomocysteinemia promotes the complications of atherothrombotic vascular disease, that is, limiting the angiogenic response to ischemia. Homocysteine-dependent changes in the vascular phenotype previously shown to contribute to impaired angiogenesis include a decrease in glutathione peroxidase expression and consequent increase in oxidant stress, leading to endothelial progenitor cell dysfunction (7, 20); decreased bioactive nitric oxide generation (2, 3); and dysregulation of MMP activity (21). Specifically, homocysteine has been shown to increase the expression and activity of MMP-2 and MMP-9 (21). In light of the results of the study by Jacovina and colleagues (17), the increase in MMP activity suggests that zymogen activation does not alone depend on cell surface-bound

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plasmin; rather, as recently shown in vitro, both Factor Xa and thrombin can also activate MMP-2 (22), thereby providing a potential mechanism by which to link the prothrombotic effects of homocysteine to matrix degradation. As MMP-dependent matrix degradation is essential for (endothelial) cell migration, the association of increased MMP activity with impaired angiogenesis may at first appear paradoxical. However, two angiogenesis inhibitors, endostatin and angiostatin, can be generated by the action of MMPs on collagen 18 and plasminogen, respectively (23, 24). Taken together, these data suggest that homocysteine-mediated, tissue factor–dependent thrombin generation not only promotes the prothrombotic state of hyperhomocysteinemia, but also contributes indirectly to angiostasis following ischemic injury.

Homocysteine and the prothrombotic state

Other evidence supports an indirect angiostatic role for thrombin. A thrombin-mediated increase in TAFI (16) not only inhibits fibrinolysis, but also inhibits endothelial tube formation and angiogenesis (25). Inhibition of fibrinolysis leads to attenuated release of proteolytic degradation fragments of fibrin that have angiogenic potential (26), such as fibrin fragment E (27). These indirect angiostatic effects of thrombin may be offset by its activation of the transglutaminase factor XIII; factor XIIIa not only crosslinks fibrin, but also crosslinks the integrin α5β3 with VEGFR2 on the endothelial cell to promote angiogenesis and suppress expression of the natural angiostatic adhesive glycoprotein thrombospondin-1 (28). While these indirect factor XIII–mediated angiogenic effects of thrombin may be active in a hyperhomocysteinemic ischemic model, they are clearly inadequate to overcome the angiostatic effects of homocysteine, leading to impaired angiogenesis in the model.

Lastly, as yet another potential level of regulation of angiogenesis by the homocysteine-dependent prothrombotic state, consider the consequences of elevated homocysteine concentrations on the methylation potential of vascular cells. In general, elevated homocysteine leads to a decrease in the S-adenosyl-methionine/S-adenosyl-homocysteine ratio, which decreases methylation potential and suppresses global DNA methylation (29). As a result, homocysteine can increase the expression of genes that inhibit fibrinolysis, such as plasminogen activator inhibitor-1 (30), and that inhibit angiogenesis, such as thrombospondin-1 (31).

Conclusions

Thus, the current study by Jacobina and colleagues (17) adds an important new feature to the adverse vascular phenotype of hyperhomocysteinemia, namely, impaired angiogenesis or angiostasis. Furthermore, this study links angiostasis to altered endothelial cell surface plasminogen activation, supporting the importance of plasmin in the angiogenic response. In addition and importantly, their results give one pause to consider the pathobiological network that defines the interface between thrombosis and angiogenesis (see Figure 1). Taken together, these findings highlight the complexity of molecular events that conspire to produce a prothrombotic and angiostatic endothelial pathophenotype from homocysteine exposure and offer novel explanations for the epidemiological association between hyperhomocysteinemia and vascular disease.

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Address all correspondence to: Joseph Loscalzo, Department of Medicine, Brigham
Mutations in the enzyme superoxide dismutase 1 (SOD1) have been linked to the neurodegenerative disease amyotrophic lateral sclerosis (ALS). In this issue of the JCI, Zhong et al. report that the endogenous anticoagulant activated protein C (APC) is able to cross the blood–spinal cord barrier in mice and signal to both neuronal and non-neuronal cells (see the related article beginning on page 3437). This signaling resulted in the suppression of mutant SOD1 synthesis and retarded disease progression in a murine model of ALS. Here we discuss the potential importance of these data and possible relevance to human neurodegenerative diseases.

Amyotrophic lateral sclerosis (ALS) is a devastating neurodegenerative disease that strikes in midlife, causing progressive weakness, disability, and death. Unfortunately, the cause of ALS is unknown, and the treatment is largely palliative. Research into the pathogenesis of and potential treatments for ALS focuses heavily on experimental models that use human genetic mutations in transgenic animals or on cellular models that express mutant proteins known to impart a high risk of developing ALS in people who carry these mutations. Mutant superoxide dismutase 1 (SOD1) is the most common protein known to cause ALS in humans; however, the mechanisms underlying mutant SOD1–related ALS are unknown. People with mutant SOD1–related ALS represent only about 20% of inherited (i.e., familial) ALS cases and about 2% of all patients with ALS (1). Nevertheless, the animal models of mutant SOD1–related ALS develop a neurological disorder that mimics the human disease, and investigations using these models have taught us a lot about motor neuron biology as well as the potential interactions between motor

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