Protective effect of marrow microenvironment. The Hematopoietic Inductive Microenvironment (HIM) niches include osteoblasts, stromal/mesenchymal cells, endothelial cells and extracellular matrix components. AML blast quiescence, proliferation and apoptosis are influenced by receptor kinases, adhesive receptors and signaling via matrix-mediated or bound chemokines and cytokines. AMD3100 and AMD3465 CXCR4 antagonists +/- sorafenib or inhibitors of VLA-4/CAM-1 interactions chemosensitize AML blasts within the HIM niches. Shh: sonic hedgehog, Notch, vascular endothelial factor VEGF, and interleukin VLA-4, CXCR4, sonic Hedgehog (shh), vascular endothelial receptors signals promote leukemic stem cell survival and expansion and can be targeted to overcome AML chemoresistance.

ERK survival pathways. CXCR4 inhibition partially abrogated the protection conferred by stromal cells, rendering these leukemic cells more susceptible to apoptosis when exposed to cytarabine. AMD3465 led to down-regulation of FLT3 receptor expression and inhibition of KIT signaling when used in conjunction with sorafenib, a permissive tyrosine kinase inhibitor in FLT3-ITD-expressing AML cell lines (see figure). The latter observations are very intriguing and may explain in part the successful treatment of a relapsed FLT3-ITD-positive AML patient with sorafenib in the posttransplantation setting.

Finally, while both reports open new avenues for overcoming in vivo drug resistance in AML, it is yet unclear whether durable complete remissions can ensue from this strategy. AML is indeed a very heterogenous disease, and successful eradication of leukemic stem/progenitor cells will require blocking multiple receptors/pathways as shown in the figure, with targeted agents focused on CD44, VLA-4, CXCR4, sonic Hedgehog (shh), vascular endothelial factors VEGF, and interleukin 3 receptor alpha chain CD123, in addition to many factors likely not yet discovered.

Progress in whole genome sequencing approaches may uncover novel target genes and signaling pathways that may radically alter our approach to therapy.

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REFERENCES

In this issue, Li and colleagues describe a new mechanism promoting thrombus dissolution involving thrombin-induced secretion and cleavage of an “orphan” metalloproteinase ADAMTS-18. This mechanism involves a dynamic process that is essential for the cessation of bleeding at sites of vascular injury. Disruption of this process can lead to excessive thrombus formation, particularly at sites of atherosclerotic plaque rupture, precipitating...
the acute coronary syndromes or ischemic stroke. The hemostatic response is regulated by competing prothrombotic and antithrombotic forces. A key player in this regard is the serine protease, thrombin. In its free form, thrombin is procoagulant, activating several components of the coagulation cascade as well as converting fibrinogen to fibrin. It is also one of the most potent platelet agonists known. Conversely, thrombin bound to thrombomodulin on vascular endothelial cells can activate protein C, a potent physiologic anticoagulant.

In this issue of Blood, Li et al describe a surprising new role for thrombin: promoting the dissolution of platelet thrombi. They demonstrate that thrombin, generated at the site of vessel injury, is able to cleave a protease of previously unknown function, ADAMTS-18, which, following secretion from activated endothelium, binds to the \( \beta_3 \) subunit of the platelet adhesion receptor integrin \( \alpha_{IIb}\beta_3 \). The thrombin-generated 45-kDa C-terminal cleavage product of ADAMTS-18 clusters the \( \beta_3 \) integrins and induces oxidative fragmentation of the platelet, leading to thrombus dissolution. This process is dependent on the sequential activation of platelet 12-lipoxygenase and NADPH oxidase, resulting in the intracellular production of reactive oxygen species (ROS) including hydrogen peroxide (H\(_2\)O\(_2\)).

Like many breakthroughs, the path leading to this discovery has been an unexpected one. It began with the seemingly unrelated observation that patients with HIV-1 infection develop autoimmune thrombocytopenia due to the presence of an antiplatelet antibody directed against the epitope GPIIIa49–66 of \( \alpha_{IIb}\beta_3 \) (GPIIb-IIIa). In general, the higher the titer of antibody, the lower the platelet count. It was determined that this antibody induced thrombocytopenia by inducing platelet fragmentation \( \textit{via} \) a process dependent on the generation of H\(_2\)O\(_2\) and other ROS. The search for a physiologic ligand that could induce oxidative platelet fragmentation was undertaken using the platelet GPIIIa49–66 peptide as bait in a phage surface display library. From 20 clones isolated, 1 had 70% identity with a C-terminal region of ADAMTS-18. Li et al demonstrate that an active C-terminal fragment of ADAMTS-18 can completely dissolve platelet aggregates formed in vitro. Moreover, this fragment lyse thrombus formed in the carotid artery of mice and reduces infarction and neurologic impairment in a cerebral stroke model. Importantly, this protection was afforded whether the fragment was infused 2 hours before or 2 hours after the 90-minute ischemia and reperfusion, with no adverse bleeding events.

These provocative findings shed new light on the physiologic processes regulating thrombus dissolution in vivo. It has long been recognized that platelet thrombus formation is a dynamic process, dependent on the equilibrium between platelet deposition and dispersal. As a consequence, impairment of thrombin/ADAMTS-18–dependent platelet dissolution could conceivably produce a prothrombotic phenotype. It is also possible that thrombin/ADAMTS-18 could act as a safeguard in situations where the anticoagulant function of thrombin is compromised, as occurs in proinflammatory states.

As with any good study, this work poses more questions than it answers. Foremost is whether thrombin-induced cleavage of ADAMTS–18 represents the dominant physiologic mechanism controlling thrombus dissolution in vivo. In this regard, the generation of ADAMTS–18 null mice or mice expressing noncleavable ADAMTS–18 will be informative. This issue aside, the authors clearly demonstrate that the C-terminal fragment of ADAMTS–18 is highly effective at promoting platelet thrombus dissolution in vivo. This may have direct clinical relevance, as platelet-rich thrombi are notoriously difficult to lyse with fibri ninolytic therapies alone. Hence, therapeutic approaches that induce platelet fragmentation may represent an innovative approach to enhance fibrinolysis. At a basic level, the study by Li et al raises interesting issues with respect to the physiologic mechanisms regulating H\(_2\)O\(_2\) and ROS production during thrombus development. These molecules are known to be generated following platelet activation, which begs the question as to what stage in the thrombotic process these molecules are generated and how their platelet-stimulating functions are coordinated with their ability to induce platelet fragmentation. Additionally, whether thrombin is the only (or major) proteolytic enzyme with the capacity to regulate the activity of ADAMTS–18 remains to be determined. Nonetheless, despite being one of the most intensely investigated proteases, new and unanticipated functions for thrombin continue to emerge, reinforcing its role as a master of thromboregulation.

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Is thrombin the problem or (dis)solution?

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