HIV-associated hematological abnormalities involve all lineages of blood cells, thus implying that the virus impairs the function of early HSCs. However, the underlying mechanisms of this defect are unknown, particularly since HSCs are largely resistant to HIV-1 infection. In this issue of the *JCI*, Prost and colleagues show that the viral accessory protein Negative factor (Nef) plays a potentially critical role in the pathogenesis of HIV/SIV-associated hematopoietic dysfunction by affecting the clonogenic potential of HSCs (see the related article beginning on page 1765). Soluble Nef induces PPARγ in uninfected HSCs, thereby suppressing the expression of STAT5A and STAT5B, two factors necessary for proper HSC function. The identification of this novel activity of extracellular Nef defines a new mechanism of HIV/SIV pathogenesis and suggests that approaches aimed at increasing STAT5A and STAT5B expression may be considered in HIV-infected individuals with prominent hematological abnormalities. The results also raise the question of whether dysregulation of hematopoiesis by extracellular Nef plays a role in the development of T cell immunodeficiency and the high levels of chronic immune activation associated with AIDS.

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this process and may ultimately facilitate the prevention and treatment of kidney disease.

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Is Nef the elusive cause of HIV-associated hematopoietic dysfunction?

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HIV-associated hematological abnormalities involve all lineages of blood cells, thus implying that the virus impairs the function of early HSCs. However, the underlying mechanisms of this defect are unknown, particularly since HSCs are largely resistant to HIV-1 infection. In this issue of the *JCI*, Prost and colleagues show that the viral accessory protein Nef plays a critical role in the pathogenesis of HIV/SIV-associated hematopoietic dysfunction by affecting the clonogenic potential of HSCs (see the related article beginning on page 1765). Soluble Nef induces PPARγ in uninfected HSCs, thereby suppressing the expression of STAT5A and STAT5B, two factors necessary for proper HSC function. The identification of this novel activity of extracellular Nef defines a new mechanism of HIV/SIV pathogenesis and suggests that approaches aimed at increasing STAT5A and STAT5B expression may be considered in HIV-infected individuals with prominent hematological abnormalities. The results also raise the question of whether dysregulation of hematopoiesis by extracellular Nef plays a role in the development of T cell immunodeficiency and the high levels of chronic immune activation associated with AIDS.

Nonstandard abbreviations used: Nef, Negative factor.

Conflict of interest: The authors have declared that no conflict of interest exists.

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several studies indicate that they are not susceptible to HIV-1 infection (7–9). Alternative mechanisms that were proposed — although without definitive experimental evidence — include an effect of HIV-1 proteins on uninfection HSCs as well as the hematosuppressive potential of certain proinflammatory cytokines that are produced at high levels during HIV infection (10–13).

### Impaired STAT5 and HSC function in SIV-infected macaques

The study from Prost and colleagues in this issue of the *JCI* (14) describes a series of experiments aimed at elucidating the mechanisms causing the hematopoietic abnormalities associated with HIV infection and AIDS. The main conclusion is that the extracellular viral accessory protein Nega 

Figure 1

Pathogenesis of the hematological abnormalities associated with HIV infection. (A) Schematic demonstrating how bone marrow–derived HSCs develop into precursors of the three main blood cell lineages (erythroid cells, myeloid cells, and platelets) as well as into the T cell precursors that will mature in the thymus and create the pool of circulating naive and memory CD4+ T cells. Ag, antigen. (B) Schematic demonstrating how the soluble viral accessory protein Nef may affect the ability of HSCs to maintain normal levels of peripheral blood cells, based on the new findings of Prost and coworkers reported in this issue of the *JCI* (14). In addition, Nef may decrease the production of T cell precursors, thus contributing to the peripheral CD4+ T cell depletion observed in HIV-infected individuals that is caused by both the direct effects of HIV on infected cells and the indirect effects of chronic immune activation.

A role for Nef in impaired hematopoiesis

Surprisingly, the Prost study (14) shows that extracellular, soluble forms of the HIV and SIVmac Nef proteins may downmodulate STAT5 expression by activating the transcriptional suppressor PPARγ, thus providing a new mechanism to explain the HIV-associated HSC defects (Figure 2B). Of note, this inhibitory effect of soluble Nef on the clonogenic potential of progenitor cells (which the authors elegantly map to a central region of HIV-1 Nef between amino acids 66 and 97) might explain why the hematological abnormalities of HIV-infected patients are correlated with disease progression even though HSCs are not directly infected by the virus. Nef is an early accessory protein of immuno-deficiency viruses that is required for efficient viral persistence and strongly accelerates disease progression in HIV-1–infected humans and in experimentally SIV-infected rhesus macaques (18). Nef performs a striking variety of activities that allow the virus to spread effi-
ciently in the infected host and to evade the immune system, including downmodulation of CD4, MHC class I, and in many cases CD3 (most SIV strains and HIV-2, but not HIV-1 and its simian precursors), upregulation of the invariant chain associated with immature MHC class II complexes, and enhancement of viral infectivity and replication (18, 19). Furthermore, Nef interacts with various kinases, modulates cell signaling pathways, and might alter cytokine secretion (20). All these activities of Nef require an intact N-terminal myristoylation signal and are exerted within the virally infected cell. However, Nef is also found in the extracellular environment and reaches concentrations of up to 10 ng/ml in sera of HIV-1–infected individuals (21). Reported effects of extracellular Nef include activation of STAT1 and STAT3 in macrophages (22), induction of apoptosis (23), and immunoglobulin class switching in bystander B cells (24). However, the true in vivo significance of these findings is difficult to assess as the above mentioned results were often obtained using rather artificial experimental conditions characterized by exceptionally high levels of Nef usually produced by bacteria. In this regard, Prost and coworkers (14) demonstrate that sera from SIVmac251-infected macaques can directly suppress STAT5A/B expression and cause HSC dysfunction (and that these effects are reduced by anti-Nef antibodies). These latter results not only support the possibility that the Nef-dependent STAT5A/B suppression occurs at “physiologic” concentrations of this protein, but also raise the question of whether humoral immune responses to Nef, which are detectable in the majority of HIV-1–infected individuals (25), may prevent the development of hematopoietic dysfunction.

**Nef-mediated suppression of hematopoiesis: viral immune evasion strategy or suboptimal virus-host adaptation?**

As mentioned above, the current study (14) might explain why the severity of the hematologic abnormalities correlates with the levels of virus replication and why functional hematopoiesis is restored by HAART. The observation that extracellular Nef impairs the function of CD34+ progenitor cells in trans also clarifies why HIV-1 does not need to directly infect HSCs to cause hematological abnormalities. However, new interesting questions are raised by the observed results. For example, Prost et al. show that peptides corresponding to residues 66–97 of the central core region of HIV-1 Nef are sufficient to reproduce the defect of HSC function. Although this region is relatively well conserved, the HIV-1 and SIVmac251 Nef sequences differ in 9 of the 31 amino acid residues. Those of other SIVs are frequently even more divergent and — in contrast to HIV-1 Nef proteins — often do not contain an intact Src homology 3–binding (SH3-binding) domain in this region, raising the question of whether this Nef activity is conserved among the different groups of primate lentiviruses. This question is crucial as SIV infection of natural hosts such as the Sooty mangabeys or African green monkeys is not associated with immunodeficiency or impaired bone marrow function despite chronically high levels of virus replication (26). From this perspective, it will be interesting to elucidate whether this effect of Nef is mainly an indicator of suboptimal virus-host adaptation in the evolutionarily recent human and experimental rhesus macaque hosts or indeed a conserved strategy of primate lentiviruses to evade the host immune system by suppressing the clonogenic potential of HSCs.

**Perspectives**

While the current study by Prost and colleagues (14) is of great interest, much work will be required to define the mechanism by which extracellular Nef induces PPARγ and to determine whether the levels of Nef in blood and bone marrow are sufficient to impair HSC function during HIV infection. It will also be important to better understand (a) to what extent this impaired hematopoiesis contributes to the T cell immunodeficiency associated with progression to AIDS, and (b) how these hematological defects relate to the state of chronic hyperimmune activation that seems to drive the loss of CD4+ T cells and progression to immunodeficiency in HIV-infected individuals. As STAT5 signaling plays a critical role in T cell development and function (27), an additional question is whether Nef also affects T cell homeostasis by suppressing STAT5A/B function in other, more mature T cell subsets (28). From a clinical point of view, the results of Prost and colleagues may potentially identify Nef and/or PPARγ as targets for new therapeutic approaches aimed at preventing or correcting the hematologic abnormalities associated with HIV-1 infection. On the other hand, the identification of PPARγ as a suppressor of HSC function might have implications for the use of PPARγ agonists, such as glitazones, in the treatment of diabetes.
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Immunosuppression in islet transplantation

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Islet transplantation can temporarily cure type 1 diabetes mellitus (T1DM) but requires simultaneous immunosuppression to avoid allograft rejection. In this issue of the JCI, Monti et al. report that immune conditioning via use of the Edmonton protocol—a treatment approach in which T1DM patients infused with pancreatic islets from multiple cadaveric donors simultaneously receive immunosuppressive drugs—results in lymphopenia that is associ- ated with elevated serum levels of the homeostatic cytokines IL-7 and IL-15, which causes in vivo expansion of the autoreactive CD8+ T cell population (see the related article beginning on page 1806). Reemergence of autoreactiv- ity is likely the main culprit underlying long-term islet graft failure, and new strategies will need to be tested to circumvent this homeostatic expansion and recurrent autoreactivity.

Nonstandard abbreviations used: GAD, glutamic acid decarboxylase; MMF, mycophenolate mofetil; TIDM, type 1 diabetes mellitus.

Conflict of interest: The authors have declared that no conflict of interest exists.


Autoimmunity, lymphopenia, and homeostasis

Autoimmunity develops when immune cells attack and destroy our own cells. For instance, type 1 diabetes mellitus (T1DM) is a disease caused by immune-mediated destruction of the insulin-secreting pancreatic β cells, leading to hyperglycemia. The immediate conse-quences of untreated T1DM (e.g., diabetic ketoacidosis) can be fatal, and, despite insu- lin therapy, long-term vascular complic-ations can significantly affect life expectancy. Autoreactive T cells specific for β cell auto-antigens are present in most individuals, but are usually kept under control by peripheral tolerance mechanisms or have a regulatory phenotype (1). It is now well established that T1DM is the result of interactions between susceptibility genes (especially those within the HLA region) and probably several envi-ronmental factors (2). The highest-risk geno-types, HLA-DR3 or HLA-DR4 class II alleles, can be responsible for the development of anti-islet autoimmunity via production of autoantibodies to insulin; to glutamic acid