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NEWS & VIEWS

CTL to HIV-1: Surrogates or sirens

The generation of cytotoxic T lymphocyte response may be an essential part of an effective immune response to HIV infection. It may also do more harm than good (pages 330–336).

The recognition that cytotoxic T lymphocytes (CTL) might be an important surrogate marker or correlate for immune protection against HIV-1 infection or disease has gradually evolved. Several observations have supported the incorporation of CTL induction as a goal in HIV-1 vaccine design. First, CD8+, MHC class Irestricted CTL can be a powerful tool in the host's armamentarium to blunt the effects of a number of viral diseases (reviewed in Koenig et al.¹, see p. 329 of this issue of Nature Medicine). Recently, a few infected persons^{2,3} (and Connick, E. et al., manuscript submitted) and monkeys4 have been documented to have detectable precursor CTL at the time of initial HIV-1 or SIV clearance. However, detectable neutralizing antibody to autologous or other primary HIV isolates from the same individuals is not observed until substantially later, if at all². Second, a few, apparently uninfected infants born to HIV-infected mothers5-7 and some seronegative subjects with high risk of sexual exposure to HIV8,9 (that is, exposed but "uninfected" subjects) have been reported to have CTL directed against selected epitopes of HIV-1 proteins. Finally, information from studies of long-term survivors with high CD4+ T lymphocyte counts suggests that CTL may function by limiting virus spread or replication^{10,11} reviewed in Schrager et al.¹² (Rinaldo, C.R. et al., Harrer, T. et al. and Riviere, Y. et al. manuscripts submitted).

In general, functionally active CD8⁺ CTL to infectious viruses can be generated quickly de novo. In HIV or SIV infections, this appears to be within days to a few weeks2.4. In addition, HIV-specific CTL activity can be detected in the peripheral circulation, without in vitro stimulation, in most HIV-infected subjects^{13,14}. A quiescent memory/precursor population can be stored in lymphoid tissues as a CTL 'reserve' for rapid recall. When functionally active, CD8⁺ CTL produce a substantial cytokine response that can profoundly depress viral replication for many viruses. HIV-specific CTL clones have been shown to produce interferon gamma and several other antiviral cytokines¹⁵. However, in an HIV-1-infected person, these cytokines may provide

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stimulatory signals for viral replication. Koenig *et al.*¹ observed a brief viral increase, despite the apparently successful delivery of the therapy.

To induce CTL, vaccine strategies must introduce viral protein epitopes into the MHC class I antigen-processing pathway. In addition to infection with the virus itself or even an attenuated virus¹⁶, AIDS vaccines designed with live, replicating viral or bacterial vectors, such as vaccinia^{17,18}, BCG^{19,20}, or Salmonella have been reported to induce CTL. Non-replicating viral vectors, such as canary pox (ALVAC) have also been reported to induce CTL (Francini, G. et al. and Riviere, Y. et al., personal communication). In addition, some adjuvants that enable antigen to enter cytoplasmic pathways have enabled CTL induction to AIDS virus antigens^{21,22}. Finally, recent studies with nucleic acid vaccines indicate that CTL can be generated by this strategy²³ (Robinson, H. et al., personal communication). Virtually all of these vaccine strategies that exhibited at least memory/precursor CTL have already failed to protect animals unequivocally (data presented by the cited groups of investigators at the Conference on AIDS Vaccine Development, November 6-10, 1994).

Somewhere in the rush to design HIV-1 vaccines that would induce CTL, the mode of action was left in the shadows, that is, CTL kill MHC class I-matched infected cells. They could never prevent infection unless they were (1) in an effector state of stimulation at the time of viral challenge, (2) at the site of infection, and most critically, (3) matched for MHC class I. Thus the question is as follows: How rapidly can virus-specific CTL come into play in a 'CTL' vaccine-primed host compared to either a "T helper (Th)" primed host or a totally naive host? Will a delay for de novo generation of CTL effectors draw the host onto rocky shores? Can this be ameliorated by some antibody that slows down the virus?

Further, as Koenig *et al.*^{*i*} illustrate, there is concern that CTL can contribute to the pathogenesis of HIV-1 disease

when the balance is tipped in favour of already widely dispersed viral an pathogen. The balance in favour of the host is further compromised, when there is a highly focused, possibly unique, immune response¹ — in this case the cloned CTL. The virus then only has to evade a single 'enemy' by simple variation, which HIV-1 has repeatedly demonstrated is its forte for monovalent drug therapy. Because the range of CTL recognition is tightly controlled by an individual's MHC genes, even when responses are successfully induced by virus infection, CTL may use only one T cell receptor beta family, or be limited to only one or two epitopes on a single viral protein²⁴.

Thus, clonal CTL, as therapies, may be like mythical Sirens, offering the ability to lyse virus-infected cells at the cost of the host's cells. The solution may lie in inducing an increased breadth of the immune response (as proposed by others¹⁴), increasing the quality of the response (that is, the precise specificity), or increasing the height of the response. The latter is problematic because data on attenuated viruses indicate that more replication produces better CTL¹⁶.

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Division of AIDS

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Filling in the matrix of kidney disease

The discovery of a gene associated with autosomal dominant polycystic kidney disease (ADPKD) was only the first step toward understanding the disease. Now researchers are determining what the gene' product normally does (pages 359–364).

Autosomal dominant polycystic kidney disease (ADPKD) affects 1 in 1,000 individuals,

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counseling is particularly vexing since the current diagnostic test that achieves 95%

making it one of the most common genetic diseases afflicting mankind, and the most common genetic cause of renal failure. The major manifestation of the disease is the formation of fluid filled cavities or 'cysts' within the kidney parenchyma. Often appearing first in neonates, the cysts expand over time, resulting in significantly enlarged kidneys and a host of related symptoms, including pain and bleeding episodes. The cysts also offer sanctuary to bacteria, and the resultant bacterial infections can be difficult to treat. Individuals afflicted with ADPKD face the prospect of renal failure, sudden death due to intracerebral bleeds and the possibility of passing the trait on to their children¹. Genetic

confidence is a computer axial tomography (CAT) scan of the kidneys using radiocontrast dye after the patient has reached age 30 (refs 2,3). Since 1957, when Dalgaard first described the disease⁴, there has been little to offer patients because of the paucity of therapeutic interventions and the incomplete understanding of the underlying molecular genetics.

In the past year, advances in molecular genetics have dramatically changed the outlook on ADPKD. Pioneering work that identified a 750-kb restriction fragment length polymorphism (RFLP) from human chromosome 16 (ref. 5) led directly to the identification of the PBP (polycystic breakpoint product) gene by

The cystic kidney shown is markedly enlarged (15 cm long) and the surface is studded with large fluid-filled sacs. Inset: Photomicrograph of a tissue section derived from a kidney with ADPKD.

Photographs courtesy of Frank Carone, Northwestern University Medical School, Evanston, Illinois, USA

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