



Published in final edited form as:

Nat Med. ; 17(8): 989–995. doi:10.1038/nm.2422.

Protective HIV-specific CD8⁺ T cells evade T_{reg} cell suppression

Shokrollah Elahi¹, Warren L Dinges^{1,2}, Nicholas Lejarcegui¹, Kerry J Laing³, Ann C Collier⁴, David M Koelle^{3,4,5,6,7}, M Juliana McElrath^{3,4,5,6}, and Helen Horton^{1,4,5}

¹Viral Vaccine Program, Seattle Biomedical Research Institute (Seattle Biomed), Seattle, Washington, USA.

²Polyclinic Infectious Disease, Seattle, Washington, USA.

³Vaccine and Infectious Diseases Division, Fred Hutchinson Cancer Research Center, Seattle, Washington, USA.

⁴Department of Medicine, University of Washington, Seattle, Washington, USA.

⁵Department of Global Health Medicine, University of Washington, Seattle, Washington, USA.

⁶Department of Laboratory Medicine, University of Washington, Seattle, Washington, USA.

⁷Benaroya Research Institute, Seattle, Washington, USA.

Abstract

Specific human leukocyte antigens (HLAs), notably *HLA-B*27* and *HLA-B*57* allele groups, have long been associated with control of HIV-1. Although the majority of HIV-specific CD8⁺ T cells lose proliferative capacity during chronic infection, T cells restricted by *HLA-B*27* or *HLA-B*57* allele groups do not. Here we show that CD8⁺ T cells restricted by 'protective' HLA allele groups are not suppressed by T_{reg} cells, whereas, within the same individual, T cells restricted by 'nonprotective' alleles are highly suppressed *ex vivo*. This differential sensitivity of HIV-specific CD8⁺ T cells to T_{reg} cell-mediated suppression correlates with their expression of the inhibitory receptor T cell immunoglobulin domain and mucin domain 3 (Tim-3) after stimulation with their cognate epitopes. Furthermore, we show that *HLA-B*27*- and *HLA-B*57*-restricted effectors also evade T_{reg} cell-mediated suppression by directly killing T_{reg} cells they encounter in a granzyme B (GzmB)-dependent manner. This study uncovers a previously unknown explanation for why *HLA-B*27* and *HLA-B*57* allele groups are associated with delayed HIV-1 disease progression.

INTRODUCTION

Cytotoxic T lymphocytes (CTLs) are crucial in controlling HIV-1 infection (reviewed in ref. 1), which is emphasized by the influence of human leukocyte antigen (HLA) class I alleles on the rate of progression to AIDS^{2,3}. Long-term nonprogressors (LTNPs) are a rare, heterogeneous population of chronically infected individuals with low viral loads and sustained CD4⁺ T cell counts who are relatively spared from disease progression even without antiretroviral therapy. The mechanisms allowing for such control of HIV-1 are not known, but a number of specific HLA allele groups, notably *HLA-B*27* and *HLA-B*57*, are

Correspondence should be addressed to H.H. (helen.horton@seattlebiomed.org).

AUTHOR CONTRIBUTIONS

S.E. designed and performed all the experiments and wrote part of the manuscript. N.L. assisted S.E. to perform some of the experiments. W.L.D. performed statistical analysis and graphing design. K.J.L. performed epitope mapping for individuals infected with HSV. D.M.K. advised on the HSV experiment. D.M.K., K.J.L., M.J.M. and A.C.C. supplied samples from subjects. H.H. designed and supervised all of the research and wrote the manuscript. All authors revised and edited the manuscript.

highly enriched in LTNP populations⁴. Possession of HIV-1–specific CD8⁺ CTLs restricted by HLA-B*27 and HLA-B*57 during early HIV-1 infection correlates with longer AIDS-free survival⁵ and better defines disease progression compared to HLA genotype alone, suggesting that HIV-specific CD8⁺ CTLs, and not the presence of a particular HLA allele, determines disease progression⁵.

LTNPs possess HIV-specific CD8⁺ CTLs restricted by HLA-B*27 or HLA-B*57 that can continue to proliferate throughout chronic infection, whereas the majority of HIV-specific CD8⁺ CTLs restricted by other HLA alleles lose their proliferative capacity^{6, 7, 8}. Proliferative ability is linked to upregulation of perforin and is, therefore, associated with the cytotoxic capabilities of virus-specific CD8⁺ CTLs⁶. We show here that HLA-B*27– and HLA-B*57–restricted HIV-specific CD8⁺ CTLs possess an additional feature in that they evade suppression mediated by T_{ref} cells. T_{ref} cells, CD4⁺ (or CD8⁺) T cells with suppressive activity, have a key role in maintaining peripheral tolerance, preventing autoimmune diseases and limiting chronic inflammatory responses (reviewed in refs. 9,10). However, they may also have detrimental effects by suppressing effective antigen-specific immune responses. Several mechanisms are used by T_{ref} cells to exert immunosuppressive function; however, our observation of differential suppression of CD8⁺ CTLs restricted by distinct HLA alleles has not been previously described to our knowledge.

Tim-3 negatively regulates T helper type 1 responses upon interaction with its ligand, galectin-9 (Gal-9, encoded by *LGALS9*) expressed on lymphocytes and other cell types¹¹. In humans, defects in Tim-3 expression contribute to multiple sclerosis pathology¹², suggesting that expression of Tim-3 on effector T cells is involved in inducing and maintaining peripheral tolerance of these T cells. T_{ref} cells constitutively express Gal-9 (ref. 13) and, thus, could be providing the ligand for inducing tolerance in Tim-3–expressing effectors. Furthermore, a recent study demonstrated that Tim-3 expression defines a distinct population of dysfunctional effector T cells in progressing HIV-1–infected individuals¹⁴.

Here we show that T_{ref} cells suppress proliferation of nonprotective HIV-specific CD8⁺ CTLs through Tim-3–Gal-9 interactions during chronic infection, but they do not suppress proliferation of HIV-specific CD8⁺ CTLs restricted by HLA-B*27 or HLA-B*57. These HLA-B*27– or HLA-B*57–restricted CTLs can continue to proliferate and kill infected targets during chronic infection, which may account for delayed disease progression in persons with *HLA-B*27* and *HLA-B*57* allele groups. In addition, this finding may explain why the *HLA-B*27* allele group is also associated with resolution of other chronic infections (for example, hepatitis C virus) and why both *HLA-B*27* and *HLA-B*57* allele groups are associated with autoimmunity^{15, 16, 17}.

RESULTS

T_{ref} cells disparately suppress CTLs based on HLA allele

To assess the susceptibility of HIV-1–specific CTLs to T_{ref} cell–mediated suppression, we measured proliferation of numerous HIV-1 epitope–specific CD8⁺ CTLs from LTNPs (individual clinical data are listed in Supplementary Table 1; epitope responses measured are listed in Supplementary Table 2) by measuring carboxyfluorescein succinimidyl ester (CFSE) dilution in the presence or absence of T_{ref} cells^{8, 18} *in vitro*. T_{ref} cells differentially suppressed proliferation of CD8⁺ CTLs restricted by different HLA alleles (Fig. 1). Epitope-specific CD8⁺ CTLs restricted by the protective HLA allele groups (HLA-B*27 and HLA-B*57) were not suppressed by T_{ref} cells ($P = 0.9$ and $P = 0.12$, respectively; median percentage T_{ref} cell suppression –1.0 (indicating that effector T cells proliferated more in the presence of T_{ref} cells than in their absence) and 13.3, respectively), whereas proliferation of epitope-specific CD8⁺ CTLs restricted by other HLA-A and HLA-B alleles (control

alleles) was significantly ($P = 0.0005$) suppressed (Fig. 1a,c). The control allele groups were *HLA-A*02*, *HLA-A*24* and *HLAB*08*, allele groups with at least three epitope responses found in at least two HIV-1 infected individuals. Of note, responses were differentially susceptible to suppression, with CTLs restricted by *HLA-A*03* being particularly susceptible ($P = 0.0005$; Fig. 1a; median percentage T_{ref} cell suppression 92.8). Of note, T_{ref} cells suppressed interferon- γ (IFN- γ) secretion from all $CD8^+$ CTLs (Fig. 1b,d; $P = 0.02$ for *HLA-B*27*-restricted responses and $P = 0.005$ for *HLAB*57*, *HLA-A*03* and control allele group). This suggests that only the proliferative ability of effector $CD8^+$ CTLs is differentially suppressed by T_{ref} cells, whereas cytokine secretion is always suppressed, regardless of the HLA restriction.

We also observed differential suppression of HIV-specific CTLs within individuals (for example, subjects NP02 and NP41, Fig. 1e). Thus HIV-specific CTLs restricted by *HLA-B*27* or *HLAB*57* were not suppressed by T_{ref} cells, whereas HIV-specific CTLs restricted by other alleles from the same individual were suppressed, suggesting that the suppression was due to a difference in the CTLs (for example, in the strength of T cell receptor (TCR) signaling) rather than the T_{ref} cells.

Although many individuals with *HLA-B*57* allele group have longer AIDS-free survival than individuals with other HLA alleles, the majority will eventually progress to disease. Indeed, it has been shown that *HLA-B*57⁺* LTNPs who eventually progress to disease have *HLA-B*57*-restricted T cells with lower proliferative capacity when compared to *HLA-B*57⁺* LTNPs who have not progressed⁸. This reduced proliferation correlated with an increased susceptibility to T_{ref} cell-mediated suppression when we compared HIV-specific, *HLA-B*57*-restricted $CD8^+$ CTLs in *HLA-B*57⁺* LTNPs versus *HLA-B*57⁺* delayed progressors ($P = 0.0002$; Fig. 1f). Longitudinal analyses before and after progression in three individuals confirmed that when clinical progression to disease occurred, previously T_{ref} cell-resistant *HLA-B*57*-restricted CTLs became susceptible to T_{ref} cell-mediated suppression ($P = 0.004$; Fig. 1g).

Differential suppression is independent of CTL frequency

*HLA-B*27*- and *HLA-B*57*-restricted $CD8^+$ CTLs proliferate more than HIV-specific $CD8^+$ CTLs restricted by other HLAs^{8, 19}. Thus, cells capable of high proliferation may escape T_{ref} cell-mediated suppression regardless of their allele restriction. We examined whether percentage suppression by T_{ref} cells (defined in Methods) correlated with either the initial precursor frequency of epitope-specific $CD8^+$ CTLs (measured by IFN- γ enzyme-linked immunosorbent spot (ELISPOT) assay) or with magnitude of proliferation. Only about 10% of the data could be explained by either initial precursor frequency ($r^2 = 0.1$, $P = 0.01$; Supplementary Fig. 1a) or magnitude of proliferation ($r^2 = 0.13$, $P = 0.003$; Supplementary Fig. 1b) suggesting that neither initial precursor frequency nor magnitude of proliferation has a substantial role in determining the susceptibility of the CTLs to T_{ref} cell-mediated suppression.

Because all of the *HLA-A*03*-restricted responses were of low magnitude (below 3% CFSE^{lo} $CD8^+$ CTLs) (Fig. 1a), we extended these analyses to look at only low-frequency responses below 3%. Even then, only 21% of the data variations were explained by the magnitude of proliferation ($r^2 = 0.21$, $P = 0.0004$; Supplementary Fig. 1c), and this weak correlation seemed to be mainly driven by *HLA-A*03*-restricted responses ($n = 11$; $r^2 = 0.82$, $P < 0.0005$), as responses restricted by *HLA-B*57* were not correlated with their ability to be suppressed ($n = 12$; $P = 0.2$). These data suggested that for some low-frequency responses (for example, restricted by *HLA-A*03*), the magnitude of proliferation might explain susceptibility to T_{ref} cell-mediated suppression, with lower frequency responses

being more susceptible. However, this does not explain why HLA-B*27- and HLA-B*57-restricted T cells are resistant to T_{ref} cell-mediated suppression.

T_{ref} cells mediate differential suppression via Gal-9-Tim-3

Upon activation with their cognate epitopes, effector CD8⁺ CTLs upregulate many inhibitory receptors (for example, programmed death-1 (PD-1) and Tim-3) as a means of self-limiting the inflammatory response. T_{ref} cells constitutively express the ligands for Tim-3 (Gal-9) and PD-1^{20, 21}. We hypothesized that the differential sensitivity to T_{ref} cell-mediated suppression could be due to differential upregulation of inhibitory receptors on CD8⁺ CTLs restricted by HLAB*27 and HLA-B*57 versus CD8⁺ CTLs restricted by other HLA alleles. Tim-3 was of particular interest, as Tim-3-knockout mice are resistant to peripheral tolerance²², and elevated frequencies of dysfunctional Tim-3⁺CD8⁺ CTLs are found in HIV-infected individuals with progressive disease¹⁴. Furthermore, although in recent years PD-1 has been the primary marker for 'exhausted' T cells²³, new data have shown that Tim-3 expression correlates better with dysfunctional CD8⁺ CTLs than expression of PD-1 (ref. 24).

Peripheral blood mononuclear cells (PBMCs) from LTNPs are not exhausted and do not constitutively express PD-1 or Tim-3. However, they do upregulate these inhibitory receptors after antigenic stimulation. We stimulated PBMCs with HIV-1 epitopes bound by various HLAs and determined the frequency of Tim-3 expression on antigen-specific CD8⁺ CTLs (as measured by either tetramer staining (Fig. 2a) or CD137 upregulation²⁵ (Fig. 2b)). As predicted, HIV-specific CD8⁺ CTLs restricted by HLA-B*27 or HLA-B*57 allele groups upregulated significantly less Tim-3 after stimulation than HIV-specific CD8⁺ CTLs restricted by other HLA alleles (Fig. 2c, $P = 0.005$), suggesting that T_{ref} cells may mediate differential suppression of effector CD8⁺ CTLs via Gal-9-Tim-3 interactions. To test this, we masked Gal-9 on T_{ref} cells by adding lactose (Fig. 3), which is a ligand of the galectin family of lectins, to which Gal-9 belongs. Addition of lactose during *in vitro* proliferation assays²⁶ prevented T_{ref} cell-mediated suppression of proliferation of HIV-specific CD8⁺ CTLs restricted by any HLA alleles other than the HLA-B*27 or HLA-B*57 allele groups but had no effect on proliferation of HIV-specific CD8⁺ CTLs restricted by HLA-B*27 or HLA-B*57 (Fig. 3a,c). Because lactose is not specific for Gal-9, but rather binds all lectins, we also isolated T_{ref} cells from PBMCs, silenced *LGALS9* using siRNA (Supplementary Fig. 2) and then added the cells back to T_{ref} cell-depleted PBMCs, followed by stimulation with their cognate HIV-1 epitopes. In agreement with the lactose studies, knockdown of *LGALS9* in T_{ref} cells prevented their suppression of proliferation of HIV-specific CD8⁺ CTLs restricted by any non-HLA-B*27 or non-HLA-B*57 allele groups but had no effect on their suppression of proliferation of HIV-specific CD8⁺ CTLs restricted by HLAB*27 or HLA-B*57 (Fig. 3b,d). Therefore, differential suppression of proliferation of HIV-specific CD8⁺ CTLs by T_{ref} cells is mediated through Gal-9-Tim-3 interactions; HLA-B*27- or HLA-B*57-restricted HIV-specific CD8⁺ CTLs upregulate less Tim-3 when they encounter their cognate epitopes than HIV-specific CTLs restricted by other HLAs, and, thus, they are less susceptible to T_{ref} cell-mediated suppression.

Protective CTLs kill T_{ref} cells via granzyme B

A recent study showed that HIV-specific HLA-B*27- and HLA-B*57-restricted CD8⁺ CTLs upregulate more granzyme B (GzmB, encoded by *GZMB*) on a per-cell basis than HIV-specific CD8⁺ CTLs restricted by other HLAs¹⁹. This upregulation is independent of their enhanced proliferative ability. Moreover, it has also been shown that CD4⁺ effectors that express high amounts of GzmB escape T_{ref} cell-mediated suppression by killing T_{ref} cells that they encounter²⁷. This mechanism of T_{ref} cell evasion has not been observed for human CD8⁺ effector cells. To determine whether this mechanism could also be involved in

the observed allele-specific differential suppression by T_{ref} cells, we performed epitope-specific CFSE dilution assays in the presence of z-AAD-CMK (z-Ala-Ala-Asp(OMe)-chloromethyl ketone), a potent and selective GzmB peptide inhibitor. When we preincubated $CD8^+$ T cells with z-AAD-CMK, $CD8^+$ CTLs restricted by HLA-B*27 and HLA-B*57 became susceptible to T_{ref} cell suppression (Fig. 4a and Supplementary Fig. 3). Similarly, if we silenced *GZMB* in $CD8^+$ CTLs with *GZMB* siRNA (Supplementary Fig. 2b) before stimulation with cognate epitope in the presence or absence of T_{ref} cells, the HLA-B*27- and HLA-B*57-restricted $CD8^+$ CTLs were rendered susceptible to T_{ref} cell-mediated suppression, whereas $CD8^+$ CTLs restricted by HLA-B*27 and HLA-B*57 treated with an irrelevant siRNA were not susceptible to T_{ref} cell suppression (Fig. 4b and Supplementary Fig. 3b). This suggests that either expression of Tim-3 or lack of expression of GzmB in $CD8^+$ effector CTLs may increase their susceptibility to T_{ref} cell-mediated suppression. Furthermore, the resistance of HLA-B*27- and HLA-B*57-restricted $CD8^+$ CTLs to T_{ref} cells resulted from their directly killing T_{ref} cells when they were stimulated with their cognate epitopes, whereas we observed low killing of T_{ref} cells in cultures stimulated with non-HLA-B*27 and non-HLA-B*57 epitopes (Fig. 5a). When we blocked GzmB activity with z-AAD-CMK, the frequency of apoptotic T_{ref} cells was reduced greatly in cultures stimulated with HLA-B*27- and HLA-B*57-restricted epitopes compared with the absence of GzmB inhibitor (Fig. 5a). In contrast, blockade of GzmB activity in cultures stimulated with non-HLA-B*27 and non-HLA-B*57 epitopes had minimal effect on the frequency of apoptotic T_{ref} cells. Thus, HLA-B*27- and HLA-B*57-restricted effector $CD8^+$ CTLs killed the T_{ref} cells they encountered in a GzmB-dependent manner. Killing mediated by HLA-B*27- and HLAB*57-restricted HIV-specific $CD8^+$ CTLs was not limited to T_{ref} cells, as they also killed non- T_{ref} $CD4^+$ cells (Supplementary Fig. 4). HLA-B*27 and HLA-B*57-restricted CTLs killed considerably more T_{ref} cells than non-HLA-B*27 and HLA-B*57-restricted CTLs (Fig. 5a). Notably, both HLA-B*27 or HLA-B*57-restricted effectors and non-HLA-B*27 or HLAB*57-restricted effectors induced more apoptosis of T_{ref} cells than of non- T_{ref} cells ($P < 0.0001$ for HLA-B*27 or HLA-B*57-restricted effectors and $P = 0.001$ for non-HLA-B*27 or HLAB*57-restricted effectors; Supplementary Fig. 4). Thus, T_{ref} cells are not seen as ' T_{ref} cells' by protective $CD8^+$ CTLs but rather as any other $CD4^+$ T cell presenting their epitope.

Maximal expression of lytic effector molecules (perforin and GzmB) requires many days of proliferation^{6, 19}. This explains why T_{ref} cells were able to suppress the IFN- γ secretion capabilities of HLA-B*27- and HLA-B*57-restricted T cells, as there would be insufficient GzmB upregulation to kill T_{ref} cells during this short assay (24 h). In agreement with this, we did not observe killing of T_{ref} cells in cultures stimulated for 24 h with HLA-B*27 and HLA-B*57 epitopes (Fig. 5b). However, after 72 h, we observed a substantial increase in the killing of T_{ref} cells by HLA-B*27-restricted $CD8^+$ CTLs (Fig. 5c).

In agreement with the data showing that HLA-B*57- restricted $CD8^+$ CTLs became susceptible to T_{ref} cell-mediated suppression of proliferation upon disease progression, HIV-specific HLAB*57-restricted $CD8^+$ CTLs from delayed progressors failed to kill T_{ref} cells ($P = 0.0003$; Fig. 5d). Analyses before and after progression in *HLA-B*57+* individuals confirmed that when clinical progression to disease occurred, HLA-B*57-restricted HIV-specific CTLs lost their ability to kill T_{ref} cells ($P = 0.008$; Fig. 5e,f).

The fact that stimulated HLA-B*27- and HLA-B*57-restricted $CD8^+$ CTLs kill T_{ref} cells suggested that there should be fewer T_{ref} cells in individuals who express these HLA alleles. Indeed, we found a significant decrease ($P = 0.0003$) in T_{ref} cell frequency in *HLA-B*27+* or *HLA-B*57+* LTNP compared with non-*HLA-B*27+* and non-*HLA-B*57+* LTNP and seronegative individuals (Fig. 5g), suggesting that deletion of T_{ref} cells occurs *in vivo* in

*HLA-B*27+* and *HLA-B*57+* individuals despite their maintenance of normal CD4⁺ T cell counts.

Our data show that *HLA-B*27-* and *HLA-B*57-*restricted CTLs are resistant to T_{ref} cell-mediated suppression irrespective of the epitopes recognized by the T cells (Fig. 1a and Supplementary Fig. 5), suggesting that resistance to T_{ref} cell-mediated peripheral tolerance is related to allele restriction rather than epitope specificity. To determine whether this occurs only during HIV infection, we assessed T_{ref} cell susceptibility of *HLA-B*27-* restricted versus *HLA-A*02-* or *HLA-B*07-*restricted herpes simplex virus (HSV-2) or Epstein Barr virus (EBV) epitopes in HIV-seronegative individuals. HSV- and EBV-specific *HLA-B*27-*restricted CD8⁺ CTLs were also not susceptible to T_{ref} cell-mediated suppression, whereas HSV-2- and EBV-specific CD8⁺ CTLs restricted by other alleles were susceptible (Supplementary Fig. 6), proving that ability to evade T_{ref} cells is due to the allele restriction of the CTL and not to the specific infection.

DISCUSSION

Although the role of human T_{ref} cells in preservation of self-tolerance is well documented, less is known about their influence on chronic viral infection. Several studies have shown that T_{ref} cell frequencies are either increased^{28, 29} or decreased in HIV infection^{30, 31}. Despite the conflicting data, some recent *in vivo* studies suggest that high frequencies of T_{ref} cells during HIV infection are detrimental. Specifically, a recent *in vivo* phase 3 trial showed that IL-2 therapy preferentially expands T_{ref} cells in infected individuals and that individuals with the greatest expansion are more likely to progress to disease^{32, 33}.

We show that there are two divergent outcomes for HIV-specific CD8⁺ CTLs during chronic infection: the majority of HIV-specific CTLs upregulate Tim-3 when they encounter their cognate epitopes and are subsequently suppressed by T_{ref} cells; however, CD8⁺ CTLs restricted by protective HLA allele groups upregulate less Tim-3 but more GzmB upon recognition of their cognate epitopes. They are subsequently less susceptible to T_{ref} cell-mediated suppression, instead killing T_{ref} cells they encounter. Thus, our data suggest a previously unknown model of how *HLA-B*27-* and *HLA-B*57-*restricted, HIV-specific CD8⁺ CTLs may evade T_{ref} cells and subsequently control HIV replication (Fig. 6). Moreover, we provide data showing that HIV-specific, *HLA-B*57-*restricted CD8⁺ CTLs can be suppressed by T_{ref} cells once progression to disease occurs. However, we cannot at present determine whether disease progression in these individuals is a cause or effect of loss of the ability to evade T_{ref} cells.

The mechanisms accounting for why *HLA-B*27-* and *HLA-B*57-*restricted T cells upregulate less Tim-3 and more GzmB than other HIV-specific T cells upon recognition of their cognate epitopes are not known. Understanding these differences in TCR signaling will be crucial for determining potential therapeutic interventions that 'switch on' virus-specific CTLs that are restricted by other HLAs. Such mechanisms may include a higher-avidity TCR on CD8⁺ CTLs restricted by *HLA-B*27* and *HLA-B*57* (ref. 34). In addition, *HLA-B*27* is an unusual allele group in that it can form heavy-chain homodimers and trimers, which may lead to alternative TCR interactions³⁵.

The observation that T_{ref} cells are killed more efficiently than non-T_{ref} CD4⁺ T cells suggests that, during HIV infection, T_{ref} cells are more susceptible to effector-induced apoptosis. This is consistent with studies demonstrating that T_{ref} cells have increased turnover and higher expression of caspase-3 during chronic infection³⁶. Increased intracellular levels of caspase-3 have been shown to increase susceptibility of T cells to apoptosis³⁷. Peripheral T_{ref} cell frequencies are similar between *HLA-B*27+* or *HLA-B*57+*

LTNPs and *HLA-B*27+* or *HLA-B*57+* uninfected subjects. Most of the LTNPs studied here have low or undetectable viral load in their peripheral blood, so there is minimal antigen to stimulate killing of T_{ref} cells by *HLA-B*27* or *HLA-B*57*-restricted CTLs. We presume that *HLA-B*27* or *HLA-B*57*-restricted CTLs will kill T_{ref} cells where antigen is present, so it is not surprising that we did not observe significant differences in T_{ref} cell frequencies in the periphery between uninfected and infected individuals. However, in lymph nodes and gut we would expect to see fewer T_{ref} cells in infected individuals with *HLA-B*27* or *HLA-B*57*. This is supported by data from another group showing fewer T_{ref} cells in lymph nodes from LTNPs than in those from infected individuals with progressive infection²⁹. Therefore, it is possible that loss of effector cell evasion of T_{ref} cells results in accumulation of T_{ref} cells in lymph nodes and disease progression.

Our study indicates that $CD8^+$ CTLs are differentially susceptible to T_{ref} cell-mediated suppression of proliferation dependent on HLA restriction but independent of the epitope specificity of the suppressed cell. Thus, HIV-specific CTLs restricted by *HLA-B*27* and *HLA-B*57* are resistant to T_{ref} cells, allowing them to continue to function maximally during chronic infection. This may explain why these HLAs are associated with delayed progression to disease. Resistance of *HLA-B*27*-restricted CTLs to T_{ref} cell-mediated suppression may also explain why this allele group is associated with better clinical outcomes of other chronic infections, such as hepatitis C virus¹⁵. Both *HLA-B*27* and *HLA-B*57* allele groups are highly associated with autoimmunity: *HLA-B*27* allele groups with ankylosing spondylitis¹⁶ and *HLA-B*57* allele groups with psoriasis³⁸. Thus, possession of these allele groups seems to be a double-edged sword. Because T cells restricted by them cannot be tolerized, these allele groups are beneficial in chronic infection but detrimental in autoimmunity. Therefore, we feel this work has far-reaching implications for both control of chronic infection and autoimmunity.

METHODS

Study group

LTNPs were enrolled through the HIV Vaccine Trial Unit (M.J.M.). They were defined as HIV infected for more than 11 years, with repeated $CD4^+$ T cell counts over 500 cells per μ l or $CD4\%$ over 28% and viral load $<10,000$ copies per ml in the absence of antiretroviral therapy. Clinical data and HLA genotypes are shown in Supplementary Table 1. An additional subject was recruited through the University of Washington Primary Infection Clinic (A.C.C.), meeting early criteria of an elite controller (viral load 58 copies per ml), infected for about ten years but recently progressed. We studied 17 LTNPs and 7 LTNPs who eventually progressed (delayed progressors), defined as HIV infected with viral load $>10,000$ RNA (copies per ml) or declining $CD4^+$ T cell count in the absence of antiretroviral therapy. All subjects were males of European descent except subject NP14 who reported African-American ethnicity. HIV-1 seronegative individuals were recruited at Seattle BioMed.

Fourteen HIV-seronegative individuals (seven males and seven females) who were seropositive for HSV-2 were recruited and HLA-typed as described³⁹ (Supplementary Table 1). They were assumed to be EBV seropositive if they had an IFN- γ response to known EBV $CD8^+$ epitopes because most adults are EBV seropositive.

The appropriate Institutional Review Boards at the University of Washington, Fred Hutchinson Cancer Research Center and Seattle BioMed approved the studies, and all individuals provided written informed consent.

Epitope mapping

HIV epitope specificities were mapped in each HIV⁺ individual as previously described⁸. Some LTNP were mapped using potential T cell epitopes provided by The National Institutes of Health AIDS Research and Reference Reagent Program. The epitope-specific responses assessed for each individual are shown in Supplementary Table 2. HSV-2⁺ individuals were mapped as previously described³⁹. HSV and EBV epitope-specific responses assessed for each individual are shown in Supplementary Table 2.

Cell separation and flow cytometry

We purified T_{ref} cells by magnetic separation (STEMCELL Technology) and assessed purity by flow cytometry using a LSRII flow cytometer (BD Biosciences). We analyzed flow data with FlowJo software v7.2.2 (Tree Star). To determine levels of T_{ref} cell apoptosis, we labeled epitope-stimulated PBMCs with annexin V (BD Bioscience).

In vitro cytokine measurements

We measured cytokine responses by performing ELISPOT or intracellular cytokine secretion (ICS) assays^{40, 41}. In brief, we cultured 2×10^5 (ELISPOT) and 1×10^6 (ICS) cells per well and stimulated them with $2 \mu\text{g ml}^{-1}$ of their cognate epitopes. For ELISPOT we called a response positive when the number of spot-forming cells was twice the background level and there were at least 50 spot-forming cells per 1×10^6 PBMCs. For ICS, we called a response positive when the percentage of bright cytokine⁺CD8⁺ T cells was twice that of the negative control.

Proliferation assays

We performed CFSE dilution assays as previously described⁸. Where indicated, we added T_{ref} cells (1:4) to PBMCs or CD8⁺ T cell cultures, although we obtained similar data when we used a ratio as low as 1:16 T_{ref} cells:PBMCs (that is, an approximate T_{ref} cell frequency of 6%, which we have shown is in the physiological range of T_{ref} cell frequencies *in vivo*; Supplementary Fig. 5). We performed some assays in the presence of lactose (Sigma-Aldrich; 30 mM) or the GzmB inhibitor z-AAD-CMK (Calbiochem; $10 \mu\text{g ml}^{-1}$).

Small interfering RNA transduction

For silencing of *LGALS9*, we performed RNA interference experiments on isolated T_{ref} cells according to the manufacturer's protocol (Santa Cruz Biotechnology). Knockdown of *LGALS9* in T_{ref} cells is shown in Supplementary Figure 2a. Control T_{ref} cells received a scrambled I duplex RNA. Seven hours after transfection, we replaced the transfection medium with complete medium and cultured T_{ref} cells for another 24 h before using them in CFSE proliferation assays. For silencing of *GZMB*, we electroporated PBMCs or CD8⁺ T cells with *GZMB* siRNA (Applied Biosystems/Ambion) following the manufacturer's guidelines. We rested the cells for 36 h before using them in CFSE proliferation assays. Expression of intracellular GzmB after electroporation is shown in Supplementary Figure 2b.

Statistical analyses

We calculated percentage suppression of proliferation by the formula (percentage CFSE^{lo}CD8⁺ without T_{ref} cells – percentage CFSE^{lo}CD8⁺ with T_{ref} cells) / percentage CFSE^{lo}CD8⁺ without T_{ref} cells × 100, grouped according to HLA restricting allele. Two responses with CFSE^{lo}CD8⁺ cells without T_{ref} cells were zero, and the lowest measured response (0.012) was substituted to prevent a division by zero error. We similarly calculated percentage suppression by lactose and *LGALS9* siRNA by substituting the differential suppression via these compounds. We calculated percentage suppression of cytokine

secretion by the formula (spot-forming cells per million PBMCs without T_{ref} cells–spot-forming cells per million PBMCs with T_{ref} cells) / spot-forming cells per million PBMCs without T_{ref} cells $\times 100$. We used signed Wilcoxon tests to compare paired data with and without the suppressing agent (T_{ref} cells, lactose, siRNA). For comparisons of grouped data, we used Kruskal-Wallis tests with a *post hoc* Dunn's test for subgroup comparisons. We used Spearman rank tests for correlation analyses. All tests were two-tailed, with a *P* value of less than 0.05 considered statistically significant. We conducted statistical analyses and graphing with GraphPad Prism 5.02 (GraphPad) and Adobe Illustrator Creative Suite 3, 13.0.2 (Adobe Systems).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

We thank our study volunteers for providing samples and supporting this work, as well as the clinical staff for their dedication to this research. This work was supported by US National Institutes of Health (NIH) grants R01 AI65328, R21 AI089373, U01 AI4674, U01 AI 46725, P01 AI57005, P30 AI27757, a New Investigator Award (for S.E.) from the University of Washington/Fred Hutchinson Cancer Research Center Center for AIDS Research (CFAR), AI30731, AI 081060, R37 AI042528 and M01-RR-00037 (the University of Washington General Clinical Research Center). This research was supported by the University of Washington Center for AIDS Research (CFAR), an NIH funded program (P30 AI027757) which is supported by the NIH National Institute of Allergy and Infectious Diseases, National Cancer Institute, National Institute of Mental Health, National Institute on Drug Abuse, National Institute of Child Health and Human Development, National Heart, Lung, and Blood Institute and National Center for Complementary and Alternative Medicine). We also acknowledge the support of the James. B. Pendleton Charitable Trust.

References

1. McMichael AJ, Rowland-Jones SL. Cellular immune responses to HIV. *Nature*. 2001; 410:980–987. [PubMed: 11309628]
2. Kaslow RA, et al. Influence of combinations of human major histocompatibility complex genes on the course of HIV-1 infection. *Nat. Med.* 1996; 2:405–411. [PubMed: 8597949]
3. Carrington M, et al. HLA and HIV-1: heterozygote advantage and B*35-Cw*04 disadvantage. *Science*. 1999; 283:1748–1752. [PubMed: 10073943]
4. Carrington M, O'Brien SJ. The influence of HLA genotype on AIDS. *Annu. Rev. Med.* 2003; 54:535–551. [PubMed: 12525683]
5. Dinges WL, et al. Virus-specific CD8+ T cell responses better define HIV disease progression than HLA genotype. *J. Virol.* 2010; 84:4461–4468. [PubMed: 20147397]
6. Migueles SA, et al. HIV-specific CD8+ T cell proliferation is coupled to perforin expression and is maintained in nonprogressors. *Nat. Immunol.* 2002; 3:1061–1068. [PubMed: 12368910]
7. Lichterfeld M, et al. Loss of HIV-1-specific CD8+ T cell proliferation after acute HIV-1 infection and restoration by vaccine-induced HIV-1-specific CD4+ T cells. *J. Exp. Med.* 2004; 200:701–712. [PubMed: 15381726]
8. Horton H, et al. Preservation of T cell proliferation restricted by protective HLA alleles is critical for immune control of HIV-1 infection. *J. Immunol.* 2006; 177:7406–7415. [PubMed: 17082660]
9. Vila J, Isaacs JD, Anderson AE. Regulatory T cells and autoimmunity. *Curr. Opin. Hematol.* 2009; 16:274–279. [PubMed: 19417650]
10. Galgani M, Di Giacomo A, Matarese G, La Cava A. The Yin and Yang of CD4+ regulatory T cells in autoimmunity and cancer. *Curr. Med. Chem.* 2009; 16:4626–4631. [PubMed: 19903146]
11. Zhu C, et al. The Tim-3 ligand galectin-9 negatively regulates T helper type 1 immunity. *Nat. Immunol.* 2005; 6:1245–1252. [PubMed: 16286920]
12. Koguchi K, et al. Dysregulated T cell expression of TIM3 in multiple sclerosis. *J. Exp. Med.* 2006; 203:1413–1418. [PubMed: 16754722]

13. Li XC, Turka LA. An update on regulatory T cells in transplant tolerance and rejection. *Nat. Rev. Nephrol.* 2010; 6:577–583. [PubMed: 20683480]
14. Jones RB, et al. Tim-3 expression defines a novel population of dysfunctional T cells with highly elevated frequencies in progressive HIV-1 infection. *J. Exp. Med.* 2008; 205:2763–2779. [PubMed: 19001139]
15. Neumann-Haefelin C, et al. Dominant influence of an HLA-B27 restricted CD8+ T cell response in mediating HCV clearance and evolution. *Hepatology.* 2006; 43:563–572. [PubMed: 16496339]
16. Mathieu A, et al. The interplay between the geographic distribution of HLA-B27 alleles and their role in infectious and autoimmune diseases: a unifying hypothesis. *Autoimmun. Rev.* 2009; 8:420–425. [PubMed: 19185064]
17. Henseler T. Genetics of psoriasis. *Arch. Dermatol. Res.* 1998; 290:463–476. [PubMed: 9808339]
18. Migueles SA, Connors M. Frequency and function of HIV-specific CD8(+) T cells. *Immunol. Lett.* 2001; 79:141–150. [PubMed: 11595301]
19. Migueles SA, et al. Lytic granule loading of CD8+ T cells is required for HIV-infected cell elimination associated with immune control. *Immunity.* 2008; 29:1009–1021. [PubMed: 19062316]
20. Wang F, et al. Tim-3-Galectin-9 pathway involves the suppression induced by CD4+CD25+ regulatory T cells. *Immunobiology.* 2009; 214:342–349. [PubMed: 19362679]
21. Francisco LM, Sage PT, Sharpe AH. The PD-1 pathway in tolerance and autoimmunity. *Immunol. Rev.* 2010; 236:219–242. [PubMed: 20636820]
22. Sabatos CA, et al. Interaction of Tim-3 and Tim-3 ligand regulates T helper type 1 responses and induction of peripheral tolerance. *Nat. Immunol.* 2003; 4:1102–1110. [PubMed: 14556006]
23. Barber DL, et al. Restoring function in exhausted CD8 T cells during chronic viral infection. *Nature.* 2006; 439:682–687. [PubMed: 16382236]
24. Sakuishi K, et al. Targeting Tim-3 and PD-1 pathways to reverse T cell exhaustion and restore anti-tumor immunity. *J. Exp. Med.* 2010; 207:2187–2194. [PubMed: 20819927]
25. Wehler TC, et al. Rapid identification and sorting of viable virus-reactive CD4+ and CD8+ T cells based on antigen-triggered CD137 expression. *J. Immunol. Methods.* 2008; 339:23–37. [PubMed: 18760281]
26. Chabot S, et al. Regulation of galectin-9 expression and release in Jurkat T cell line cells. *Glycobiology.* 2002; 12:111–118. [PubMed: 11886844]
27. Ashley CW, Baecher-Allan C. Cutting Edge: Responder T cells regulate human DR+ effector regulatory T cell activity via granzyme B. *J. Immunol.* 2009; 183:4843–4847. [PubMed: 19801510]
28. Andersson J, et al. The prevalence of regulatory T cells in lymphoid tissue is correlated with viral load in HIV-infected patients. *J. Immunol.* 2005; 174:3143–3147. [PubMed: 15749840]
29. Nilsson J, et al. HIV-1-driven regulatory T-cell accumulation in lymphoid tissues is associated with disease progression in HIV/AIDS. *Blood.* 2006; 108:3808–3817. [PubMed: 16902147]
30. Tsunemi S, et al. Relationship of CD4+CD25+ regulatory T cells to immune status in HIV-infected patients. *AIDS.* 2005; 19:879–886. [PubMed: 15905668]
31. Eggena MP, et al. Depletion of regulatory T cells in HIV infection is associated with immune activation. *J. Immunol.* 2005; 174:4407–4414. [PubMed: 15778406]
32. Abrams D, et al. Interleukin-2 therapy in patients with HIV infection. *N. Engl. J. Med.* 2009; 361:1548–1559. [PubMed: 19828532]
33. Weiss L, et al. *In vivo* expansion of naive and activated CD4+CD25+FOXP3+ regulatory T cell populations in interleukin-2-treated HIV patients. *Proc. Natl. Acad. Sci. USA.* 2010; 107:10632–10637. [PubMed: 20498045]
34. Almeida JR, et al. Superior control of HIV-1 replication by CD8+ T cells is reflected by their avidity, polyfunctionality, and clonal turnover. *J. Exp. Med.* 2007; 204:2473–2485. [PubMed: 17893201]
35. Antoniou AN, Ford S, Taurog JD, Butcher GW, Powis SJ. Formation of HLA-B27 homodimers and their relationship to assembly kinetics. *J. Biol. Chem.* 2004; 279:8895–8902. [PubMed: 14684742]

36. Xing S, et al. Increased turnover of FoxP3^{high} regulatory T cells is associated with hyperactivation and disease progression of chronic HIV-1 infection. *J. Acquir. Immune Defic. Syndr.* 2010; 54:455–462. [PubMed: 20585263]
37. Sabbagh L, et al. The selective increase in caspase-3 expression in effector but not memory T cells allows susceptibility to apoptosis. *J. Immunol.* 2004; 173:5425–5433. [PubMed: 15494489]
38. Feng BJ, et al. Multiple Loci within the major histocompatibility complex confer risk of psoriasis. *PLoS Genet.* 2009; 5:e1000606. [PubMed: 19680446]
39. Laing KJ, et al. Diversity in CD8⁺ T cell function and epitope breadth among persons with genital herpes. *J. Clin. Immunol.* 2010:703–722. [PubMed: 20635156]
40. Horton H, et al. Induction of human immunodeficiency virus type 1 (HIV-1)-specific T-cell responses in HIV vaccine trial participants who subsequently acquire HIV-1 infection. *J. Virol.* 2006; 80:9779–9788. [PubMed: 16973582]
41. Horton H, et al. Optimization and validation of an 8-color intracellular cytokine staining (ICS) assay to quantify antigen-specific T cells induced by vaccination. *J. Immunol. Methods.* 2007; 323:39–54. [PubMed: 17451739]

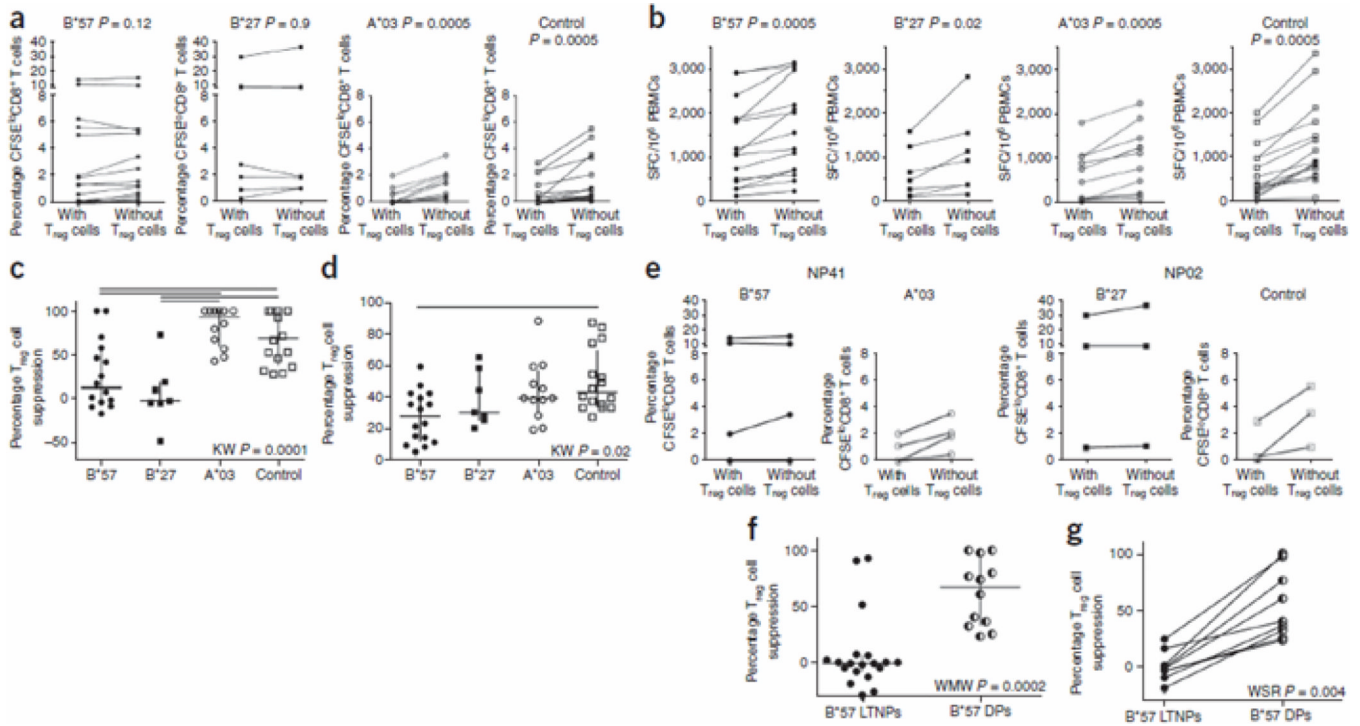
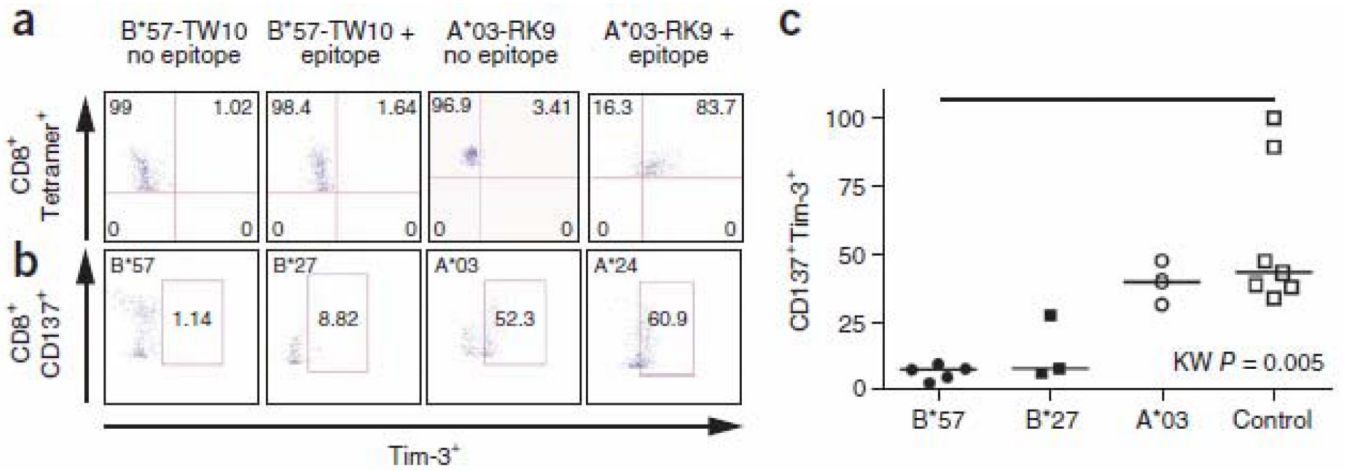
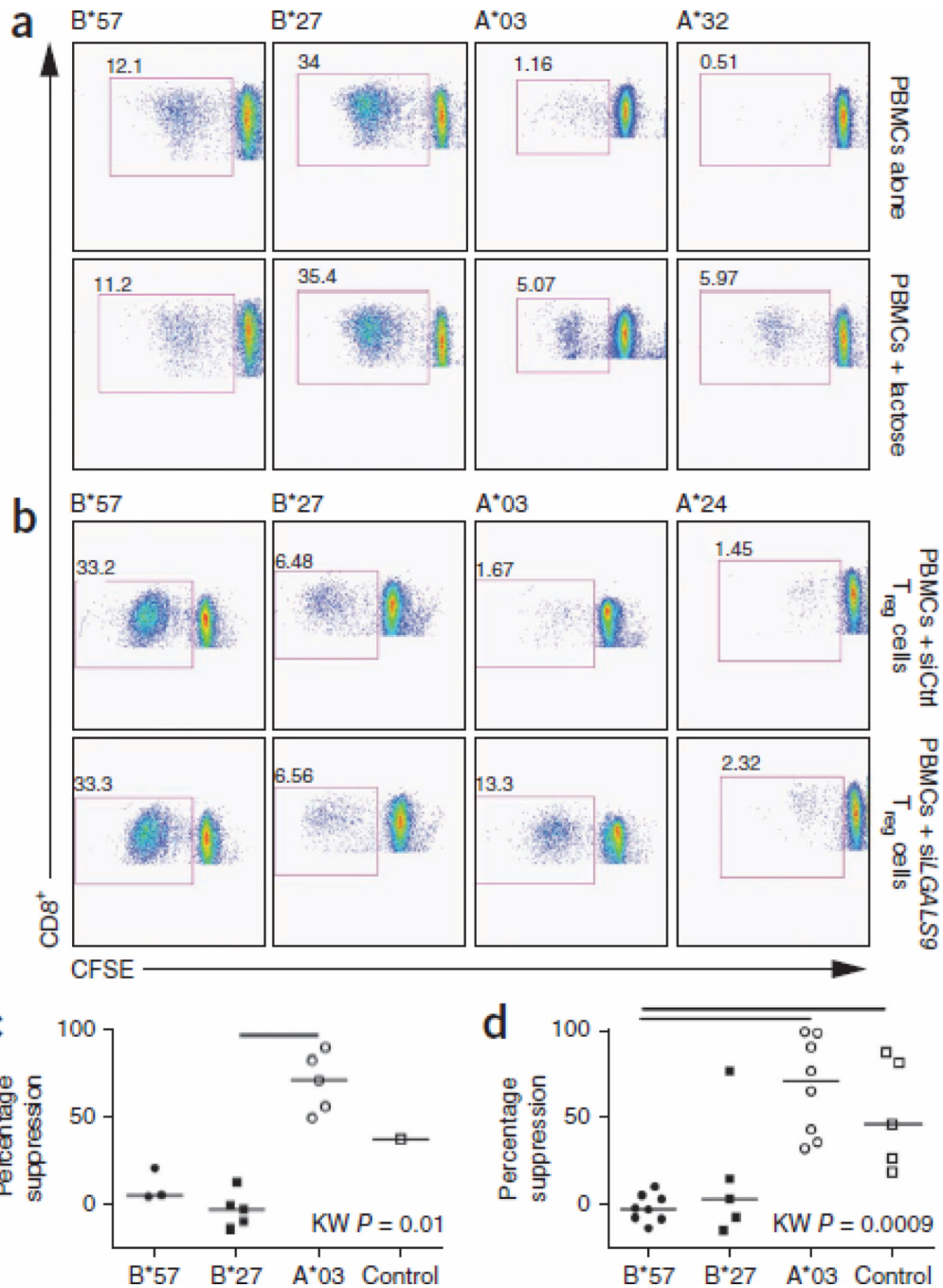


Figure 1.

Treg cell suppression of *in vitro* proliferative ability or cytokine secretion of CD8+ T cells restricted by HLA-B*57, HLA-B*27, HLA-A*03 and control HLAs (HLA-A*02, HLA-A*24 and HLA-B*08). (a) Background-subtracted percentage CFSE^{lo}CD8+ T cells in PBMCs with or without T_{reg} cells when cultured in the presence of HIV-1 epitopes recognized by CD8+ T cells restricted by various HLA alleles. (b) Background-subtracted HIV-specific IFN- γ ELISPOT responses in the presence and absence of Treg cells. HLA-B*27 or HLA-B*57-restricted and non-HLA-B*27- and HLA-B*57-restricted responses are shown after stimulation with their cognate epitopes. SFC, spot-forming cell. In a and b, Wilcoxon signed-rank (WSR) test was used. (c) Percentage suppression of proliferation grouped according to HLA restricting allele. (d) Percentage suppression of cytokine secretion grouped according to HLA restricting allele. In c and d, Kruskal-Wallis (KW) test was used for grouped comparisons with a *post hoc* Dunn's test showing significant subgroup comparisons with horizontal lines. (e) Differential suppression of proliferation of HLA-B*27- and HLA-B*57- versus HLA-A*03- and control HLA-restricted HIV-specific CD8+ CTLs within the same person. NP02 and NP41 are two LTNPs. (f) Percentage suppression of proliferation by Treg cells of HLA-B*57-restricted CD8+ CTLs in *HLA-B*57+* LTNP versus *HLA-B*57+* delayed progressors (DP). (g) Longitudinal analyses of percentage suppression of proliferation by Treg cells of HLA-B*57-restricted CD8+ CTLs before and after progression in *HLA-B*57+* individuals.

**Figure 2.**

Frequency of CD8+Tim-3+ T cells following stimulation with their cognate epitopes. **(a)** Percentage of Tim-3+ CD8+ T cells using allophycocyanin-labeled HLA-A*03–RLRPGGKKK tetramer or phycoerythrin-labeled HLA-B*57-TSTLQEIQGW tetramer staining of PBMCs before and after stimulation with their cognate epitopes. Top right quadrant shows percentage of Tim-3+tetramer+ CD8+ CTLs. **(b)** Percentage of Tim-3+ on CD8+ T cells using CD137 to identify antigen-specific T cells after stimulation with their cognate epitopes. **(c)** Percentage of CD137+Tim-3+ T cells after stimulation of PBMCs from different individuals with their corresponding epitopes.

**Figure 3.**

CFSE dilution data showing inhibition of Gal-9–Tim-3 interactions by lactose and siRNA.

(a) Examples of proliferation of PBMCs stimulated with their corresponding epitopes, showing percentage CFSE^{lo}CD8⁺ T cells in the absence or presence of lactose. (b) Examples of proliferation of CFSE-labeled, Treg cell–depleted PBMCs stimulated with their cognate epitopes in the presence of Treg cells treated with either *LGALS9* siRNA or siControl (at 1:0.25 ratio). The measures of coculture suppression by Treg cells in the presence or absence of lactose or siRNA are shown for a representative experiment from three repeat experiments for each approach. (c) Percentage of Treg cell suppression calculated after stimulation of CD8⁺ T cells with their corresponding epitopes in the

presence of lactose. **(d)** Percentage of Treg cell suppression calculated after stimulation of CD8⁺ T cells with their cognate epitopes in the presence of *LGALS9* siRNA-treated Treg cells (at 1:0.25 ratio).

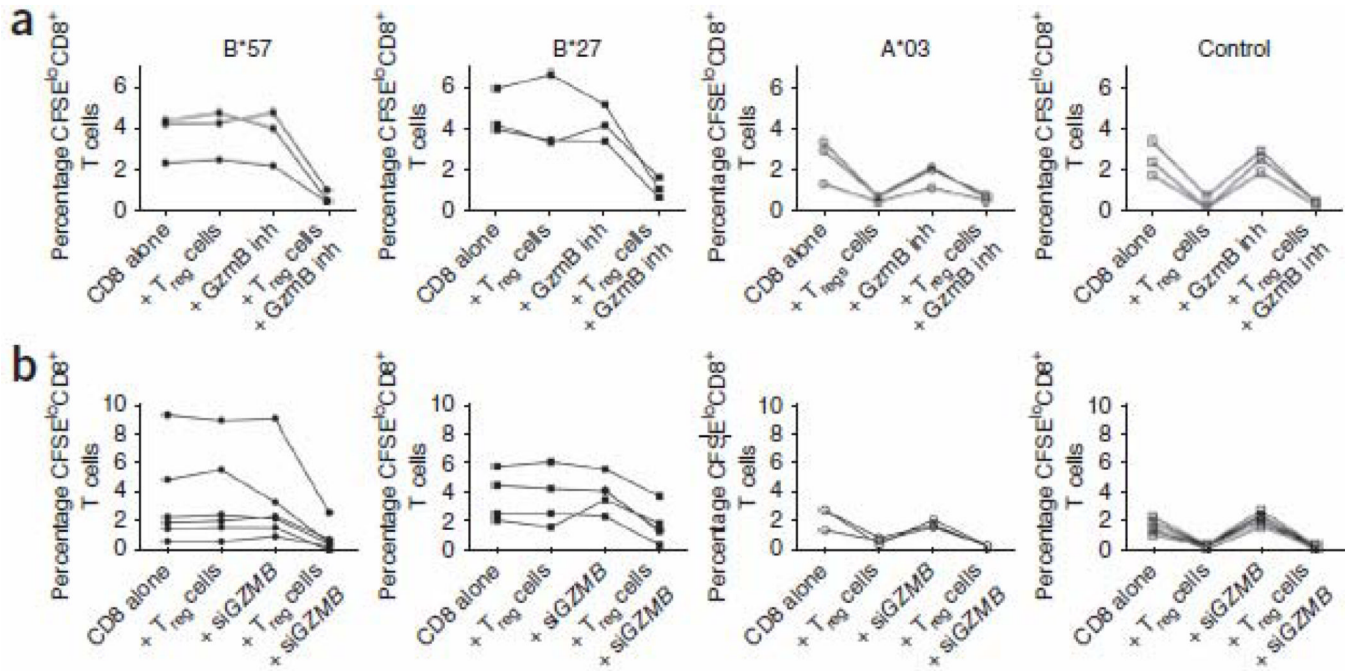
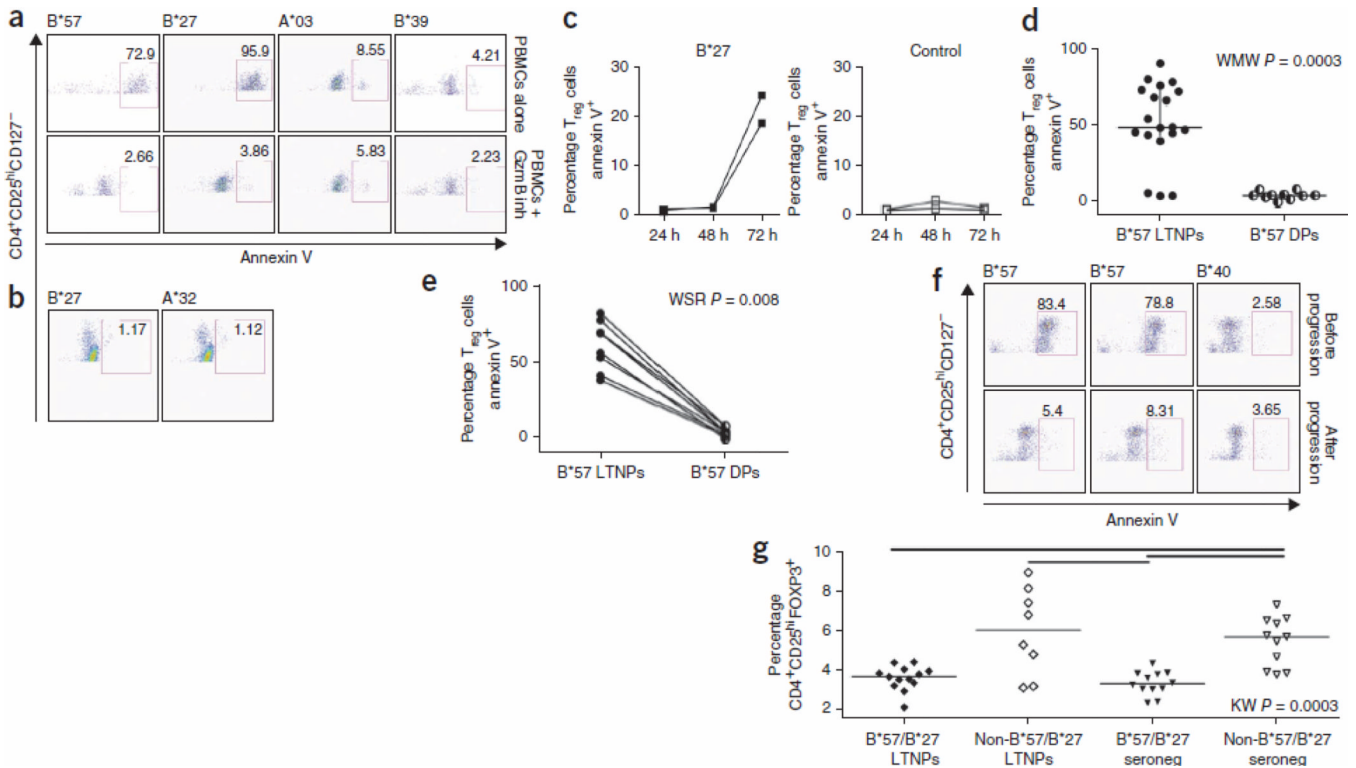


Figure 4.

CFSE dilution data showing CD8⁺ T cells restricted by HLA-B*57 and HLA-B*27 resist Treg cell-mediated suppression in a GzmB dependent manner. **(a)** Percentage CFSE^{lo}CD8⁺ T cells after CFSE-labeled isolated CD8⁺ T cells were stimulated with their corresponding epitopes alone or together with Treg cells (at 1:0.25 ratio), and also in the presence or absence of a GzmB peptide inhibitor (z-AAD-CMK). Examples of flow data are shown in Supplementary Figure 3a. **(b)** Percentage CFSE^{lo}CD8⁺ T cells after electroporation with *GZMB* siRNA or nonhybridizing negative control (siControl) siRNA oligonucleotides and stimulation with their cognate epitopes alone or with Treg cells (at 1:0.25 ratio). Examples of flow data are shown in Supplementary Figure 3b.

**Figure 5.**

CD8⁺ T cells restricted by HLA-B*27 and HLA-B*57 induce Treg apoptosis in a GzmB-dependent manner. (a–c) Percentage annexin V⁺ Treg cells (CD3⁺CD4⁺CD25^{hi}CD127^{lo}) in PBMCs stimulated for 4 d (a), 24 h (b) or 24–72 h (c) with HLA-B*27⁻, HLA-B*57⁻, HLA-B*39⁻ or HLA-A*03–restricted epitopes in the presence or absence of GzmB peptide inhibitor. These data are representative of three separate experiments from different LTNPs. (d) Percentage annexin V⁺ Treg cells in PBMCs stimulated with HLA-B*57–restricted epitopes from *HLA-B*57*⁺ LTNP versus *HLA-B*57*⁺ DPs. (e,f) Percentage annexin V⁺ Treg cells in PBMCs stimulated with HLA-B*57–restricted epitopes before and after progression to disease. (g) Treg cell frequencies in HIV-1–seronegative individuals versus HIV-1 infected *HLA-B*27*⁺ or *HLA-B*57*⁺ and *HLA-B*27*⁻ or *HLA-B*57*⁻ LTNPs. Percentages of CD4⁺CD25^{hi}FOXP3⁺ Treg cells are shown in PBMCs from 12 HIV-seronegative *HLA-B*27*⁺ or *HLA-B*57*⁺ individuals, 12 *HLA-B*27*⁻ or *HLA-B*57*⁻ individuals, 13 *HLA-B*27*⁺ or *HLA-B*57*⁺ LTNPs and 8 *HLA-B*27*⁻ or *HLA-B*57*⁻ LTNPs.

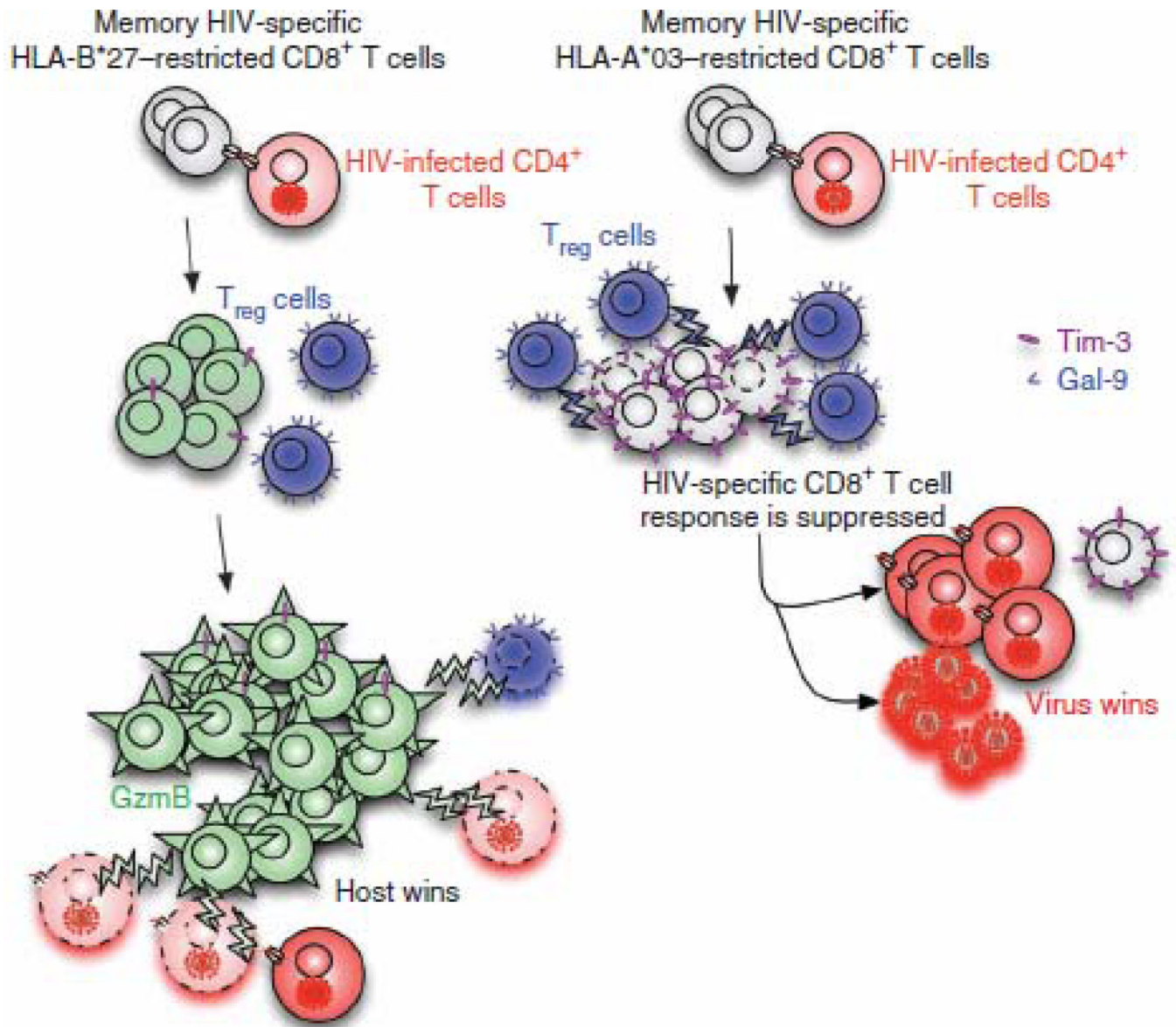


Figure 6.

Model depicting how HLA-B*27- or HLA-B*57-restricted HIV-specific CD8+ T cells evade Treg cell suppression and subsequently control HIV replication. HIV-specific, HLA-B*27-restricted CD8+ T cells do not upregulate surface expression of Tim-3 upon recognition of their cognate epitopes on HIV-infected CD4+ T cells, whereas HIV-specific, HLA-A*03-restricted CD8+ T cells upregulate high surface expression of Tim-3. Treg cells suppress HLA-A*03-restricted CD8+ T cells owing to their high expression of Tim-3 but cannot suppress proliferation of HLA-B*27-restricted CD8+ T cells. Highly proliferating HLA-B*27-restricted CD8+ T cells upregulate high levels of GzmB and kill not only infected CD4+ T cells but also infected Treg cells that they encounter. Thus, HLA-B*27-restricted CD8+ T cells can control HIV replication during chronic infection, whereas HLA-A*03-restricted CD8+ T cells cannot.