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HIV Immunopathogenesis: Inhibition of CD4+ T Cell Activation

Abstract

Human immunodeficiency virus (HIV) infection causes profound impairment of CD4+ T cell immunity. Anti-retroviral therapy (ART) restores CD4+ T cell responses to common antigens, but HIV-specific responses remain deficient. The immunization of chronically HIV infected, ART treated subjects also leads to poor HIV-specific CD4+ T cell responses. The mechanisms are not fully understood. In this thesis, I demonstrate that HIV envelope glycoprotein (Env), when delivered in the form of a vaccine or when present on free viral particles, suppresses antigen-stimulated CD4+ T cell proliferation. I investigate the potential involvement of human T regulatory cells (Treg) using an in vitro model system. I show that Env exposure neither changes the frequency nor the suppressive activity of Treg cells in human PBMC, and that Env-induced suppression of CD4+ T cell proliferation is independent of Tregs. The studies of HIV-induced inhibition of CD4+ T cell immunity were then moved to an in vivo model to determine physiological significance. A macaque model of HIV-infected individuals treated with ART during chronic infection was used to study the effect of SIV antigen stimulation in lymph nodes early after immunization. CMV seropositive rhesus macaques were infected with pathogenic SIVmac251 and after 4 months, treated with D4T and PMPA for viral control and immune reconstitution. I studied the immune and viral responses to SIV and CMV antigen immunization in draining lymph nodes. Animals were immunized with both SIV gag and CMV pp65 encoding plasmids in both arms and legs, which allowed draining lymph nodes for each antigen to be obtained at the same time, thus allowing direct comparisons of the effect of each antigen stimulation in the same animal. I observed that both SIV and CMV antigen immunizations stimulated antigen-specific T cell responses in draining lymph nodes. The CMV-specific responses were found in the periphery for 50 days post-immunization, while the SIV-specific responses transiently appeared. The SIV antigen stimulation also induced transient SIV viral replication in the draining lymph nodes, suggesting a mechanism for the early loss and poor HIV-specific CD4+ T cell response observed in HIV-infected progressors.

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HIV IMMUNOPATHOGENESIS: INHIBITION OF CD4⁺ T CELL ACTIVATION

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ABSTRACT

HIV IMMUNOPATHOGENESIS: INHIBITION OF CD4⁺ T

CELL ACTIVATION

Haitao Hu

Dissertation Supervisor: Drew Weissman, MD, PhD

Human immunodeficiency virus (HIV) infection causes profound impairment of CD4⁺ T cell immunity. Anti-retroviral therapy (ART) restores CD4⁺ T cell responses to common antigens, but HIV-specific responses remain deficient. The immunization of chronically HIV infected, ART treated subjects also leads to poor HIV-specific CD4⁺ T cell responses. The mechanisms are not fully understood. In this thesis, I demonstrate that HIV envelope glycoprotein (Env), when delivered in the form of a vaccine or when present on free viral particles, suppresses antigen-stimulated CD4⁺ T cell proliferation. I investigate the potential involvement of human T regulatory cells (Treg) using an in vitro model system. I show that Env exposure neither changes the frequency nor the suppressive activity of Treg cells in human PBMC, and that Env-induced suppression of CD4⁺ T cell proliferation is independent of Tregs. The studies of HIV-induced inhibition of CD4⁺ T cell immunity were then moved to an in vivo model to determine physiological significance. A macaque model of HIV-infected individuals treated with ART during chronic infection was used to study the effect of SIV antigen stimulation in lymph nodes early after immunization. CMV seropositive rhesus macaques were infected with pathogenic SIVmac251 and after 4 months, treated with D4T and PMPA for viral

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CHAPTER 1

Introduction

HIV and the global epidemic

Human immunodeficiency virus (HIV) is a lentivirus that causes acquired immunodeficiency syndrome (AIDS) in humans^{48, 162}. As of December 2008, between 31.1 million and 35.8 million people were believed to be living with HIV. The number of people newly infected in 2008 was estimated to be 2.7 million and approximately 2 million people died from HIV infection (AIDS epidemic update 2009, UNAIDS).

HIV is composed of two copies of single-stranded genomic RNA that are enclosed by capsid and contain nine genes encoding 19 proteins. The HIV replication cycle is summarized in Figure 1.1¹⁴⁴. First, the glycoprotein complex present on the surface of virus, known as Env or gp160, enables the virus to attach to target cells to initiate entry and infection²⁷. Entry of HIV into the host cell depends on the interaction between the viral envelope and CD4 plus a co-receptor (generally either CCR5 or CXCR4) on the surface of target cells. Binding of gp120 to CD4 leads to a conformational change in Env and the subsequent exposure of the co-receptor binding site, which allows gp120 to interact with a target chemokine receptor to mediate fusion and entry^{28, 169}.

Figure 1.1

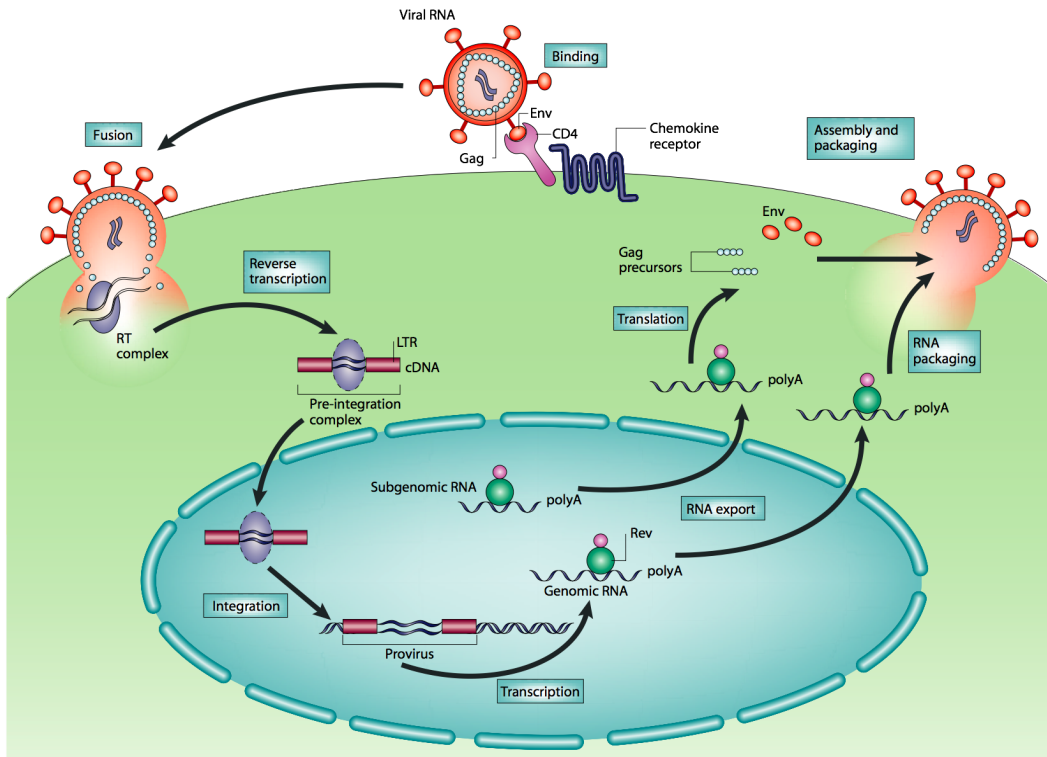


Figure 1.1. Schematic illustration of HIV replication cycle in an infected cell. LTR: long terminal repeat; RT: reverse transcriptase ¹⁴⁴

The viral RNA genome is reverse transcribed shortly after the virus enters the cell and the preintegration complex is transported into the cell nucleus and integrates into host cell's genome. Infection of CD4⁺ T cells has 2 main outcomes, lytic and latent. In latent infection, the integrated viral DNA may lie "dormant" in resting memory cells. When such cells become activated, the integrated DNA provirus starts viral gene expression and viral replication. The HIV genome contains multiple splice sites, resulting in alternatively spliced mRNAs. They can be divided into three groups: unspliced, singly

spliced, and doubly spliced (Figure 1.2). In brief, the proviral DNA is first transcribed into a primary viral RNA transcript, which is about 9 kb in length and serves as both genomic RNA for progeny virus and the unspliced viral mRNA encoding viral Gag and Gag-Pol proteins. Alternative retention or splicing of the primary transcript and nuclear export of the unspliced transcript are well regulated. During the early phase of viral replication, two doubly spliced viral transcripts are produced and exported into cytoplasm, where they are translated into the regulatory and accessory proteins Tat, Rev and Nef. These viral proteins regulate subsequent viral gene expression and cellular function. As the newly generated Rev proteins accumulate in the nucleus, they bind to the transcribed viral mRNA and facilitate the export of singly spliced and unspliced mRNAs, which encode the viral structural proteins Env and Gag. The full-length unspliced viral genomic RNA binds to Gag polyproteins which traffic to the cell membrane for viral assembly. Among the numerous cellular factors reported to be involved in HIV assembly and budding, the ESCRT cellular machinery (Endosomal Sorting Complex Required for Transport), which was initially discovered to be important for the biogenesis of multivesicular bodies (MVB), recruits the Gag p6 domain and plays a key role in formation and release of new particles. It is currently thought that the resulting late-budding structure is released by the cellular ESCRT machinery severing the membrane tether connecting it to the producer cell ^{13, 104}.

Figure 1.2

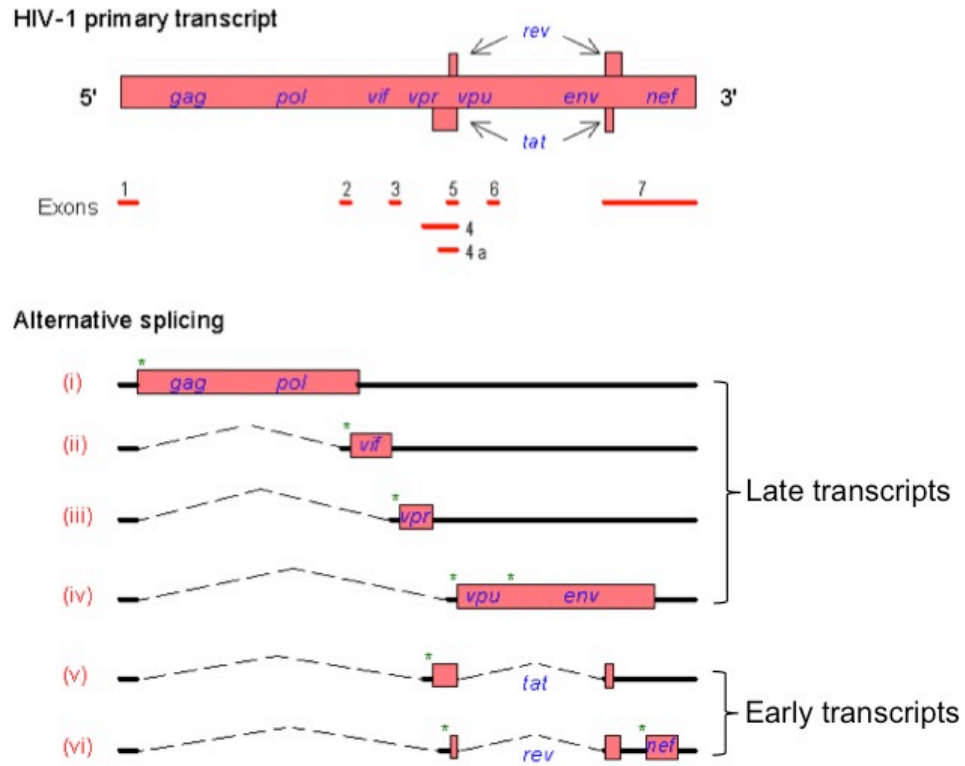


Figure 1.2. Schematic illustration of alternative splicing of the HIV-1 primary transcript. (i) is unspliced, (ii) to (iv) are singly spliced, (v) and (vi) are doubly spliced. The resulting mRNA (i), (iv) and (vi) are bicistronic. The star "*" indicates the location of the initiation codon (AUG). (From Web-book: <http://www.web-books.com/>)

Clinical course of HIV infection

The clinical course of HIV infection generally includes three phases or stages: 1) acute infection; 2) chronic or latent infection; and 3) AIDS-defining illness. Such a course of infection is characteristic of the so-called typical progressor who represents the majority of HIV-infected individuals (Figure 1.3). Three major clinical parameters are

associated with HIV disease progression: viral load, immune activation, and peripheral CD4⁺ T cell number. In acute infection, rapid viral replication occurs leading to acute viremia, followed by a massive depletion of CD4⁺ T cells^{33, 124}. Activation of CD8⁺ T cells is then detected and believed to significantly control acute viral replication¹¹⁶. The disease transition from primary infection to the clinical latent stage is characterized by downregulation of viremia and resolution of symptoms, if present. A set point for viral load is usually formed in this stage. HIV disease is still active with persistent replication of virus and progressive loss of CD4⁺ T cells. Finally, when CD4⁺ T cell numbers decline below a critical level cellular immunity drops to a point leading to infections with a variety of opportunistic pathogens defined as AIDS^{51, 117}.

Figure 1.3

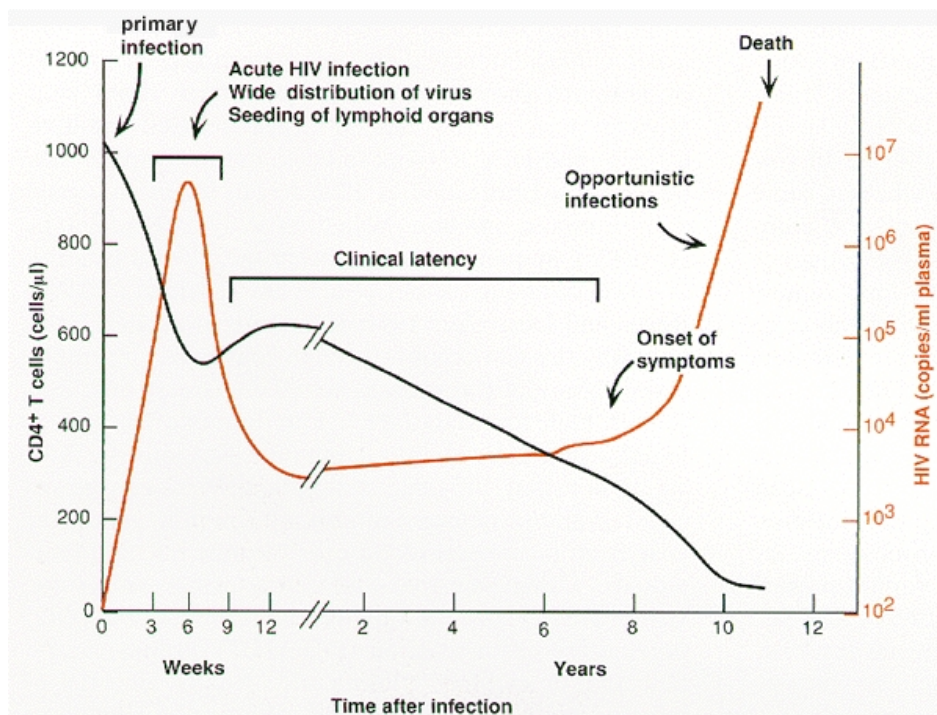


Figure 1.3. Natural course of HIV infection. The diagram demonstrates the relationship between viral load and peripheral CD4⁺ T cell count.

Impairment of CD4⁺ T cell immunity in HIV

CD4⁺ T cells play a central role in maintaining effective cellular and humoral immune responses by providing help to CD8⁺ T cells, B cells and innate effectors²². In HIV infection, the number of circulating CD4⁺ T cells is commonly used as a measure of global “immune competence” and provides a predictor for the immediate risk of opportunistic infections¹⁰¹. The importance of cellular immunity in HIV infection has been identified in many studies, even though the specific protective immune correlates have not been fully elucidated. Enhanced cellular immunity towards HIV has been linked with better control of viral replication in patients who show slow or delayed disease progression^{67, 106, 118}.

However, one of the key features for HIV infection is the profound impairment of CD4⁺ T cell immunity, attributed both to the massive loss of CD4⁺ T cells mediated by direct viral infection during early infection and the subsequent progressive loss of CD4⁺ T cells during the chronic stage^{47, 125}. In addition to cell depletion, functional abnormalities of CD4⁺ T cells also contribute significantly to the impaired CD4⁺ T cell immunity in HIV. T cells from HIV-infected individuals showed reduced proliferation following stimulation with antigens or mitogens *in vitro*³⁴. Diminished expression of IL-2

is readily demonstrated in these cells and may be related to the proliferation defects¹³⁹. Furthermore, a sequential loss of immune responsiveness to recall antigens, followed by alloantigens and then mitogens has been described for CD4⁺ T cells from HIV-infected individuals³⁴. Further studies using TCR stimulation to characterize the proliferation defects in CD4⁺ T cells indicate that such defects are associated with early-phase cell-cycle arrest¹⁴⁰ and are commonly observed in patients with sustained CD4⁺ T cell loss¹⁴¹. Since CD4⁺ T cells play a key role in regulating antibody responses, CTL responses and other innate effectors, the loss of these cells constitutes a major impairment in immune capacity in HIV infection.

HIV Envelope protein and inhibition of CD4⁺ T cell response

As described above, HIV infection causes profound CD4⁺ T cell impairment leading to immune suppression and the development of AIDS (49). Mechanisms have been suggested for this including direct cytopathic effects of HIV, accumulation of unintegrated viral DNA or toxic viral proteins, virus-induced changes in target cell membrane permeability, and apoptosis of HIV-infected CD4⁺ T cells (50). Other proposed mechanisms include lysis of CD4⁺ T cells by cytolytic T cells and tryptophan depletion by indolamine 2,3-dioxygenase (51, 52). However, in many patients only a small fraction of CD4⁺ T cells appear to be infected and the degree of cell loss greatly exceeds the number of infected cells, indicating that virus-induced bystander dysfunction plays a role in CD4⁺ T cell deficiency.

Among the HIV proteins, Env, expressed on the surface of free virions and infected cells, is a potent immune inhibitory molecule and has been demonstrated to play a predominant role in suppressing CD4⁺ T cells by multiple mechanisms (61, 62, 63). CD4 is a transmembrane glycoprotein that plays an important role in T helper cell function by interacting with MHC class II during T cell activation⁷³. However, it has been shown that cross-ligation of CD4 molecules prior to TCR stimulation primes the cells for apoptosis or a non-responsiveness state upon TCR activation^{146,63}. Env is a trimeric protein with each of its monomers possessing a CD4 and co-receptor binding site, and it has been demonstrated to produce a similar effect on CD4 ligation. Env multimers created in vitro with antibodies against monomeric gp120 subunits can mediate cross-linking of CD4 and/or co-receptor leading to signaling that causes aberrant activation of CD4⁺ T cells^{10, 132}. These observations from in vitro culture systems are consistent with the studies of HIV-infected patients demonstrating that similar interactions between Env on infected cells or virions can cause apoptosis of uninfected bystander CD4⁺ T cells, which occurs primarily in lymphoid tissues where antigen presentation and T cell activation occur⁵⁵. A proposed hypothesis is that when bystander, uninfected HIV-specific CD4⁺ T cells have CD4 pre-ligated by Env, they undergo apoptosis after they are activated via TCR stimulation by the cognate antigen presenting cells (APCs)³⁷. For Env-induced apoptosis of bystander CD4⁺ T cells, multiple apoptotic pathways have been proposed. Ligation of CD4 receptor by Env prior to TCR stimulation leads to upregulation of Fas in CD4⁺ T cells, and the subsequent expression of Fas-L upon TCR activation thereby renders the T cells susceptible to

apoptosis by Fas-FasL pathway¹⁴⁶. Fas independent apoptosis and other death pathways, including tumor necrosis factor (TNF)-TRAIL pathway, have also been reported as mechanisms for Env-induced apoptosis^{15, 58, 77, 113, 127}. In addition to apoptosis, lack of proliferation¹³² and induction of IL-2 reversible anergy⁹⁹ have also been suggested as forms of Env-induced inhibition of CD4⁺ T cells.

Although inhibition of CD4⁺ T cells by Env has been well established, the underlying mechanisms in terms of cellular signaling defects and the receptors utilization are not fully understood. CD4 binding has been shown to play a major role in mediating suppression in most studies^{3, 19, 82}. Treatment of CCR5Delta32 homozygote PBMCs with a CCR5-specific envelope induced apoptosis in T cells, demonstrating that CD4 could mediate suppression in the absence of co-receptor⁷. In contrast, involvement of co-receptor binding in Env-induced inhibition of CD4⁺ T cells was also reported. Stimulation of CCR5 by R5 Env triggers apoptosis of CD4⁺ T cells⁴. Another study showed that bystander apoptosis of CD4⁺ T cells required direct contact with virions and Env/CXCR4 binding⁷¹.

As previously mentioned, Env could deliver aberrant signals to CD4⁺ T cells through interaction with CD4 and/or coreceptors. Therefore, it is likely that T cell signaling cascades (TCR signaling and cytokine signaling) could be affected by Env as the molecular mechanisms for CD4⁺ T cell inhibition. Env-induced defects in JAK/STAT signaling pathways have been reported in multiple studies^{89, 135}. It was demonstrated that

exposure of CD4⁺ T cells to Env caused a failure of normal JAK-1 and JAK-3 activation and the unresponsiveness of STAT5a and STAT5b to IL-2⁸⁹. In addition, many studies have demonstrated defects in TCR signaling induced by Env. Binding activities of important transcription factors to target promoters, including NFAT, NFκB and AP-1, were observed to be significantly decreased in T cells with exposure to Env⁷⁵. Pre-exposure of a T cell line to gp120 led to the inhibition of Lck activation in response to TCR/CD3 stimulation¹⁰⁹. In a study with primary human CD4⁺ T cells, pre-treatment of these cells with gp160 resulted in marked inhibition of tyrosine phosphorylation of p59 (Fyn), PLC-gamma1 and ras activation, providing insight into the mechanism by which the Env may alter TCR/CD3-activation-induced signal transduction¹⁴⁵.

Collectively, the interference of TCR or cytokine signaling by Env is complex and the biochemical mechanisms are not fully understood. Env protein is a key component of an effective vaccine because it exclusively induces neutralizing antibodies⁷². Accordingly, understanding the mechanism for Env-induced inhibition is critical for designing effective Env-based HIV vaccine candidates and approaches.

T regulatory cells (Treg) and HIV

The concept of suppressor T cells was first described by Gerson et al when they suggested that a population of T cells could suppress immune responses⁶¹. Later on CD25 was identified as a surface marker for a population of suppressor T cells that could control autoimmune responses in mice, and these cells were termed regulatory T cells

(Treg). The discovery of the transcription factor FoxP3, which is highly expressed in both mouse and human Treg cells and is required for Treg cell function and development, has led to intensive studies of Treg biology in mouse models as well as for more reliable identification of Tregs in human ^{21, 29, 56, 95, 166}.

Several subclasses of Treg cells have been identified. Tregs expressing FoxP3, CD4 and high levels of CD25 are termed natural regulatory T cells and are thought to primarily originate in the thymus, although studies have shown that these cells can be generated in the periphery from non-Treg cells under certain conditions ¹⁸. The other type of Treg cell is known as T regulatory 1 cells (Tr1) that express little or no FoxP3 but preferentially produce large amounts of IL-10 ⁷⁴. The TGF- β -producing intraepithelial T cells in gut associated lymphoid tissues (GALT) are known as T helper 3 (Th3) with suppressive function ¹⁶⁰. The current knowledge on Treg classification, phenotype and development is still limited and more research is needed. Treg cells can exert suppression by multiple mechanisms, although much remains to be elucidated ^{156, 172}. Treg cells could induce suppression by direct cell-cell contact or soluble factors like IL-10 and/or TGF- β ^{31, 137}. Whether Treg cells release IL-10 and thus exert contact-independent suppression is not clear, but it should be noted that IL-10 is primarily found in adaptive regulatory T cells, such as Tr1 cells, and it is indicated to play an important role in mediating suppression for these cells ¹⁵⁶.

Treg cells are traditionally believed to protect the host from autoimmune reactions by down-regulating autoimmune responses. However, there is accumulating evidence supporting an active role for Tregs in modulating immune responses directed to infectious agents, including HIV². Treg cells isolated from the peripheral blood and lymphoid tissues of HIV-infected patients have been demonstrated to maintain a suppressive activity on HIV-specific CD4⁺ and CD8⁺ T cell response^{85, 86, 161}. Both increased and decreased numbers of Treg cells have been reported in HIV infection, and a number of factors could influence the result, such as the tissues studied, the stage of infection, and the institution of ART. In addition, it becomes more complicated by the use of different Treg detection methods by different groups^{5, 6, 49, 52, 85-87, 108, 112, 115, 149, 161}. In studies using peripheral blood, both increased^{1, 85, 161} and reduced^{49, 115} frequencies of CD4⁺CD25⁺ Treg cells were reported, leading to two converse hypotheses that HIV drives expansion of Treg cells or HIV induces a selective loss of Treg cells. Subsequent studies using LN or gastrointestinal mucosal tissue demonstrated an increased frequency of suppressive Treg cells in HIV-infected patients, as detected by FoxP3 expression, supporting the hypothesis that Treg cells, rather than undergoing a selective loss induced by virus, accumulate at sites of viral expression, where they can exert suppression on Ag-specific T cell activation^{6, 52, 112}. How HIV regulates the frequency and function of Treg cells during infection and how this regulation relates to HIV pathogenesis and host immunity, remains highly controversial. There is limited evidence, so far, demonstrating the direct interaction of Treg cells and HIV or its gene products. Human Treg isolated from healthy donors express the HIV coreceptor CCR5 and were reported to be the target

of HIV infection with high susceptibility¹¹⁵. In contrast, it has been demonstrated that HIV increases Treg viability through Env-CD4 interaction, resulting in accumulation of Treg cells¹¹². Collectively, the regulation of Treg frequency and function by HIV and the physiological relevance in the context of HIV infection or vaccination are not fully understood.

Immunization and immune dysfunction in chronic HIV/SIV infection during ART

The introduction of anti-retroviral therapy (ART) results in delayed disease progression and a remarkable decrease in AIDS related deaths. However, the current regimens for HIV treatment have major limitations. First, it fails to eradicate the virus. It has also been demonstrated that even during the most potent regimens of retroviral suppression, viral production continues, as detected by some ultrasensitive assays^{43, 70, 122}. In addition, immune recovery subsequent to ART often appears to be partial and limited to certain T cell responses, and does not include HIV-specific CD4⁺ T cell responses^{8, 83, 129}.

Cell-mediated immunity is involved in controlling HIV replication and correlates with better control of disease progression, which provides a rationale for active immunization of chronically infected individuals to augment the existing T cell responses. A number of approaches attempting to stimulate cell-mediated responses by immunization have been studied in SIV-infected rhesus macaques during ART with mixed results reported regarding immunologic and virologic benefits^{42, 57, 69, 96, 150, 157, 173}.

Rhesus macaques with long-standing SIVmac251 infection were immunized three times with a poxvirus vector expressing the SIVmac structural and regulatory genes during ART. Following antiretroviral treatment interruption, the viral set point was significantly lower in immunized macaques than in control macaques ¹⁴⁸. A strategy using autologous dendritic cells (DC) pulsed either with viral peptides or chemically inactivated virus, which could elicit strong immune control of the virus, has been tested in the SIV macaque model and demonstrated promising result ^{42,96}. Immunizations with DNA plasmids encoding multiple SIV antigens also offered long lasting immunologic and virologic benefits to SIVmac251 infected and ART treated macaques ¹⁵⁷. In contrast to these promising results, very limited immunologic and virologic benefits by immunization in ART-treated, SIV infected macaques have also been reported. A study using the SIVmac239 model demonstrated that immunization with SIVmac239-derived Env, Gag/Pol, and Tat/Rev/Nef plasmids by in vivo electroporation elicited T-cell immune responses, but no control of SIVmac293 viremia ¹⁷³. Collectively, using various immunization strategies including DNA, poxvirus vectors, and antigen-pulsed DCs, SIV-specific immune responses were evoked in SIV-infected rhesus macaques during ART. The virologic benefits of these immunizations after release of macaques from ART were variable from no control to temporal control or long-lasting control.

However, similar immunization (either vaccine or treatment interruption) for chronically HIV-infected, ART-treated humans has been disappointing, as no significant HIV-specific T cell responses were induced by immunization in this setting ⁶⁶, which is

consistent with the observation that treatment with effective ART results in regained T cell responses to common antigens, but poor responses to HIV-specific antigens, suggesting the long-lived defects in the generation of HIV-specific CD4⁺ T cell immunity in HIV^{8, 83, 129}. The mechanisms are not fully characterized or understood. Compared to peripheral blood, which has been commonly used to study the antigen-specific T cell responses to HIV/SIV infection or immunization, the secondary lymphoid tissues represent an important site where antigen presentation, antigen-specific T cell response and the viral replication occur. APCs, primarily DCs, function by obtaining antigens in the periphery, migrating to draining lymph nodes (LNs), and interacting with antigen-specific T cells, leading to the T cell activation and antigen-specific responses. However, in the context of HIV infection, the interaction between viral antigen-expressing APC and HIV-specific T cells could blunt the immune response through multiple mechanisms^{65, 168}. It has been suggested that DC-T cell interaction can stimulate viral replication in DC, CD4⁺ T cells and the cell conjugates, which represent an explosive site of HIV infection that could escape from ART¹²⁸. Multiple mechanisms have been proposed for this suppressed T cell activation induced by the inappropriate deleterious DC-T interactions in LN during HIV infection, including T cell death secondary to activation of latent infection, aberrant signals from DCs expressing inhibitory HIV proteins, and CD4⁺ T cell fusion with infected DCs or DCs expressing Env. Collectively, the mechanisms for the impaired antigen-specific T cell response in chronic HIV are complex and not fully understood.

Aims of thesis

In this thesis, I have examined the mechanisms for the inhibition of CD4⁺ T cell immunity in HIV. An in vitro system modeling the interaction between HIV Env and human primary CD4⁺ T cells will be used to study the inhibitory effect of Env protein, when delivered as a vaccine on APC and non-APC, or as free virions, on antigen-stimulated bystander CD4⁺ T cell proliferation. I investigate the mechanisms by studying various forms of T cell inhibition, including anergy, apoptosis and cell death. Some important TCR and cytokine signaling pathways are also investigated in CD4⁺ cells with Env exposure. Considering the suppressive effect of human Treg cells on autologous CD4⁺ and CD8⁺ T cell proliferation and its ability to interact with HIV Env, I study the regulation of Treg frequency, activation phenotype, and suppressive activity in human peripheral blood by Env, and the functional involvement of Treg cells in Env-induced inhibition.

In the third chapter of this thesis, I move the studies to an in vivo model. Using chronically SIV and CMV co-infected, ART-treated rhesus macaques, I study the early immunologic and virologic consequences of immunization with SIV and CMV antigen in draining lymph nodes, where antigen presentation, antigen-specific T cell responses and viral replication occur during infection. The systemic response in peripheral blood after immunization is also studied to track the fate of immunization-induced antigen-specific T cells.

CHAPTER 2

HIV Envelope Suppresses CD4⁺ T Cell Activation Independent of T Regulatory Cells

Introduction

One of the hallmarks of HIV-associated defects in cell-mediated immunity is the progressive loss of CD4⁺ T cell responses, even before a significant decrease in peripheral blood CD4⁺ T cell numbers occurs¹⁰⁵. The Env protein of the virus is a key component of an effective vaccine because it exclusively induces neutralizing antibodies, which are protective in the macaque model¹⁰⁰. However, Env has been demonstrated to suppress CD4⁺ T cells by multiple mechanisms. Inhibition has been characterized by a lack of proliferation^{82, 132}, the induction of anergy^{93, 99}, and various apoptotic mechanisms^{10, 165}. The role of Env interactions with CD4 and coreceptors in mediating the various forms of suppression is under debate with studies demonstrating a role for each^{7, 15, 154}. R5 Env caused apoptosis of CD4⁺ T cells from a CCR5Δ32 homozygous donor, whereas, in another study, Env interaction with CXCR4 and a nonsignaling CD4 induced CD4⁺ T cell apoptosis¹³⁰. The mechanism of suppression observed and the receptors involved varied according to the nature of Env used and the cell culture system studied. Env is a trimeric protein with each of its monomers possessing a CD4 binding site. Thus, an Env trimer¹³² or aggregates created by antibodies against monomeric gp120 subunits¹⁰ can mediate cross-linking of CD4 and/or coreceptor leading to signaling. In patients infected with HIV, trimeric Env capable of suppression is found on

the surface of infected cells as well as on the virus itself. The use of Env in HIV vaccines, especially in commonly used DNA and viral vectors, could be envisioned to deliver similar CD4⁺ T cell suppressive signals.

Although APCs such as DCs are prime targets for vaccines, other cell types at the site of vaccine delivery, and potentially the major vaccine antigen expressing cell types, are non-APCs such as myocytes, fibroblasts, subepithelial, and epithelial cells for muscle, dermal, subcutaneous, and mucosal delivery, respectively, as well as cells in the draining lymph nodes^{30, 38}. CD4⁺ T cells that are recruited to the site of vaccine delivery as well as to the draining lymph node in response to vaccine-induced inflammation may then interact with and be suppressed by interaction with Env expressed on the surface of all of these cells.

Naturally occurring regulatory T cells (Treg) are a subset of CD4⁺ T cells co-expressing CD25 and FoxP3 and have the ability to suppress autologous CD4⁺ and CD8⁺ T cell activation and proliferation. The role of Treg cells in HIV has been intensively studied and there are accumulating evidence supporting the active engagement of Treg cells in modulating immune response towards HIV infection^{1, 86, 112, 161}. Issues with regard to whether and how HIV regulates the frequency and function of Treg cells during infection, and how this regulation relates to HIV pathogenesis and host immunity, remains highly controversial. There is limited evidence, so far, demonstrating the direct interaction of Treg cells and HIV or its gene products. One study showed that HIV

interacts directly with Treg cells through Env-CD4 signaling and increases Treg frequency in PBMC through selectively enhancing Treg viability and the subsequent accumulation of Treg cells ¹¹².

In this chapter, using an in vitro system that models the in vivo interaction between CD4⁺ T cells and Env, we investigated the mechanisms for how Env suppresses CD4⁺ T cells concentrating on signaling pathways and the involvement of Tregs.

Materials and Methods

Cells and HIV-1

Human embryonic kidney 293T cells (American Type Culture Collection) were maintained in DMEM (Invitrogen Life Technologies) supplemented with 10% FCS (HyClone) and glutamine (Invitrogen Life Technologies). Monocyte-derived dendritic cells were obtained from purified monocytes (Center for AIDS Research (CFAR) Immunology core) and plated in 6-well plates at 3×10^6 cells in 3 mls of AIM V serum-free media (Life Technologies, Rockville, MD), supplemented with GM-CSF (50 ng/ml) and IL-4 (100 ng/ml). After 3 days, cells were fed by adding 1.5 mls of Aim V supplemented with GM-CSF and IL-4. Peripheral blood mononuclear cells (PBMC) from HIV-negative donors were obtained through an IRB-approved protocol and maintained in complete RPMI 1640 (Invitrogen Life Technologies) containing 10% FCS and glutamine. Aldrothiol-2 (AT-2) inactivated ¹³¹ HIV-1 NL4-3 virions were obtained from University of Pennsylvania CFAR virology core.

Plasmid constructs and RNA synthesis

HIV Envs IIIB, YU2, and 89.6 were cloned into the RNA expression plasmid pT7TS that contains a T7 promoter, 5' and 3' UTR sequences of Xenopus-globin, and a 30-nuc long poly (A) tail⁸⁰. Plasmids were linearized with restriction enzymes and in vitro transcribed with T7 polymerase (mMessage T7 Ultra Kits (Ambion)), extended poly-A tail was added. RNA was maintained in nuclease-free water at a concentration of 1 µg/µl and stored at -20°C.

DC infection with vaccinia virus

DCs were suspended at a concentration of 2 million/mL and infected with vaccinia-encoding IIIB, MN, or 92UG037.8 Env (NIH AIDS Research and Reference Reagent Program), T7 Polymerase (VTF7.3), or wild-type WR strain for 2 hours, washed, and incubated for 4 hours. Infected DCs were cocultured with 5- and 6-carboxyfluorescein diacetate, succinimidyl ester (CFSE)-labeled PBMCs at a ratio of 1:10 and activated with 0.1 µg/mL toxic shock syndrome toxin-1 (TSST-1) superantigen.

Exposure of CD4⁺ T cells to Env and CD4⁺ T cell proliferation

Transcribed mRNA encoding HIV Env was complexed to lipofectin for delivery to cells as described previously⁸⁰. 293T cells seeded in 12-well plates, approximately 90% confluent (5×10^5 cells), were transfected with Env (IIIB, YU2, and 89.6) encoding mRNA (5 µg/ml) or control poly(AC) RNA (Sigma-Aldrich), incubated for 4 h to allow

protein expression, and cocultured with 2×10^6 CFSE loaded PBMC in the presence of stimulation with soluble anti-CD3 mAb (1 $\mu\text{g/ml}$; Ortho Biotech) or 0.1 $\mu\text{g/ml}$ superantigen TSST-1 (Sigma) and IL-2 (20 U/ml; AIDS Reference and Reagent Program) with or without soluble CD4 (sCD4) (10 $\mu\text{g/ml}$; AIDS Reference and Reagent Program). Increasing or decreasing the ratio of Env-expressing 293T cells to PBMC did not affect suppression of CD4^+ T cell proliferation. For exposure to free virions, 2×10^6 CFSE-labeled PBMC were incubated with AT-2-inactivated HIV-1 NL4-3 virions at concentrations of 1, 0.3, or 0.1 μg HIV p24 Gag equivalents/ml, which corresponds to a gp120 concentration of 1, 0.3, and 0.1 nM, respectively ¹¹². Four days after Env exposure, PBMC were collected, washed with PBS containing 1% FCS, and stained with anti-CD4-allophycocyanin (RPA-T4; BD Pharmingen). Cells were analyzed on a FACS-Calibur flow cytometer, and proliferation of CD4^+ T cells was analyzed by CFSE dilution using Cellquest software (BD Pharmingen). A minimum of 10,000 events were analyzed. The percent of stimulated CD4^+ T cells that proliferated was calculated as described ¹⁶⁴.

Detection of apoptosis and activation in CD4^+ T cells exposed to Env

PBMCs were cocultured with Env-expressing 293T cells or 700 μM cycloheximide (Sigma) and activated with 0.1 $\mu\text{g/ml}$ TSST-1 or 1 $\mu\text{g/ml}$ anti-CD3 mAb. Apoptosis was examined 6, 17, and 36 hours after coculture and identified as annexin V-positive (BD Pharmingen, San Diego, CA) and 7-AAD-negative (Sigma) cells. Total cell death was measured by positive staining for 7-AAD in ungated cells. For studies measuring up-regulation of activation markers in Env-treated PBMCs, CD69

peridinin chlorophyll protein (day 1), HLA-DR fluorescein isothiocyanate and CD25 phycoerythrin cyanine 7 (BD Immunocytometry Systems, San Jose, CA) (day 2) were used.

Detection of phosphorylation of signaling molecules in CD4⁺ T cells

Purified CD4⁺ T cells were exposed to IIIB Env-expressing 293T cells for 1 day, stimulated with 1 µg/mL anti-CD3 antibody for 10 minutes for p38 or 2 days followed by 20 U/mL IL-2 for 2 to 20 minutes for STAT5a, fixed, and analyzed by flow cytometry using anti-phospho-MAPK (pT180/ pY182) Alexa Fluor 647 or anti-phospho-STAT5 (pY694) Alexa Fluor 647 (BD Pharmingen), respectively. For AKT phosphorylation, PBMCs were cocultured with IIIB Env-expressing 293T cells for 2 hours in complete DMEM and then for 18 hours in DMEM with 0.5% serum, stimulated with 1 µg/mL anti-CD3 antibody for 10 minutes, fixed, and stained with anti-phospho-AKT (ser 473) Alexa Fluor 488 (Cell Signaling Technologies, Danvers, MA) and anti-CD4 v4 (clone L120) phycoerythrin (BD Immunocytometry Systems).

CD25⁺ cell depletion and T cell proliferation assay

CD25⁺ cells were depleted from PBMC using magnetic beads (Dynabeads CD25; Invitrogen Life Technologies). Briefly, PBMC and magnetic beads were washed twice with PBS containing 1% FBS. Then 1 x 10⁷ cells/ml were mixed with a 40:1 ratio of beads to CD25⁺ cells assuming 5% of cells expressed CD25, and rotated at 4°C for 45 minutes. A second round of depletion was performed with a decreased volume of beads

(half of that used in the first round) to achieve near complete depletion of CD25⁺ cells.

Whole and CD25-depleted PBMC were CFSE labeled, and cultured with Env-expressing and control 293T cells in presence of TCR stimulation.

Treg sorting and suppression culture

Whole PBMC or CD25-depleted PBMCs were cultured with Env-expressing or control 293T cells. During Env exposure, whole PBMC were unstimulated, whereas CD25-depleted PBMC were stimulated with anti-CD3 and IL-2 to expand induced Tregs. A total of 48 h after Env exposure, PBMC were removed from 293T monolayers and stained with anti-CD4-FITC, anti-CD25-PE-Cy5, and anti-CD127-PE (BD PharMingen). CD4⁺, CD127⁻, CD25⁺ cell sorting was performed on a FACS/Aria (BD bioscience) to obtain Tregs. A total of 2 x 10⁵ CFSE labeled CD25-depleted autologous PBMC responders were mixed with increasing amounts of FACS sorted CD4⁺, CD127⁻, CD25⁺ Tregs (Suppressor to Responder ratios of 1:3 and 1:6) in the presence of stimulation with anti-CD3 and IL-2. Four days after coculture, the proliferation of responder cells was determined by flow cytometry.

Treg analyses

PBMC were exposed to Env (either on 293T as AT-2-treated virions) as described above. For exposure to 293T cells, PBMC were either resting or stimulated by anti-CD3 and IL-2. For incubation with AT-2 inactivated virions, resting PBMCs were used. At various time points after Env exposure (days 1, 3, and 5), PBMC were collected, washed

with PBS containing 1% FCS, and stained with anti-human CD4-FITC (SK3; BD Pharmingen) and anti-CD25-PE-Cy5 (M-A251; BD Pharmingen). Cells were then washed, fixed with 2% paraformaldehyde (eBioscience), permeabilized with PBS 1% FCS containing 0.1% saponin (eBioscience), incubated with normal rat serum at 4°C for 15 min, and stained with anti-human FoxP3-allophycocyanin (PCH101; eBioscience) on ice for 30 min according to the manufacturer's instructions. The appropriate isotype-matched Abs were used to define background staining. Expression of CD4, CD25, and FoxP3 in cells were determined by flow cytometry. A minimum of 10,000 events were analyzed.

IL-10 and TGF- β ELISA

IL-10 and TGF- β concentrations in culture supernatants were determined by commercially available ELISA kits (BD OptEIA) according to the manufacturer's instructions.

Statistics

Mean, standard error of the mean (SEM), and student's t-test were performed using Microsoft Excel software.

Results

Env expressed on 293T cells induces suppression of CD4⁺ T cell proliferation

Following vaccine delivery, a number of cell types other than APCs take up and express vaccine-derived antigens. Expression by these cells presents Env in the absence of costimulatory molecules. 293T cells were transfected with Env-encoding mRNA as consistent, and low-level protein expression is obtained in most cells¹⁶³, allowing uniform exposure of CD4⁺ T cells to Env. PBMCs were used as a source of CD4⁺ T cells and cocultured with 293T cells expressing X4-tropic IIIB, dual tropic 89.6, and M-tropic JR-F1, YU-2, and SF162 Envs. Cells were stimulated with a concentration of anti-CD3 mAb that typically stimulates one quarter of the CD4⁺ T cells. To monitor proliferation, PBMCs were stained with CFSE⁹⁷. CD4⁺ T cells that encountered any of the Envs had lower proliferation levels compared with cultures exposed to control RNA-transfected 293T cells (Figure 2.1A). Similar inhibition was observed when cells were activated by the superantigen TSST-1, a model antigen for activation via the T-cell receptor (TCR), which requires the same costimulatory and adhesion molecules as conventional antigens¹³⁸. This suggested that suppression of proliferation was not dependent on a particular mode of activation. In addition, negatively selected purified CD4⁺ T cells exposed to Env also failed to proliferate, arguing against a role for CD8⁺ T cells and natural killer cells in mediating suppression (Figure 2.1B).

To characterize the signal delivered by Env, we examined whether suppression could be overcome by blocking Env interaction with CD4. sCD4 almost completely

abrogated Env-induced suppression (Figure 2.1A). P-values were calculated using results from 3 independent experiments. A paired Student t-test with 2 tails was performed using the percentage of change from control across experiments (no Env) or sCD4 treatment and is shown in Table 1.

Expression of functional trimeric Env on 293T cells was confirmed in fusion assays demonstrating that mRNA delivery resulted in surface expression of fusion-competent Env (Figure 2.1C-D). Delivery of Env-containing vaccines in vivo results in a variety of expression levels that are governed by the cellular vaccine target and the vaccine delivery method. In our experimental system that mimics vaccination with Env, we chose delivery of Env by encoding mRNA to 293T cells, which results in expression of Env in nearly 100% of cells at lower levels compared with other methods of delivery, such as DNA. This was important as lymphocyte trafficking does not occur in our system. Thus, if only 25% of the cells expressed Env, then only about 25% of the CD4⁺ T cells added would be exposed. This low level of expression was sufficient to mediate suppression, and, in fact, a 5-fold reduction in the quantity of Env encoding mRNA delivered, which results in a 5-fold lower level of protein expressed, resulted in similar levels of suppression (data not shown). Thus, at the range of expression levels seen in our culture system, large variations in the amount of Env expressed did not alter the degree of suppression exerted. This could explain the lack of correlation observed between the degree of suppression mediated by an Env strain (Figure 2.1A) and the amount of fusion mediated (Figure 2.1C-D).

Several reports have found that trimeric or aggregated forms of Env are required to mediate suppression^{10,93}. However, others have found that monomeric gp120 could exert suppressive effects^{82,165}. In our system, CD4⁺ T cells were exposed to trimeric cell-surface expressed Env as well as monomeric gp120 released into the supernatant, likely from dissociation of cell-surface gp160. PBMCs exposed exclusively to exogenous monomeric gp120 derived from IIIB or SF162 failed to suppress CD4⁺ T cell proliferation (Table 2). The added gp120 retained the ability to block binding of CD4 mAb (leu-3a) demonstrating functional binding to CD4. This confirmed that trimeric surface expressed Env-mediated suppression in our system. Thus, transfection of 293T cells with encoding mRNA resulted in surface expression of functional Env, which mediated inhibition of antigen-specific CD4⁺ T cell proliferation in a CD4-dependent manner.

Figure 2.1. HIV Env suppresses anti-CD3-activated CD4+ T cell proliferation. (A)

293T cells expressing IIB, 89.6, YU-2, SF162, and JR-F1 Envs or no Env poly (A,C) RNA transfected) were cocultured with CFSE-labeled PBMCs with or without sCD4 (10 $\mu\text{g/ml}$). T cells were activated with anti-CD3 mAb (0.5 $\mu\text{g/mL}$), and monitored for proliferation 4 days after coculture by flow cytometry with staining with anti-CD4 mAb.

(B) Negatively selected purified CD4+ T cells were cocultured with 293T cells expressing IIB Env or no Env and activated using anti-CD3 mAb (0.5 $\mu\text{g/mL}$).

Quadrants are labeled with the percent of events in each. Fusion assays demonstrated surface expression of functional Env. Encoding mRNA-transfected 293T cells were mixed with QT6 cells expressing CD4 and CXCR4 (C) or CCR5 (D), and fusion of the 2 cell populations was measured by luciferase production. Experiments shown are representative of 3 independent experiments.

Table 1. Statistical analysis of suppression of CD4⁺ T cells proliferation by Env

Env strain	<i>P</i> of suppression compared with no Env control	<i>P</i> of suppression compared with sCD4 treatment
IIIB	< .005	< .005
89.6	.01	.02
JR-F1	.02	.04
Yu-2	.01	.02

P values were calculated using results from 3 independent experiments. Percentage of change from control (no Env) or sCD4 treatment was used in a paired Student *t* test with 2 tails.

Table 2. Effect of monomeric gp120 on CD4⁺ T-cell proliferation

	No Env	IIIB gp120	SF162 gp120
CD4 ⁺ cells undergoing proliferation, %	15.3	14.4	12.5
CD4 ⁻ cells undergoing proliferation, %	7.7	7.4	7.3

Anti-CD3 mAb (0.5 μg/mL) stimulated CFSE-labeled PBMCs were exposed to 2 μg/mL IIIB or SF162 gp120 or left untreated. CD4⁺ T cells were monitored for proliferation 4 days after coculture by flow cytometry and expressed as percentage of total CD4⁺ T cells.

DCs expressing Env suppress CD4⁺ T cell proliferation

Presentation of vaccine-derived antigens by APCs activates immune responses and is the basis of an effective vaccine. APCs, such as DCs in their mature form, express a variety of costimulatory molecules on their surface and provide a strong activation signal for T and B cells. Certain studies have shown that activation, including potent costimulation, could overcome suppression induced by CD4 cross-linking¹⁴⁶. Other studies have not supported this, demonstrating that HIV-infected DCs were able to suppress CD4⁺ T cells in an Env-dependent fashion. Currently, viral vector or nucleic

acid delivery of encoding antigen to DCs (either directly in vivo or after in vitro culture and loading of DCs) is in human trials and represents a potent vaccine approach^{62, 111}. We infected DCs with vaccinia virus encoding various Envs and tested their ability to affect CD4⁺ T cell proliferation in response to superantigen. DCs expressing the X4 Envs IIIB and MN and the R5 Env 92UG037.8 (a Clade A primary isolate) were found to inhibit CD4⁺ T cell proliferation to antigen, which was overcome by sCD4 (Figure 2.2). The p-values comparing IIIB Env to no Env or sCD4 treatment were less than 0.005. Expression of Env by these types of vectors in vitro has been previously demonstrated^{59, 76, 121}. Very low multiplicities of infection (MOIs) of vaccinia were used as T-cell toxicity was observed at MOIs typically used in fusion assays. However, even at low MOIs (0.3-0.6) a minority of the suppression could not be overcome with sCD4. This suppression of T-cell proliferation was also observed with 2 control viruses, vTF7-3 encoding T7 Polymerase and WR, the strain used to create all of the Env-expressing and control viruses, suggesting a vaccinia mediated suppression was also present. We concluded that vaccinia viruses had a small but significant level of toxicity in our system, which accounts for the inability of sCD4 to completely overcome the suppression of proliferation. In addition, the ability of Env to interact with CD4 molecules on DCs and suppress their maturation and function cannot be ruled out in our system, as it has been observed in other systems^{54, 167}. Expression of functional Env was confirmed in fusion assays between DCs and Quail cells (data not shown). Thus, mature DCs expressing Env mediate suppression of antigen-specific CD4⁺ T cell proliferation despite expressing multiple costimulatory molecules.

Figure 2.2

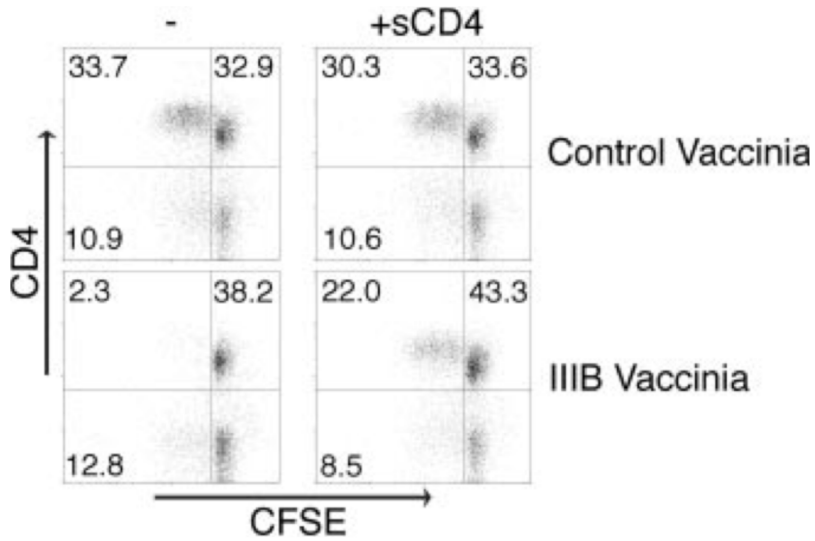


Figure 2.2. DCs expressing Env and presenting antigen suppress CD4⁺ T cell

proliferation. DCs were infected with vaccinia virus encoding IIIB Env or control vaccinia encoding T7 polymerase and cocultured with CFSE-labeled PBMCs with 0.1 $\mu\text{g}/\text{mL}$ TSST-1 with or without sCD4 (10 $\mu\text{g}/\text{ml}$). This concentration of superantigen requires presentation by DCs as no proliferation is observed in their absence.

Proliferation of CD4⁺ T cells was measured 5 days later by flow cytometry. Quadrants are labeled with the percent of events in each. Experiments shown are representative of 3 independent experiments.

Neither apoptosis nor IL-2-dependent anergy mediate CD4⁺ T cell suppression

Env-induced CD4⁺ T cell dysfunction has been described as anergy, apoptosis, cell death, or a failure to proliferate. We observed that Env suppresses CD4⁺ T cells in our system by hindering proliferation. We next investigated whether anergy or apoptosis

occurred. PBMCs were cocultured with Env-expressing 293T cells and were activated using TSST-1 superantigen. 6, 18, and 36 hours later, CD4⁺ T cells were analyzed for early apoptosis using annexin V staining of 7-AAD–negative cells. Cyclohexamide is a protein synthesis inhibitor that induces apoptosis of PBMCs and was used as a positive control. Env failed to induce apoptosis of CD4⁺ T cells above the level seen in cells unexposed to Env (Figure 2.3A). In addition, an increase in cell death (7-AAD uptake in ungated cells) in Env-exposed CD4⁺ T cells was not observed (Figure 2.3B). Thus, apoptosis and cell death are not major contributors to the suppression of proliferation induced by Env in our system. T-cell anergy is defined as a state of nonresponsiveness to activation with antigen. The addition of exogenous IL-2 overcomes most but not all types of anergy¹³³. Selliah and Finkel observed that CD4⁺ T cell suppression induced by Env could be overcome with exogenous IL-2¹³⁴. To investigate this mechanism, CFSE-stained PBMCs were cultured with control or Env-expressing 293T cells and activated with TSST-1 in the presence of increasing amounts of exogenous IL-2 (20, 40, or 80 U/mL). IL-2 was unable to prevent suppression of proliferation of CD4⁺ T cells by Env (Figure 2.3C, and data not shown), arguing against this form of anergy as the mechanism of suppression.

Figure 2.3

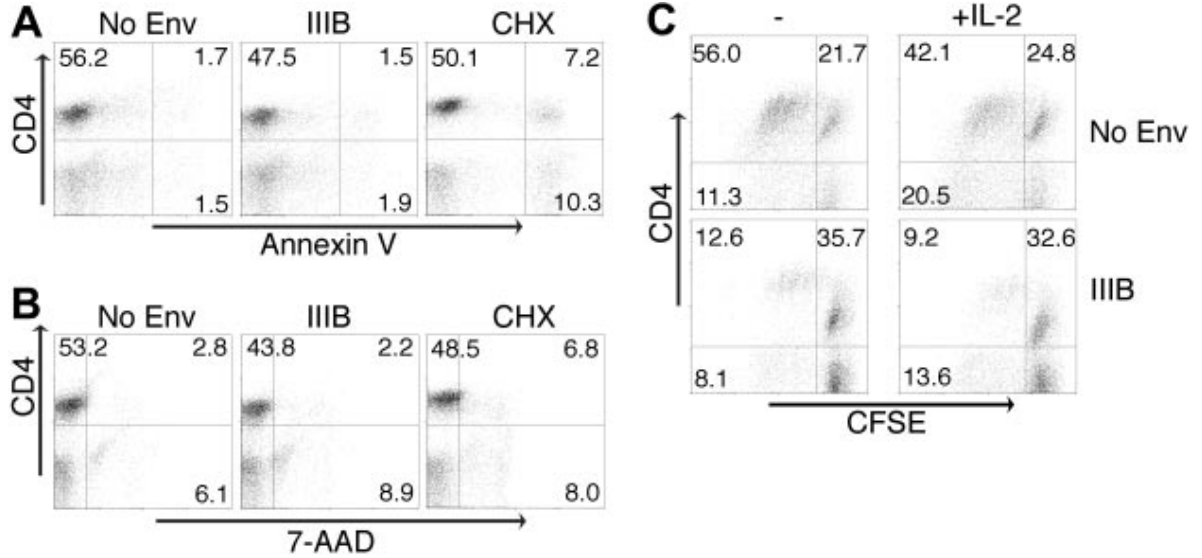


Figure 2.3. Neither apoptosis nor IL-2-dependent anergy mediate Env-induced

CD4⁺ T cell suppression. PBMCs were cocultured with IIIB Env or luciferase-

expressing 293T cells or 700 μ M cycloheximide (CHX) and activated with 0.1 μ g/mL

TSST-1. Thirty-six hours later, apoptotic CD4⁺ T cells were analyzed by flow cytometry

and defined as Annexin V positive and 7-AAD negative, early apoptotic cells (A) and 7-

AAD positive for dead cells (B). (C) CFSE-labeled PBMCs were cocultured with IIIB

Env-expressing 293T cells in the presence of 80 U/mL IL-2 and activated with 0.1 μ g/mL

TSST-1. CD4⁺ T cell proliferation was measured 5 days later by flow cytometry.

Quadrants labeled with percent of events in each. Experiments shown are representative

of 3 independent experiments.

CD4⁺ T cells activated in the presence of Env phosphorylate AKT, MAPK, and STAT5a but fail to completely up-regulate activation markers

CD4⁺ T cells exposed to Env failed to proliferate in response to antigen but did not apoptose, anergize, or die. We next sought to determine the location of the block in the activation cascade. In response to T cell activation, phosphorylation of key signaling molecules p38 MAPK and AKT relay the TCR signal to downstream mediators^{44, 133} CD4⁺ T cells exposed to Env and activated by anti-CD3 phosphorylated AKT and MAPK similar to control treated cells (Figure 2.4). IL-2 is an important cytokine involved in T-cell proliferation, and its binding to both the low and high affinity IL-2R leads to the phosphorylation of STAT5a⁷⁸. Exogenous IL-2 was unable to circumvent the suppression of proliferation induced by Env (Figure 2.3C), but cells exposed to and suppressed by Env were able to phosphorylate STAT5a in response to IL-2 (Figure 2.4B), suggesting that early signaling mediators in the IL-2 cascade also were not impaired. On activation, CD4⁺ T cells go through a series of events prior to proliferation, including up-regulation of markers of activation such as CD69, HLA-DR, and CD25³⁹. PBMCs were exposed to IIIB Env-expressing 293T cells and activated using anti-CD3 or superantigen. Up-regulation of CD69 (day 1) and HLA-DR and CD25 (day 2) was monitored using flow cytometry. CD4⁺ T cells exposed to Env were unable to completely up-regulate their activation markers (Table 3). Thus, although exposure of CD4⁺ T cells to Env did not suppress the phosphorylation of immediate signaling mediators, it suppressed the upregulation of early and late cell-surface markers of activation.

Figure 2.4

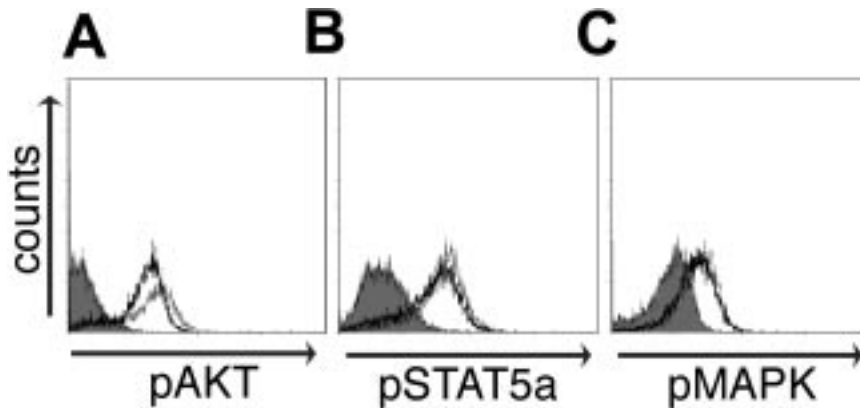


Figure 2.4. CD4⁺ T cells activated in the presence of Env phosphorylate AKT, p38 MAPK, and STAT5a. PBMCs were cocultured with Env or control RNA-expressing 293T cells for 24 hours. Phosphorylation of AKT or MAPK 10 minutes after anti-CD3 (1.0 μ g/mL) stimulation was measured using flow cytometry with gating on CD4⁺ T cells. Purified CD4⁺ T cells were exposed to control RNA or Env-expressing 293T cells and activated using anti-CD3 in the absence of IL-2. Two days later, cells were stimulated with 20 U/mL IL-2 and analyzed for STAT5a phosphorylation 7 minutes later. CD4⁺ T cells exposed to IIIB Env-transfected 293T cells and stained with isotype control mAb (shaded gray) or specific phosphoprotein mAb (thin line) or specific phosphoprotein mAb-stained control-transfected 293T cells (thick line) are shown for each signaling molecule. Experiments shown are representative of 3 independent experiments.

Table 3. Effect of HIV Env on activation marker expression

	No Env	IIIB Env
CD4⁺ cells expressing, %		
CD69	78.9	56.6
HLA-DR	23.5	12.9
CD25 (IL2R- α)	60.4	31.6
CD4⁻ cells expressing, %		
CD69	59.2	61.8
HLA-DR	43.7	41.6
CD25 (IL2R- α)	33.3	22.2

PBMCs were exposed to control or IIIB Env expressing 293T cells and activated with anti-CD3. Expression of CD69 on day 1 and HLA-DR and CD25 on day 2 on CD4⁺ and CD4⁻ T cells was determined using flow cytometry.

Env expressed on free virions suppresses CD4⁺ T cell proliferation

In HIV infection, Env present on infected cells and free virions can contact CD4⁺ T cells and induce dysfunction⁵⁵. Above, we demonstrate that Env expression on cells as occurs during vaccination with encoding vectors such as DNA or viruses induces an inhibition to antigen-specific CD4⁺ T cell proliferation in a CD4 dependent manner. This system used human PBMC as a source of CD4⁺ T cells that were cultured with Env expressing mRNA transfected 293T cells or vaccinia infected DC in the presence of either anti-CD3 or superantigen (TSST-1) stimulation. To test whether Env on virions is also able to induce suppression of anti-CD3 stimulated CD4⁺ T cell proliferation, we incubated PBMC with increasing amounts of AT-2 inactivated HIV-1 (NL4-3), which is infection deficient, but retains a functional (able to bind CD4) Env, modeling the noninfectious interactions between circulating free virions and CD4 in HIV infection¹¹². We found that inactivated virus could inhibit anti-CD3 stimulated CD4⁺ T cell proliferation in a dose dependent manner, which was overcome by sCD4 (Figure 2.5A-

B), similar to that observation for Env expressed on cells. Variability in the amount of suppression was observed for different preparations of AT-2 virus ranging from 29% (Figure 2.5A-B) to greater than 60% (data not shown). Decreased fluorescence intensity of anti-CD4 SK3 mAb staining for virion-incubated PBMC¹⁴ verified a functional interaction between CD4 and Env (data not shown). Interestingly, we found that proliferation of CD4 negative cells was also abrogated after incubation with 1 µg/ml virions (Figure 2.5A), which might be due to impairment of CD4⁺ T cell helper functions by Env, leading to decreased growth cytokine production (IL-2 and others) and less activated APCs in PBMC that, in turn, cause a decrease in antigen-stimulated CD8⁺ T cell proliferation.

Figure 2.5

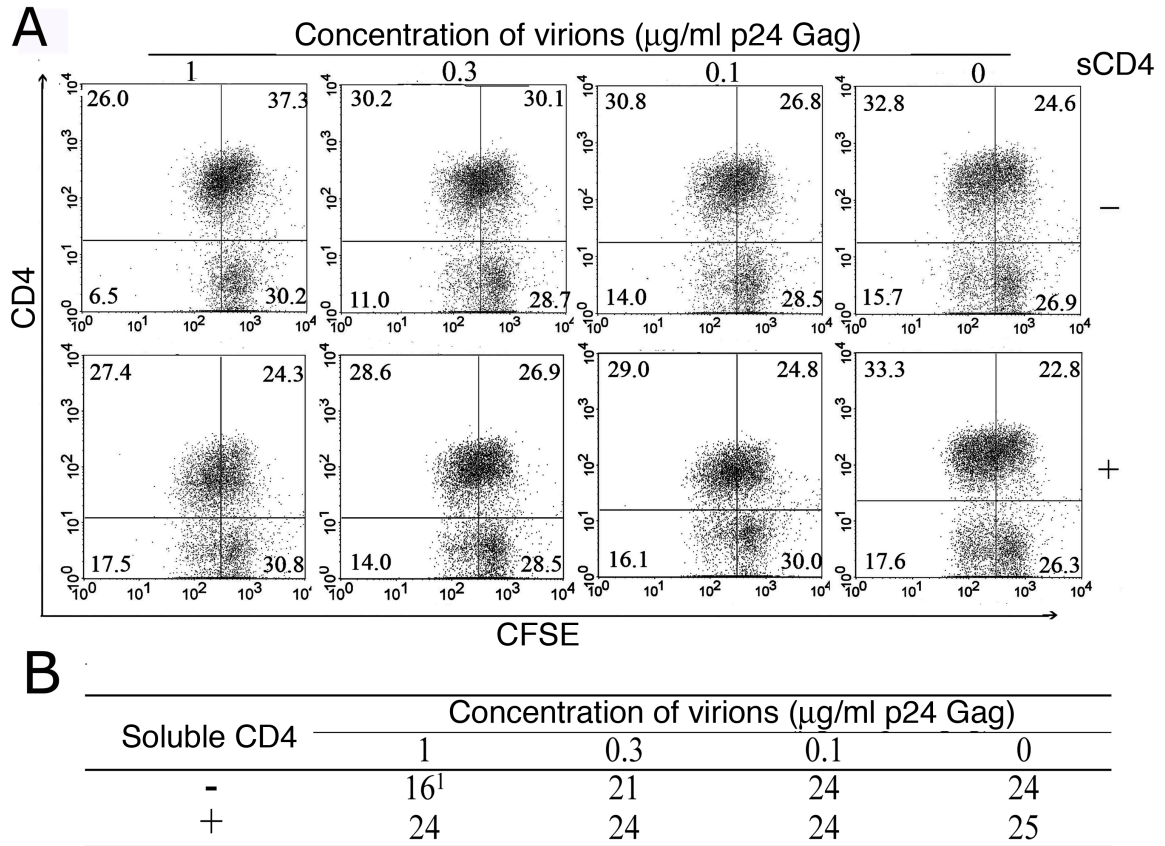


Figure 2.5. HIV Env present on free virions induces suppression of anti-CD3 stimulated CD4⁺ T cell proliferation through CD4 binding. Freshly isolated human PBMC were CFSE labeled and exposed to AT-2-treated HIV-1 virions (A and B) in the presence or absence of sCD4 (10 $\mu\text{g/ml}$). The cultures were stimulated with anti-CD3 mAb and IL-2. Four days after Env exposure and stimulation, PBMC were stained for CD4 and the proliferation of CD4⁺T cells was monitored by flow cytometry (A). The percent of input CD4⁺ T cells that proliferated (B) was calculated as described¹⁶⁴. Data are representative of three experiments. B, ¹p < 0.05. p-values calculated using 2-tailed

homoscedastic student's t-test comparing AT-2 virus-treated CD4⁺ T cell proliferation to no virus treated.

CD25⁺ Treg cells are not required for Env induced suppression

We then investigated whether Treg cells are involved in Env induced suppression in our system. CFSE stained total or CD25 depleted PBMC were cultured with Env-expressing or control 293T cells in the presence of anti-CD3 and IL-2 stimulation. CD25 staining verified that more than 95% of CD25⁺ cells were successfully removed from PBMC (Figure 2.6A). As expected, CD25 depletion significantly enhanced the proportion of proliferating CD4⁺ T cells in both Env exposed (From 8.1% to 18%) and control cultures (From 13% to 32.7%, Figure 2.6B). However, exposure to Env caused almost a 50% decrease in the proportion of proliferating CD4⁺ T cells in CD25 depleted PBMC (From 18% to 32.7%, Figure 2.6B), suggesting that pre-existing CD25⁺ Treg cells were not required for Env induced suppression of CD4⁺ T cell proliferation.

Figure 2.6

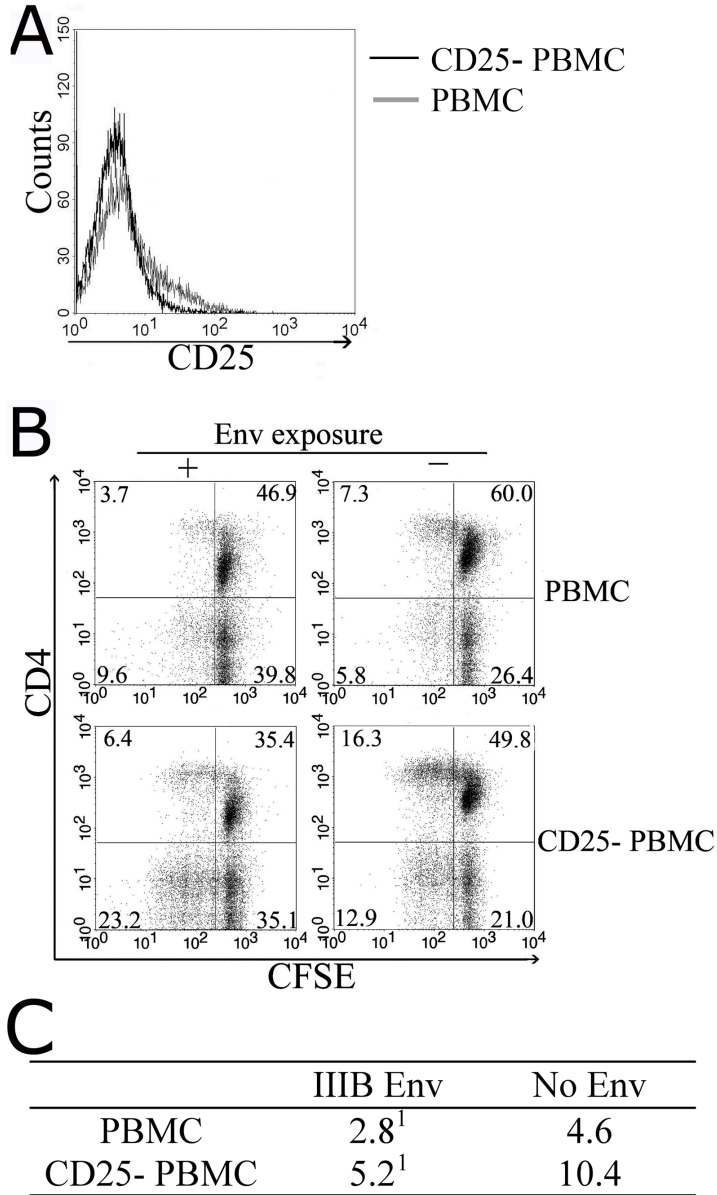


Figure 2.6. CD25⁺ Treg cells are not required for Env-induced suppression of proliferation. CD25⁺ cells were depleted from human PBMC using two rounds of CD25 magnetic beads. Flow cytometric analysis demonstrated depletion of greater than 95% of CD25⁺ cells (A). CFSE labeled whole (top, B) or CD25-depleted (bottom, B) PBMC

were exposed or not to IIIB Env-expressing 293T cells in the presence of anti-CD3 and IL-2 stimulation. CD4⁺ T cell proliferation 4 days after Env exposure is shown. (C) The percent of input CD4⁺ T cells that proliferated is shown for each condition in B.

¹=p<0.01. p-values were calculated using a 2-tailed homoscedastic student's t test comparing Env-treated CD4⁺ T cell proliferation to no virus-treated. Data are representative of at least four repetitions.

Effect of Env on frequency of CD4⁺, CD25⁺, FoxP3⁺ cells in PBMC

We show that CD25⁺ Treg cells are not required for Env to induce suppression of CD4⁺ T cell proliferation. One important question that remains highly controversial is whether and how HIV, or its associated gene products, regulates Treg cells. Both increased and reduced frequency of Treg cells have been reported in HIV⁺ patients compared to uninfected people ^{1, 149, 161}. To test whether Env, as the major surface antigen of HIV interacting with CD4⁺ T cells, directly induces or activates Treg cells, we first determined and compared the percentage of CD4⁺, CD25⁺, FoxP3⁺ cells in both stimulated and resting human PBMC with or without exposure to Env expressed on 293T cells. Compared to resting PBMC, stimulation caused a one- to two-fold increase in the amount of CD4⁺, CD25⁺, FoxP3⁺ T cells in PBMC at all time points studied (Figure 2.7A), which is consistent with a study demonstrating activation induced FoxP3 expression in human T cells ⁶⁰. Of importance, for both resting and stimulated PBMC, Env exposure did not induce significant changes in the percentage of CD4⁺, CD25⁺, FoxP3⁺ T cells (Figure 2.7A). Analyses of T cell proliferation in the activated cells

demonstrated a significant reduction of CD4⁺ T cell division when cultured with Env mRNA transfected 293T cells, verifying Env expression on 293T cells and its inhibitory effect.

We then investigated whether Env present on free virions affected the percentage of FoxP3⁺ Treg cells in PBMC, as we observed inhibition of CD4⁺ T cell proliferation by free virions (Figure 2.5A-B). PBMC were incubated with AT-2 inactivated HIV-1 virions at 1 µg/ml p24 gag protein equivalent for various period of time. Surface and intracellular staining demonstrated no significant difference in the number of CD4⁺, CD25⁺, FoxP3⁺ cells between virion incubated and control PBMC at all studied time points (days 1, 3 and 5) (Figure 2.7B), which is in contrast with the observation that exposure of purified CD4⁺ T cells to AT-2 treated HIV resulted in significantly increased FoxP3 expression at both mRNA and protein levels, resulting from Env induced enhancement in cell viability and consequent accumulation of FoxP3⁺ cells¹¹². Taken together, our data suggested that neither Env delivered on cells as would be found during infection or after encoding vector delivery as part of a vaccine nor Env present on free virions could directly induce or activate T regulatory cells in resting or stimulated PBMC.

Figure 2.7

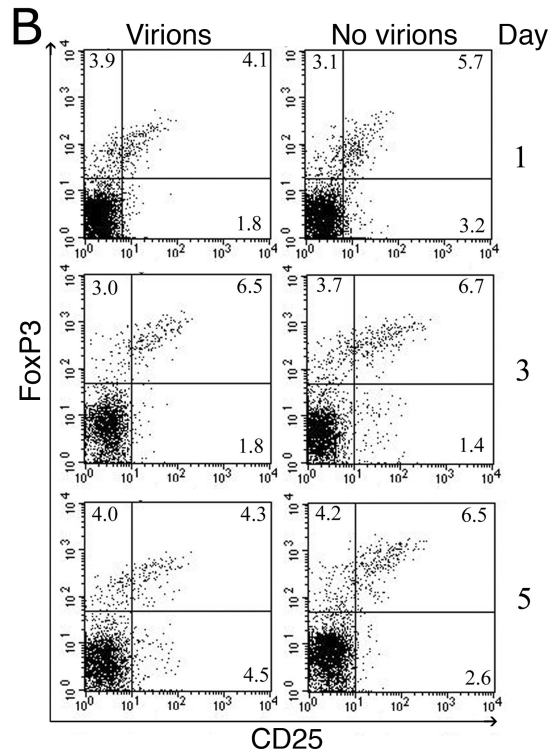
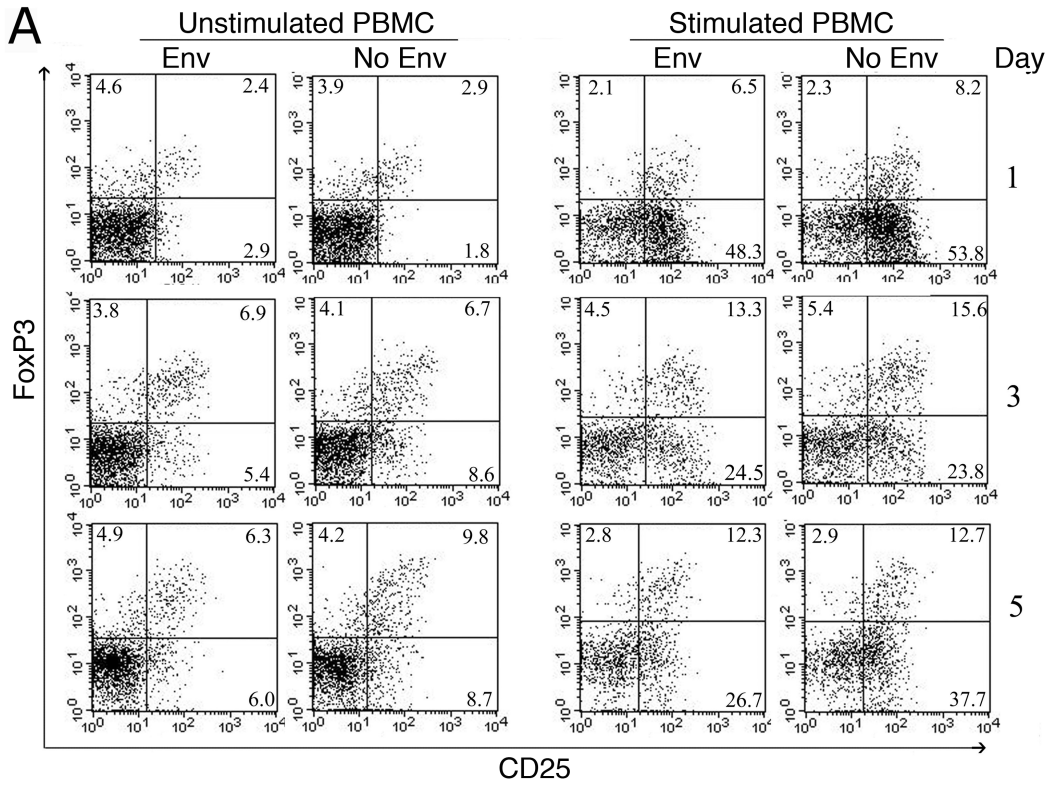


Figure 2.7. Effect of Env delivered by cell surface expression or on virions on FoxP3⁺, CD25⁺, Treg cells in unstimulated and stimulated PBMC. Unstimulated or anti-CD3 stimulated PBMC were analyzed by flow cytometry for CD25 and FoxP3 expression on CD4⁺ T cells at days 1, 3, and 5 after exposure to HIV Env expressed by 293T cells (A) or AT-2-treated virions (B). Histograms shown are gated on live CD4⁺ cells. Data are representative of at least four repetitions.

IL-10 and TGF- β production in Env exposed PBMC

Immune suppression mediated by Treg cells can occur through either cell contact or suppressive cytokines, particularly IL-10 and TGF- β ^{31, 156}. To test whether Env exposure, rather than inducing an increased number of FoxP3⁺ cells, stimulates an enhanced production of IL-10 and/or TGF- β by Tregs, supernatants of cultures with and without Env exposure were collected at various time points (days 1, 3, and 5) after initiation of co-culture, followed by quantitation of IL-10 and TGF- β by ELISA. TGF- β was persistently undetectable at any of the studied time points in both Env exposed and control supernatants. Significant amounts of IL-10 were produced and accumulated in the supernatants starting at 12 hours after stimulation. However, no difference was observed in the amount of IL-10 production between Env-exposed and control cultures (Figure 2.8). Considered together, these data suggest that Env does not induce a more activated phenotype of Treg cells leading to increased suppressive cytokine production.

Figure 2.8

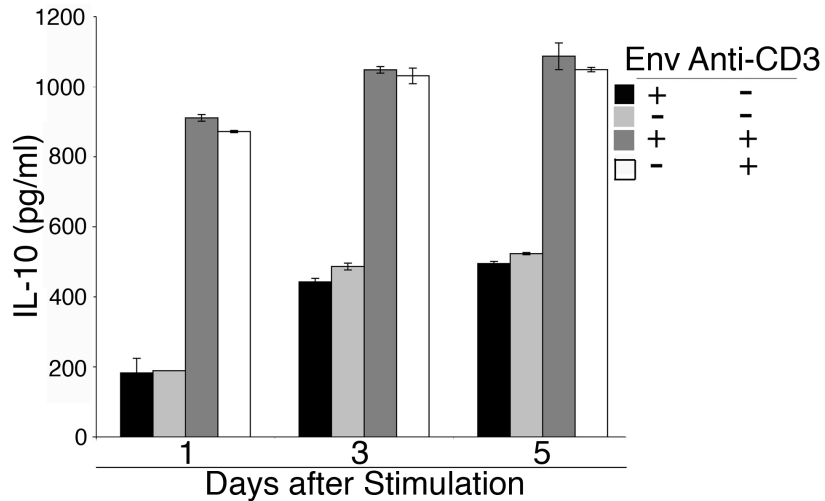


Figure 2.8. Effect of Env exposure on IL-10 production. 293T cells expressing IIB Env or control transfected were cultured with PBMC in the presence or absence of anti-CD3 and IL-2 stimulation. Supernatants were collected at the indicated time points, and supernatant associated IL-10 was determined by ELISA. Data are representative of at least five experiments. Error bars are the SEM for triplicate measurements.

Env does not enhance suppressive activity of CD4⁺, CD25⁺, CD127⁻ T cells

We demonstrate that Env induces neither an increased number of CD4⁺, CD25⁺, FoxP3⁺ cells nor an enhanced IL-10 and TGF- β production in PBMC and their depletion does not negate Envs ability to suppress CD4⁺ T cell proliferation. To test whether Env activates Treg cells, resulting in an enhanced suppressive activity independent of IL-10 or

TGF- β , an in vitro suppression assay was developed in which Treg cells isolated from Env exposed and control PBMC were cultured with autologous responder cells in the presence of anti-CD3 stimulation. The suppressive activity of Treg cells, on a per-cell basis, was evaluated by their capacity to inhibit anti-CD3 stimulated responder cell division. In addition to CD25, CD127, a subunit of the IL-7R, has also been shown to be a useful cell surface marker for defining Tregs. We sorted Tregs from control RNA or Env RNA transfected 293T cell exposed PBMC based on CD4⁺, CD25⁺, and CD127⁻ expression⁹⁴. Greater than 90% of the sorted cells expressed FoxP3 (Figure 2.9A). In addition to natively expressed FoxP3 in a subset of CD4⁺ T cells in PBMC, an additional population of FoxP3⁺ cells appears after activation. These cells are derived from FoxP3⁻, CD4⁺ cells and their ability to suppress immune responses is controversial⁶⁰. We attempted to use this same sorting strategy to isolate Tregs after Env exposure and anti-CD3 stimulation, but it should be noted that activation induces up-regulation of CD25 as well as down-regulation of CD127 on non-Treg CD4⁺ T cells, making the isolation of FoxP3⁺ cells less accurate. However, approximate 30-50% of the sorted cells from stimulated PBMCs expressed FoxP3 (data not shown).

Sorted CD4⁺, CD25⁺, CD127⁻ suppressor cells inhibited responder CD4⁺ T cell divisions in a dose dependent manner (Percentages of proliferating CD4⁺ T cells as the proportion of CD4⁺ T cells are 48% and 74% for Suppressor: Responder ratio of 1:3 and 1:6, respectively) (Figure 2.9B-C). However, Env-exposed and control CD4⁺, CD25⁺,

CD127⁻ T cells were equally suppressive, since no significant difference in the proportion of dividing responder cells was found between them (Figure 2.9B-C).

We then determined whether Env induces a more suppressive activity of activation induced FoxP3⁺ cells by sorting CD4⁺, CD25⁺, CD127⁻ T cells from PBMC, which were first depleted of CD25⁺ cells and then co-cultured with 293T cells and activated by anti-CD3. Activation induced CD4⁺, CD25⁺, CD127⁻T cells (approximately 30-50% FoxP3⁺), although less suppressive than naturally FoxP3⁺ cells isolated from unstimulated total PBMC, also inhibited CD4⁺ T cell proliferation in a dose dependent manner (Figure 2.9D-E). Of importance, no difference in suppressive ability between Env exposed and control activation-induced Treg cells was observed (Figure 2.9D-E). Taken together, these data indicate that Env exposure does not enhance the functional suppressive activity of either pre-existing Treg or activation induced FoxP3⁺ cells.

Figure 2.9. Endogenously present or activation induced (plus FoxP3, CD25) FoxP3 Treg cell suppressive activity is not modulated by Env. (A) Isolation of FoxP3⁺ Treg cells from PBMC using CD4, CD25, and CD127. CD4⁺ live lymphocyte sized cells (gate in left panel of A) were sorted for high CD25 and no CD127 expression (gate in middle panel of A). The sorted cells were stained for FoxP3 (A, right panel) and demonstrated greater than 90% positivity. (B) As a source of suppressors, PBMC were cultured with Env expressing or control 293T cells, followed by isolation of CD4⁺, CD25⁺, CD127⁻ T cells by flow cytometric sorting (detailed in A) 48 h after Env exposure. CFSE-labeled responder cells (autologous PBMC depleted of CD25⁺ cells) were cocultured with graded numbers of sorted CD4⁺, CD25⁺, CD127⁻ suppressor cells in the presence of anti-CD3 and IL-2 stimulation. CFSE intensity of responder cells with CD4 staining was monitored by flow cytometry at day 4 to quantify responder cell proliferation. (C) Percent of input CD4⁺ T cells that proliferated was calculated for each condition. ¹=p < 0.007. p-values calculated using 2-tailed homoscedastic student's t-test comparing 293T Env-treated CD4⁺ T cell proliferation to no Env-treated. (D) As a source of suppressors, CD25-depleted PBMC were cultured with Env expressing or control 293T cells in the presence of anti-CD3 and IL-2 stimulation, followed by isolation of CD4⁺, CD25⁺, CD127⁻ T cells 48 h after stimulation, as described in A. CFSE-labeled responder cells (autologous PBMC depleted of CD25⁺ cells) were cocultured with graded numbers of sorted CD4⁺, CD25⁺, CD127⁻ suppressor cells in the presence of anti-CD3 and IL-2 stimulation. CFSE intensity of responder cells with CD4 staining was monitored by flow cytometry at day 4 to quantify responder cell proliferation. (E) Percent of input CD4⁺ T cells that

proliferated was calculated for each condition. $^1p < 0.009$. p-values calculated using 2-tailed homoscedastic student's t-test comparing 293T-Env-treated CD4⁺ T cell proliferation to no Env-treated. Data are representative of at least three repetitions.

Discussion

The system studied here models the encounter of CD4⁺ T cells with both APCs and non-APCs expressing Env delivered as a vaccine, the interaction between CD4⁺ T cells and infected cells in an HIV-infected individual, and the interaction between Env expressed on a virion and CD4⁺ T cells. Both X4 and R5 Envs suppressed the activation and proliferation of CD4⁺ T cells via specific interaction with CD4. Suppression was characterized by a lack of proliferation despite normal activation of the main signaling pathways involved in T-cell activation and proliferation, AKT, MAPK, and STAT5a but a deficient upregulation of early and late cell-surface activation markers. The suppression of antigen-induced proliferation was not due to IL-2-dependent anergy, apoptosis, or cell death.

The role of CD4 versus the coreceptors CXCR4 and CCR5 in mediating Env-induced suppression is controversial^{7, 15, 71, 154}. Some reports have suggested a role for coreceptors in mediating suppression in conjunction with CD4^{15, 154}. Others have suggested that Env can mediate suppression solely via CD4 or coreceptors^{7, 71} {Arthos, 2002 #110}. In this study, we show that Env delivered as a vaccine, as a component of an infected cell, or as part of a virion mediates a suppression of activation and proliferation

regardless of its coreceptor usage. Although CD4 interaction is required for this suppression, both coreceptors could have played a role. To test this, we examined suppression of X4 Envs in the presence of AMD3100, which blocks Env interaction with CXCR4⁴⁵. AMD3100 caused a non-specific increase in proliferation in both Env and control transfected 293T-exposed CD4⁺ T cells, which may be attributed to blocking inhibitory chemokine signaling through CXCR4¹⁵⁵. Similar studies with TAK-779⁹, which blocks R5 Env interaction with CCR5, did not overcome suppression. This suggests that in our system studying Env in the absence of HIV infection, inhibition of CD4⁺ T cell proliferation occurred in a CD4-dependent, chemokine receptor-independent manner.

Previous studies demonstrating the suppressive effects of Env have focused on a single aspect of CD4⁺ T cell suppression such as apoptosis or anergy and used systems to model interactions that occur during HIV infection. In this chapter, I present a comprehensive study examining multiple facets of Env-induced suppression when it is used as an immunogen, present on a virion, or expressed in a manner that models HIV-infection of cells. DCs that acquire a vaccine containing HIV Env will express protein from all the delivered antigens. When the protein is processed and presented to antigen-specific CD4⁺ T cells by the APCs, suppression by surface-expressed Env would occur, thus hindering immune responses to all antigens in the cocktail. Because Env is an essential component in an HIV vaccine that aims to generate neutralizing antibodies, inclusion of nonsuppressive forms of Env will enhance immune responses.

Accumulating studies have implicated the involvement of CD25⁺ Treg cells in the inhibition of anti-HIV CD4⁺ and CD8⁺ T cell responses in HIV infection^{1, 5, 52, 85, 86, 112, 161}. The part of thesis was also designed to address three questions: 1) whether HIV virion expressed Env, like that expressed on the cell surface, induces inhibition of polyclonally stimulated CD4⁺ T cell proliferation; 2) whether Treg cells are responsible, at least in part, for Env induced suppression; and 3) whether Env is able to directly induce, or activate, Treg cells. The answers to these questions are critical to determine whether the frequency and function of Treg cells are regulated through direct interaction with virus, and how this regulation of Treg cells relates to dysfunction of T cell immunity in HIV infection. Here, I demonstrate that Env expressed on either virions or cells induces suppression of CD4⁺ T cell activation, and that Env does not induce or activate Treg cells as a mediator for this inhibition.

In HIV infection, in addition to infected cells, circulating free virions also bear functional trimeric Env. Although protease-defective, gp120-containing HIV-1 particles were reported to induce apoptosis in human peripheral blood T cells, whether such virions use Env, like its counterpart expressed on cells, to induce inhibition to CD4⁺ T cell proliferative responses remains unclear⁷⁹. Using AT-2 inactivated HIV-1 viral particles that are non-infectious but maintain integrity of functional Env^{112, 131}. I demonstrate that Env on HIV virions inhibits CD4⁺ T cell proliferation through gp120-CD4 signaling.

HIV impairs host CD4⁺ T cell immunity through multiple mechanisms. Although the involvement of Treg cells in suppressing anti-HIV specific T cell responses has been documented in a growing body of literature, there is controversy as to whether HIV, or its gene products, inhibits host CD4⁺ T cell activation through the direct induction or activation of Treg cells. An interaction between Treg cells and HIV through gp120-CD4 signaling has been proposed¹¹², which provides a basis for our in vitro system where human PBMC, as source of CD4⁺ T cells, were exposed to Env expressing 293T cells or AT-2 inactivated virions, modeling the in vivo interaction between Env and CD4⁺ effector as well as CD4⁺ Treg cells, during an immune response. In this thesis, I show that depletion of pre-existing CD4⁺, CD25⁺ Treg cells from PBMC increases the proliferation of polyclonal TCR stimulated CD4⁺ T cells, which is consistent with observations in HIV infected patients that virus exposed Treg cells maintain suppressive activity. However, Env-induced inhibition remained in the absence of CD25⁺ Treg cells, strongly suggesting that Treg cells are not functionally involved in Env-induced suppression. This result implies that the suppressive effect of Treg cells on anti-HIV T cell responses, which has been observed in HIV infected patients, may be a general feature of Treg cells.

The regulation of Treg cells in HIV remains highly controversial. Studies using peripheral blood reported increased^{1, 149, 161} as well as reduced^{49, 115} numbers of Treg cells in HIV infection. In contrast, later studies using LN or gastrointestinal mucosa of

infected patients demonstrated up-regulation of FoxP3 and other markers of Treg cells, supporting Treg accumulation in the tissue at sites of viral replication^{5, 6, 52, 112}. Factors including markers used to identify Treg cells, disease stage in selected patients, and ART treatment, might contribute to the various findings from different studies. Our results demonstrate that regardless of the form of Env delivery (Env-expressing cells as would be observed during infection or after encoding vaccine delivery or free virions) and stimulation status of T cells, the percentage of FoxP3⁺, CD25⁺, CD4⁺ T cells was not affected by Env, which is in contrast to the observation that an increase in FoxP3⁺ cells was detected in CD4⁺ T cells after exposure to AT-2 inactivated virions, resulting from Env induced enhancement in Treg cell viability¹¹². The difference in cells used (PBMC vs purified CD4⁺ T cells) could have contributed to the different results in these two studies. The purification of CD4⁺ T cells could have increased their level of apoptosis, which was blunted by Env acting on Tregs or increased by Env acting on non-Treg CD4⁺ T cells. We observed a low level of apoptosis in our unstimulated PBMC with no difference between Env exposed and control (data not shown). Although, if this were the case, we would have expected to see a difference after anti-CD3 stimulation (Figure 2.7A), which increases apoptosis.

It has been reported that, unlike murine Foxp3⁻ T cells, a subset of human FoxP3⁻ T cells could be induced to transiently express FoxP3 upon activation^{60, 98}. In agreement with these studies, we found that anti-CD3 stimulation induced increased FoxP3 expression in CD4⁺ T cells, peaking approximately 48 hours after activation (data

not shown). Furthermore, we measured FoxP3 expression in CD25-depleted PBMC, and found that activation induced FoxP3 expression, which is in agreement with a published report⁵³. These activation induced FoxP3⁺, CD4⁺ T cells have suppressive activity on T cell proliferation, but this activity was not modulated by Env exposure.

One of the functional characteristics of Treg cells is its capacity to suppress antigen specific CD4⁺ and CD8⁺ T cell proliferation in a dose-dependent manner¹⁵⁶. Since FoxP3 is expressed intracellularly, several surface antigens in addition to CD25 have been identified as alternative markers for isolation of Treg cells, one of which is CD127, a subunit of the IL-7R⁹⁴. Analysis of sorted CD4⁺, CD25⁺, CD127⁻ T cells, which contained 90% FoxP3⁺ cells, demonstrated that Env exposure did not alter the suppressive activity of peripheral blood Treg cells, which is consistent with a previous finding that Treg cells exposed to AT-2 treated HIV virions are equally suppressive compared with untreated Treg cells¹¹².

In summary, I used models of the encounter of CD4⁺ T cells with both APCs and non-APCs expressing Env and virion expressed Env to study Env mediated suppression of CD4⁺ T cell proliferation. Both X4 and R5 Envs suppressed the activation and proliferation of CD4⁺ T cells via specific interaction with CD4. Suppression was characterized by a lack of proliferation despite normal activation of the main signaling pathways involved in T cell activation and proliferation, AKT, MAPK, and STAT5a but a deficient upregulation of early and late cell surface activation markers. The suppression

of antigen-induced proliferation was not due to IL-2 dependent anergy, apoptosis, or cell death. My data also demonstrates that HIV Env neither induced nor activated Treg cells, and that CD25⁺ Treg cells are not responsible for Env induced suppression of CD4⁺ T cell proliferation. Our findings do not demonstrate that HIV does not regulate Treg number or function, only that Env binding does not, in model in vitro assays. Treg cells could still be depleted by direct infection or bystander effects or their number increased by other gene products or downstream effects of infection, such as chronic immune stimulation. Further work is needed to elucidate the in vivo regulation of Treg cells and their physiological relevance in the context of HIV infection and vaccination.

CHAPTER 3

In vivo analysis of immune and viral responses in retroviral suppressed

SIV infected rhesus macaques

Introduction

CD4⁺ T cells play a central role in maintaining effective cellular and humoral immune responses by providing help to CD8⁺ T cells, B cells and innate effectors. One of the hallmarks of HIV infection is the progressive loss of CD4⁺ T cells and their associated antigen-specific responses. This is characterized as both a decline in the number of CD4⁺ T cells and an early loss of the functional activity of cells with certain antigenic specificities^{41, 90, 105}. In particular, HIV antigen-specific CD4⁺ T cell responses are dysregulated to a greater extent than responses to other common antigens^{88, 91, 158}. It has been demonstrated that chronically HIV-1 infected progressing patients fail to mount a potent HIV-specific CD4⁺ T cell response even in the setting of viral suppression by ART. Immunization of chronically infected ART treated patients with HIV antigens leads to poor HIV-specific responses⁵⁰. The mechanisms for the inability to generate HIV-specific CD4⁺ T cell responses in chronic infection are not fully characterized or understood.

To initiate an immune response, DC bearing antigens interact with antigen-specific CD4⁺ T cells that migrate through lymphoid tissue, leading to CD4⁺ T cell activation and expansion. Signals from the DC induced by the pathogen instruct the type

of T cell response, characterized as Th1, Th2, Th17, Tr1, and Th3/Tr2^{12, 36, 64}. However, in the presence of HIV, the interaction between DC and CD4⁺ T cells enhances viral replication by multiple mechanisms¹⁶⁸. It has been demonstrated that even in the presence of potent regimens of viral suppression, a very low level of viral production could still be detected^{43, 70, 122}. It is generally believed that this viral replication does not represent acute infection of cells, as no sequence evolution is observed^{32, 84, 122}, but the source of the low level of virus may be derived from DC mediated activation of latent virus in memory T cells or other reservoirs¹²⁶.

I use a macaque model of HIV-infected individuals treated with ART during chronic infection to study the effect of SIV antigen stimulation in lymph nodes early after immunization. CMV seropositive rhesus macaques were infected with SIVmac251 and after 4 months, treated with D4T and PMPA to artificially control viral loads. Animals were immunized in both arms and legs with SIV or CMV antigen, which allowed draining lymph nodes for each antigen to be obtained from the same animal at the same time, allowing a direct comparison of the effect of SIV and CMV antigen stimulation (Figure 3.1). Immunologic and virologic responses were measured in draining lymph nodes and peripheral blood to analyze differences in responses induced by an SIV antigen compared to a control immunization.

Materials and Methods

Ethics statement

All animal experiments were performed in strict accordance with good animal practice following guidelines for relevant national and regional animal welfare bodies. The studies were approved by the University of Pennsylvania and Tulane Institutional Animal Care and Use Committees (IACUC).

Immunogens

Plasmid DNA expressing the SIV Gag core protein from Mac239 (pSIVgag) was used. It is a Rev-independent expression vector that has been codon optimized for a high level of expression, as previously described²⁰. Plasmid DNA expressing the Rhesus CMV pp65 protein was used¹⁷⁰. DNA was formulated in 0.15 M citrate buffer and 0.25% bupivacaine at a pH of 6.5.

Infection of rhesus macaques and overview of study

The animal study timeline is shown in Figure 3.1. Three CMV seropositive Mamu-A*01 positive rhesus macaques were intravenously infected with SIVmac251 at time zero. Four months after SIV infection, ART was introduced (SQ PMPA 20 mg/kg qD and PO D4T 10 mg/kg qD) and continued until the end of the experiment. The macaques were followed for 4 months on ART for recovery of periphery CD4⁺ T cell count and viral load suppression. Eight months after infection and 4 months after the initiation of ART, each monkey received 2 immunizations with SIV gag encoding DNA

i.m. (2 mg per injection), one in the left arm and one in the left leg and 2 immunizations with CMV pp65 encoding plasmid (2 mg/injection) in the right arm and leg. After immunization, draining lymph nodes were sampled at two-time points for each animal by first removing inguinal LNs on each side followed by axillary LN removal from both sides (FH40 - D3 and D9, DD05 - D5 and D11, and CT64 - D7 and D14). We were unable to collect pre-immunization LNs due to technical limitations. D60 LNs were collected as no-immunization control. PBMC and plasma samples were collected every 2 to 4 weeks throughout the experiment.

Measurement of peripheral viral load and CD4⁺ T cell counts

Viral RNA in plasma was quantified by a bDNA signal amplification assay from Bayer Inc., version 4.0, specific for SIV, which has a threshold detection limit of 125 viral copies per ml plasma¹⁵¹.

CD4⁺ T cell counts using whole blood collected in EDTA were analyzed with anti-CD3 (Biosource International, Camarillo, CA), anti-CD4 and anti-CD8 staining (anti-Leu3a and anti-Leu2a, Becton Dickinson, San Jose, CA) as previously described³⁵.

PBMC isolation and LN biopsy

Peripheral blood lymphocytes were isolated from EDTA-blood using Ficoll-diatrizoate gradient centrifugation and analyzed for CD4's by flow cytometry or cryopreserved in 90% FCS 10% DMSO. LNs were obtained under anesthesia. The LNs

were teased carefully to generate single-cell suspensions. The single-cell suspensions from lymph nodes were divided into two parts; one was cryopreserved and stored in liquid nitrogen and one was frozen as a pellet.

In vitro stimulation

Cryopreserved LN cells or PBMC were thawed and washed twice with pre-warmed complete RPMI 1640 supplemented with 10% heat-inactivated FCS (HyClone) and L-glutamine (Invitrogen Life Technologies). Cells were resuspended in complete RPMI medium and rested for two hours at 37°C before being washed again and used. Unless otherwise noted, cells were prepared at concentration of 1×10^6 cells/ml for in vitro stimulation. For characterizing the SIV- and CMV-specific T cell responses, LN cells or PBMC were incubated with either an SIVmac239 gag 15-mer peptide pool with 11-amino acid overlap, 2 µg/ml each peptide (NIH AIDS Research and Reference Reagent Program, Bethesda, MD) or a rhesus CMV pp65 peptide pool (15-mers overlapping by 11 amino acids) at 2 µg/ml for each peptide¹⁷⁰ at 37°C for 6 hours in the presence of Golgi-Stop (0.7 µg/ml) and Golgi-Plug (1 µg/ml) and 1 µg/ml of co-stimulatory antibodies anti-CD28 and anti-CD49d (BD Bioscience, CA). Negative controls with no stimulation and positive controls with PMA/Ionomycin (50 ng/ml, Sigma-Aldrich) were used.

Cell staining

After 6 hours of stimulation, cells were first washed with 2 ml PBS containing 1% FCS and stained with aqua blue dye for cell viability (Invitrogen, Eugene, OR). Cells were then stained with pre-titrated amounts of fluorochrome-conjugated surface staining antibodies (anti-CD4-PerCP Cy5.5, anti-CD8-FITC, anti-CD14-Pac Blue, anti-CD16-Pac Blue, and anti-CD95-PE-Cy5, anti-CD20-Pac Blue (eBioscience), and anti-CD28-ECD (Bechman Coulter)) and incubated at 4°C for 20 minutes. Cells were then washed and fixed in 250 µl BD Fixation/Permeabilization solution (BD Biosciences) for 20 minutes at 4°C. After fixation, cells were permeabilized with 1 x BD Perm/Wash buffer and stained with pre-titrated fluorochrome-conjugated antibodies (anti-CD3-APC-Cy7, anti-IL-2-PE, anti-IFN- γ -APC, anti-TNF- α -PE-Cy7 (BD Biosciences) at 4°C for 45 minutes. Cells were then washed with Perm/Wash buffer and re-suspended in 300 µl PBS 1% FBS.

Flow cytometric analysis

Cells were analyzed by polychromatic flow cytometry on an LSRII (BD Biosciences) equipped for the detection of 18 fluorescence parameters. In this study, for all 12-color flow analyses, 200,000 to 500,000 events were acquired for each sample. Antibody capture beads (BD Biosciences) stained separately with individual mAbs were used to conduct electronic compensation. Flow Jo version 8.8.7. (Tree star, Ashland, OR) was used to analyze the polychromatic flow data with all the analytic gating as described before^{16,25}. Single cells were gated according to Forward Scatter Height (FSC-H) and Forward Scatter Area (FSC-A). Dead cells, B cells, NK cells, and monocytes were

excluded based on staining with aqua blue LIVE/DEAD fixable violet dead cell stain or CD14, CD16 and CD20 staining. Lymphocytes according to FSC-A and SSC-A, CD3⁺, CD3⁺CD4⁻, and CD3⁺CD4⁺ were sequentially selected for cytokine analysis. Expressions of IL-2, TNF- α and IFN- γ were identified in CD3⁺CD4⁻ and CD3⁺CD4⁺ T cell populations. In CD3⁺ lymphocyte population, memory and naive cells were identified through CD28 and CD95 staining as previously described^{16,25}.

RNA extraction and quantitative PCR

Total RNA from LN mononuclear cells was extracted using Trizol according to manufacturer's instruction (Invitrogen) and subject to real-time PCR on an ABI 7500 (Applied Biosystem, Carlsbad, CA). Doubly spliced, singly spliced, and unspliced SIV viral RNA and the housekeeping gene GAPDH were analyzed. The primers and MGB probes specific for these genes were obtained from Applied Biosystem (Table 4). Changes in the expression of individual viral RNAs with GAPDH normalization were calculated utilizing delta cycle threshold (ΔC_T) values, and the comparisons were made between LNs draining SIV and CMV immunizations from the same animal collected at the same time on the indicated days post immunization.

Levels of IFN- γ mRNA were quantitated by RT-PCR against a standard curve derived from serial dilutions of in vitro made transcripts, using primers and probe listed on Table 4. Copies of IFN- γ mRNA were expressed per 1 μ g of RNA.

Statistics

Mean, Standard deviation, and student's t-test were performed using Microsoft Excel software.

Table 4. Primers and Probes used in real-time PCR for quantitation of SIV viral RNA and macaque IFN- γ

RNA	Primers and Probe
SIV Doubly Spliced	Forward: 5' - AGGCTAATACATCTTCTGCATCAAAC-3' Reverse: 5' - GGGTCCTGTTGGGTATGAGTCTA-3' Probe: 5' - CCACCCTCTTATTTCC-3'
SIV Singly Spliced	Forward: 5' - AGAGGCCTCCGGTTGCA-3' Reverse: 5' - CCTTCCCCTTCCACAATAGC-3' Probe: 5' - ACTGTGGAAGGGACC-3'
SIV Unspliced	Forward: 5' - TTGCAGCACCCACAACCA-3' Reverse: 5' - TGATCCTGACGGCTCCCTAA-3' Probe: 5' - CTCCACAACAAGGACA-3'
Macaque IFN- γ	Forward: 5' - GTGTGGAGACCATCAAGGAAGAC-3' Reverse: 5' - CGACAGTTCAGCCATCACTTGGAT-3' Probe: 5' - ACTGACTCGAATGTCCAACGCAAAGC-3'
Macaque GAPDH	Forward: 5' - GGCATCCTGGGCTACACTGA-3' Reverse: 5' - AGGAGTGGGTGTCGCTGTTG-3' Probe: 5' - AGGTGGTCTCCTCTGAC-3'

RESULTS

PART 1

Infection and immunization of rhesus macaques and study overview

Pathogenic SIVmac251 infection of rhesus macaques has been well described for studying viral dynamics, immune responses, and anti-retroviral therapy^{102, 103, 107, 114, 153, 159}. All animals were Mamu A*01 positive to reduce MHC variation in disease course and T cell responses and naturally infected with Rhesus CMV. The study was designed to infect animals (1000 TCID₅₀ of SIVmac251 by intravenous injection) and allow them to reach steady state viral loads (4 months) followed by ART treatment (PMPA and D4T) for 4 months. Animals were then immunized with expression plasmids encoding non-mutated SIV gag or rhesus CMV pp65 in both arms and legs and lymph nodes draining either SIV gag or CMV pp65 plasmid injections from the same animal at the same time were obtained 3, 5, 7, 9, 11 and 14 days post immunization (Figure 3.1). Serum and PBMCs were obtained every 2 to 4 weeks throughout the experiment.

Figure 3.1

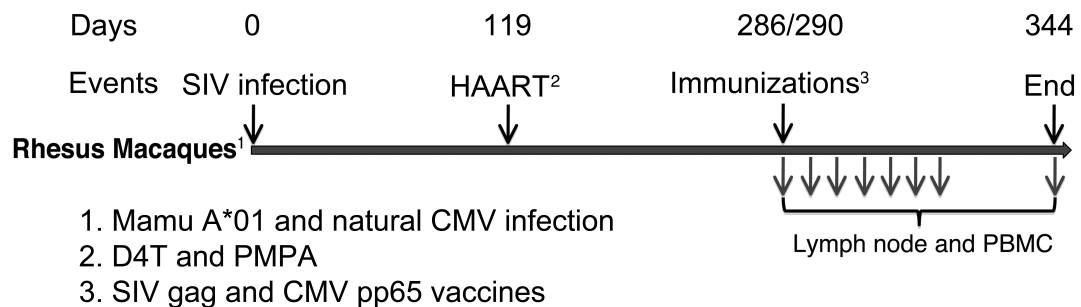


Figure 3.1. Experimental protocol for infection and immunization of rhesus

macaques. Mamu A*01 positive rhesus macaques naturally infected with rhesus CMV were intravenously inoculated with SIVmac251, followed by ART treatment (DT4 and PMPA) from days 119 post infection through the end of experiment. On days 286 or 290 post infection, monkeys received 2 immunizations with SIV gag encoding DNA i.m. (2 mg per injection), in the left arm and left leg, and 2 immunizations with CMV pp65 encoding DNA i.m. (2 mg per injection) in the right arm and leg. LN and PBMC were collected from the immunized animals.

PART 2

Establishment of chronically SIV-infected rhesus macaque with retroviral suppression and CD4⁺ T cell recovery by ART

ART suppresses viral replication leading to recovery of peripheral CD4⁺ T cell counts

SIV infection was established in all three animals with kinetics typical of SIV replication during primary infection in naïve rhesus macaques (Figure 3.2A)^{68, 143}.

Introduction of ART 4 months post infection, when set point viral loads had been established, efficiently suppressed viral replication to undetectable levels. One macaque demonstrated occasional blips in viral load but returned to undetectable levels after each without any change in therapy (Figure 3.2A).

Absolute CD4⁺ T cell counts in the peripheral blood demonstrated increases after introduction of ART in all animals with sustained levels of more than 500 cells/ μ l (Figure

3.2B). These findings demonstrate that chronic SIV infection was achieved and ART successfully suppressed viral replication leading to recovery of peripheral CD4⁺ T cell counts.

Figure 3.2

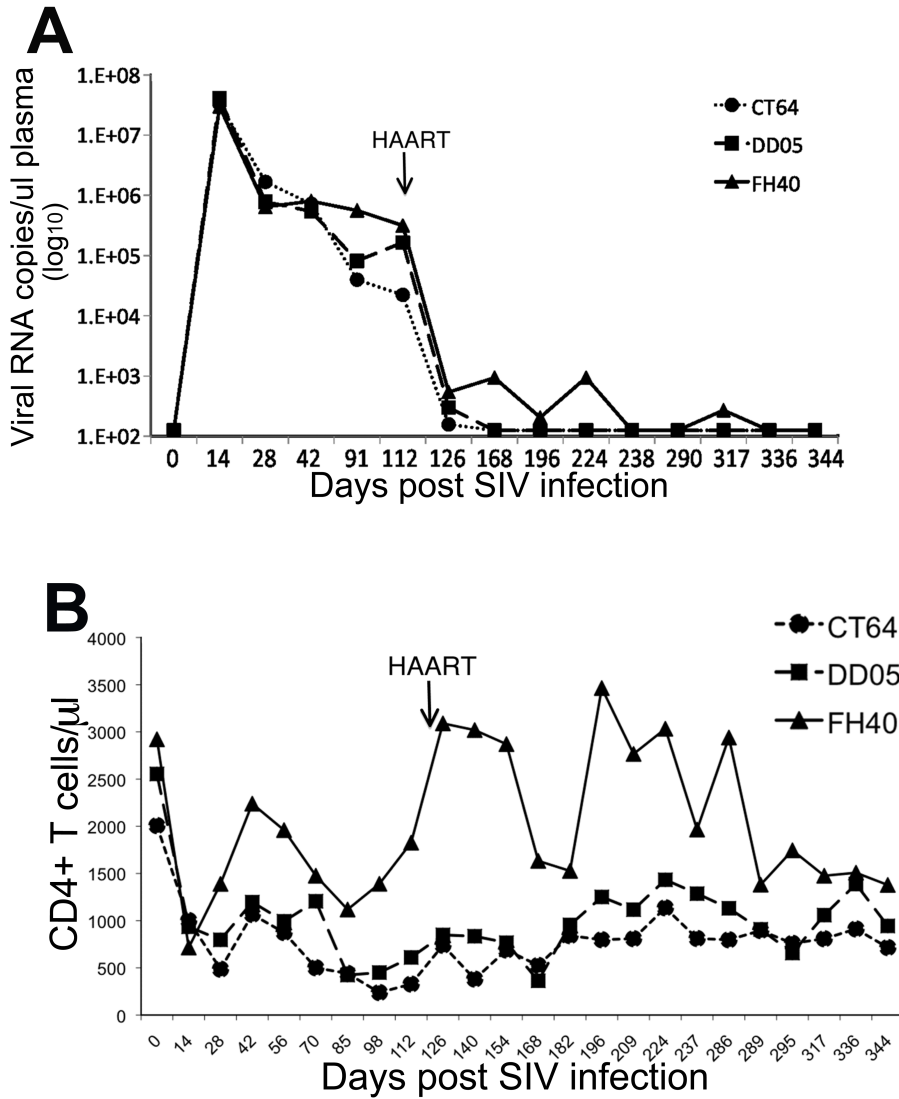


Figure 3.2. Summary of viral loads and peripheral CD4⁺ T cell counts in rhesus macaques. Viral loads and CD4 counts were measured every 2 to 4 weeks throughout the

experiment. (A) Viral RNA in plasma was quantified by a bDNA signal amplification assay and expressed as RNA copies per ml plasma. ART treatment controlled the viral loads in all three animals. (B) Peripheral CD4⁺ T cell counts increased after the initiation of ART. Macaque blood samples were stained for anti-CD3, anti-CD4, and anti-CD8 antibodies and the number of CD3⁺, CD4⁺ T lymphocytes were determined by flow cytometry. CD4⁺ T cell counts were expressed as CD4⁺ T cells per μ l blood.

PART 3

Immunization induced immunologic responses in draining lymph nodes

3.1. Immunization-induced immune activation in draining LNs

Animals were immunized with SIV gag encoding DNA in the arm and leg on one side and CMV pp65 encoding DNA on the other side at the same time. LNs, one draining an SIV immunization and one draining a CMV immunization, were excised from each animal at 2 time points post immunization. All comparisons were made between LNs from the same animal obtained at the same time that differed only in whether they drained an SIV or a CMV immunization. I was unable to obtain LNs prior to immunization due to limitations on the number of surgeries in a short time period. I obtained LNs from both the left and right sides of all animals 60 days post immunization. The measurement of immune responses did not demonstrate any difference between LNs draining HIV gag or CMV pp65 and these LNs were assumed to represent preimmune conditions.

LN mononuclear cells were assessed for the amount of IFN- γ mRNA, a major effector cytokine for adaptive immunity, as a measurement of total immune activation induced by immunization. The expression of IFN- γ was maximal 3 days post immunization in LNs draining both SIV gag and CMV pp65, which was followed by a decline at day 5 and a decrease in IFN- γ expression to the levels observed in 60 day post immunization LNs (Figure 3.3.1).

Figure 3.3.1

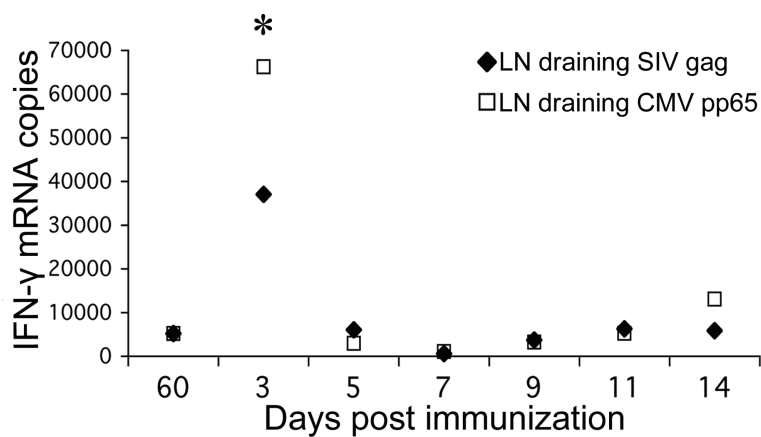


Figure 3.3.1. Total IFN- γ mRNA expression in LNs. Total RNA from LN mononuclear cells draining either SIV-gag or CMV-pp65, collected on indicated days post immunization were subject to real-time PCR with IFN- γ specific primers and probes for absolute quantitation of macaque IFN- γ RNA expression. The results were normalized and expressed as the RNA copy number per μ g total RNA. The comparisons between SIV- and CMV-immunization were made in LNs collected from the same

animal at the same time on indicated days. The difference on Day 3 was significant with a $p=0.036$ noted by a *.

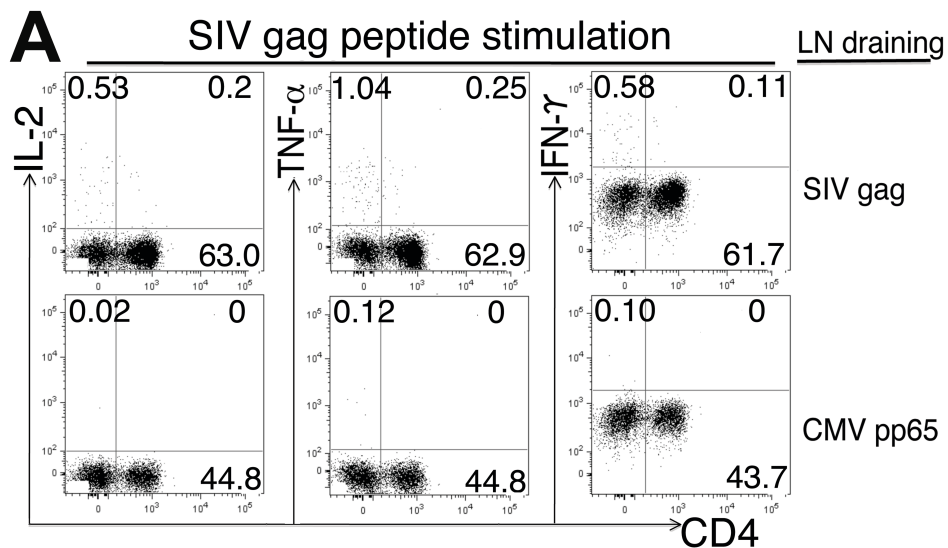
3.2. SIV immunization induced SIV gag-specific T cell responses in LN

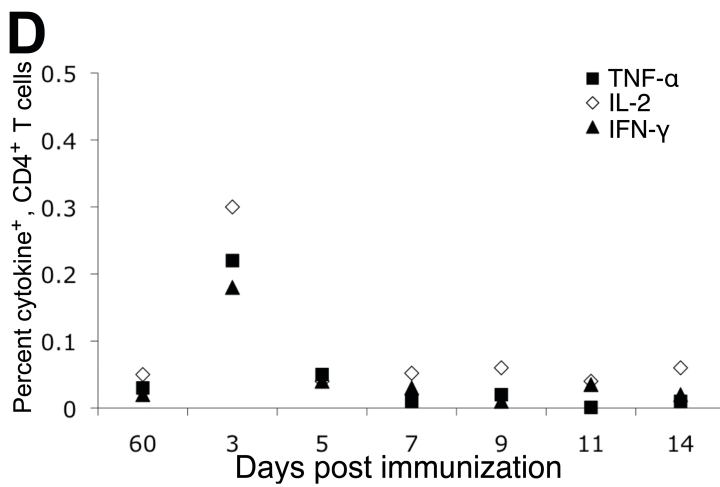
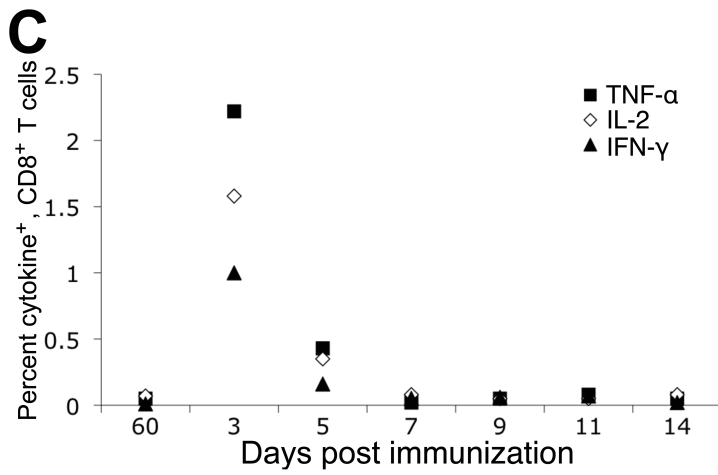
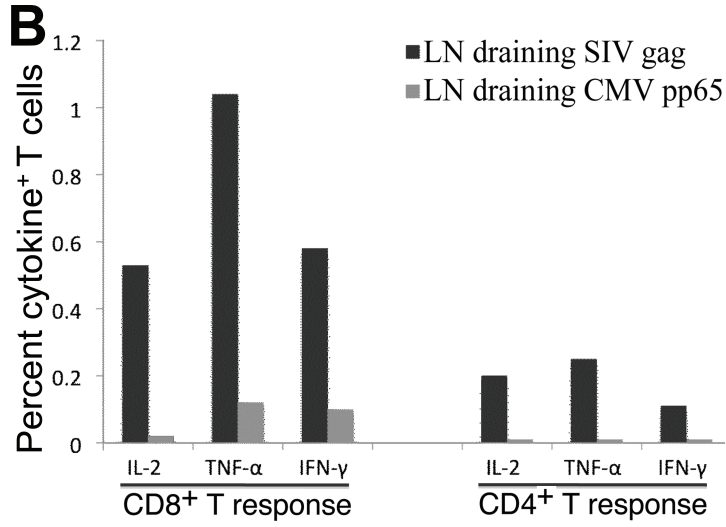
We next evaluated antigen-specific T cell responses by stimulation and intracellular cytokine staining. LN cells were stimulated in vitro with either SIV gag or CMV pp65 peptide pools and the percents of IL-2, TNF- α and IFN- γ -producing T cells were determined by polychromatic flow cytometry. Data of the LN draining SIV gag immunization on Day 3, when the immune activation, as measured by IFN- γ mRNA was the highest, demonstrated a potent response to gag peptide stimulation according to IL-2, TNF- α and IFN- γ productions (Figure 3.3.2A and B). In contrast, the LN draining the CMV pp65 immunization on day 3, when stimulated by gag peptides, demonstrated no significant response (Figure 3.3.2A and B).

SIV immunization-induced gag-specific CD8⁺ (Figure 3.3.2C) and CD4⁺ (Figure 3.3.2D) T cell responses in LNs for all time points post immunization are summarized. As shown, the overall patterns for CD8⁺ and CD4⁺ T cell responses are similar in that an immediate and potent memory T cell response is induced on Day 3 post immunization, which is followed by a decline on Day 5 and a further decrease to the levels observed in 60 day post immunization LNs (Figure 3.3.2C and D), consistent with the data obtained by measuring IFN- γ mRNA (Figure 3.3.1).

We next assessed the T cell subsets that mediated immunization-induced SIV gag-specific T-cell responses in Day 3 LN mononuclear cells. According to previous studies, CD28 and CD95 could be used to separate macaque T cells into naïve, central memory (CM) and effector memory (EM) populations. As shown in Figure 3.3.2E, the SIV-specific T cell responses were primarily mediated by central memory T cells in Day 3 LN. We then analyzed the polyfunctionality of SIV immunization-induced gag-specific T cells and found that both CD8⁺ and CD4⁺ SIV-specific T cell responses demonstrated a polyfunctional profile with the ability to produce 2 (IL-2/TNF- α , TNF- α /IFN- γ and IL-2/IFN- γ) and 3 cytokines (IL-2/TNF- α /IFN- γ), simultaneously (Figure 3.3.2F).

Figure 3.3.2





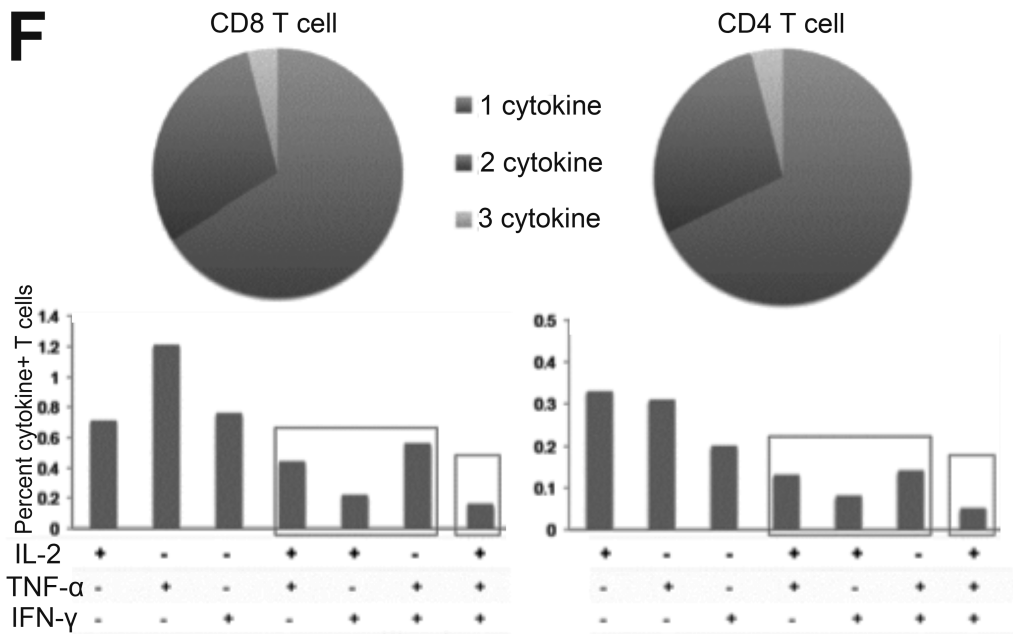
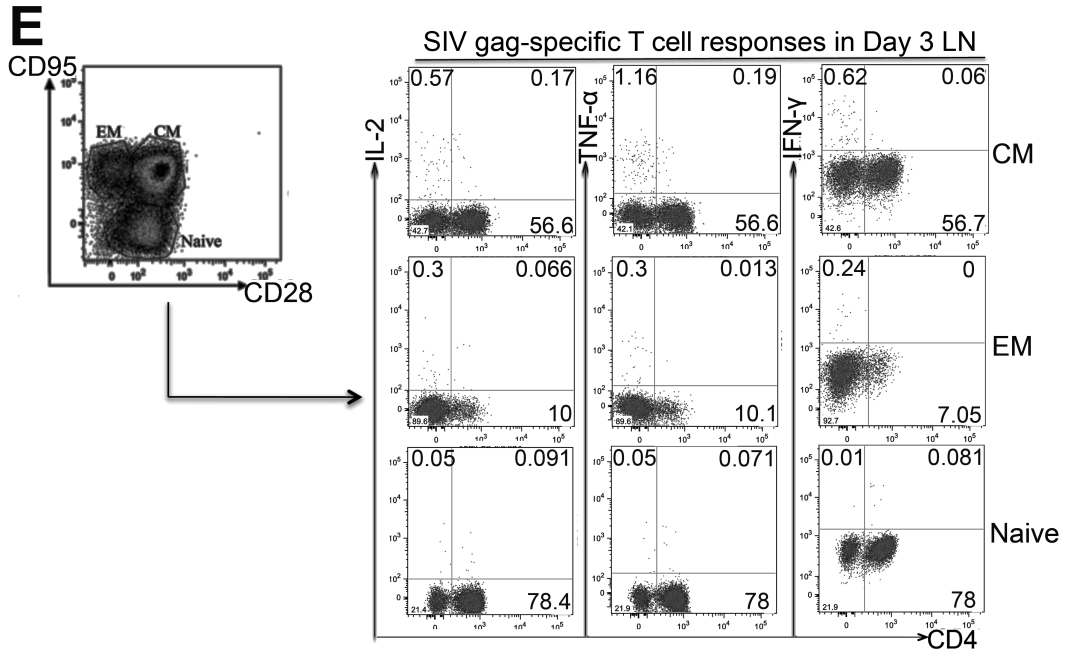


Figure 3.3.2. SIV immunization induced SIV gag-specific T cell responses in LNs.

(A) Representative multi-color flow cytometry data for SIV gag-specific T cell responses in LN draining SIV-gag DNA (Top) or CMV-pp65 (Bottom) on Day 3 post

immunizations. LN mononuclear cells were stimulated with SIV-gag 15-mer overlapping peptide pools and stained intracellularly for IL-2, TNF- α and IFN- γ . The SIV-specific cytokine-producing CD4⁺ and CD8⁺ T cells were determined by multi-color flow cytometry. (B) The percent of SIV gag-specific cytokine-producing CD4⁺ and CD8⁺ T cells in Day3 LN draining SIV gag and CMV pp65 immunization. (C, D) Summary for the percent of SIV gag-specific cytokine-producing CD8⁺ (C) and CD4⁺ (D) T cells in LNs on indicated days post immunization. (E) SIV gag-specific responses in LN were primarily mediated by central memory T cells. Macaque LN mononuclear cells were separated into naïve, central memory (CM) and effector memory (EM) cells according to CD28 and CD95 expressions. The individual cytokine expressions in these populations are shown. (F) Polyfunctionality of SIV-specific CD4⁺ and CD8⁺ T cell responses. The percent of CD4⁺ and CD8⁺ T cells able to produce 1, 2, and 3 cytokines is shown.

3.3. CMV immunization induced CMV pp65-specific T cell responses in LN

We evaluated the CMV-specific T cell responses in a LN draining the CMV-pp65 immunization. Unfortunately, due to cell viability and limited available cells, this was the only day we were able to analyze. LN mononuclear cells on day 5 post immunization were in vitro stimulated with a CMV-pp65 peptide pool, and the pp65-specific cytokine-producing T cells were assessed by multi-color flow cytometry. Compared to the LN draining SIV-gag immunization (Top panels, Figure 3.3.3A), the LN draining CMV-pp65 immunization demonstrated significantly higher frequencies of IL-2-producing CD4⁺ and CD8⁺ T cells in response to pp65 peptides (Bottom panels, Figure 3.3.3A). The

comparison for the percent of IL-2, TNF- α and IFN- γ -producing CD4⁺ and CD8⁺ T cells in Day 5 LN draining SIV or CMV immunization is shown in Figure 3.3.3B.

Figure 3.3.3

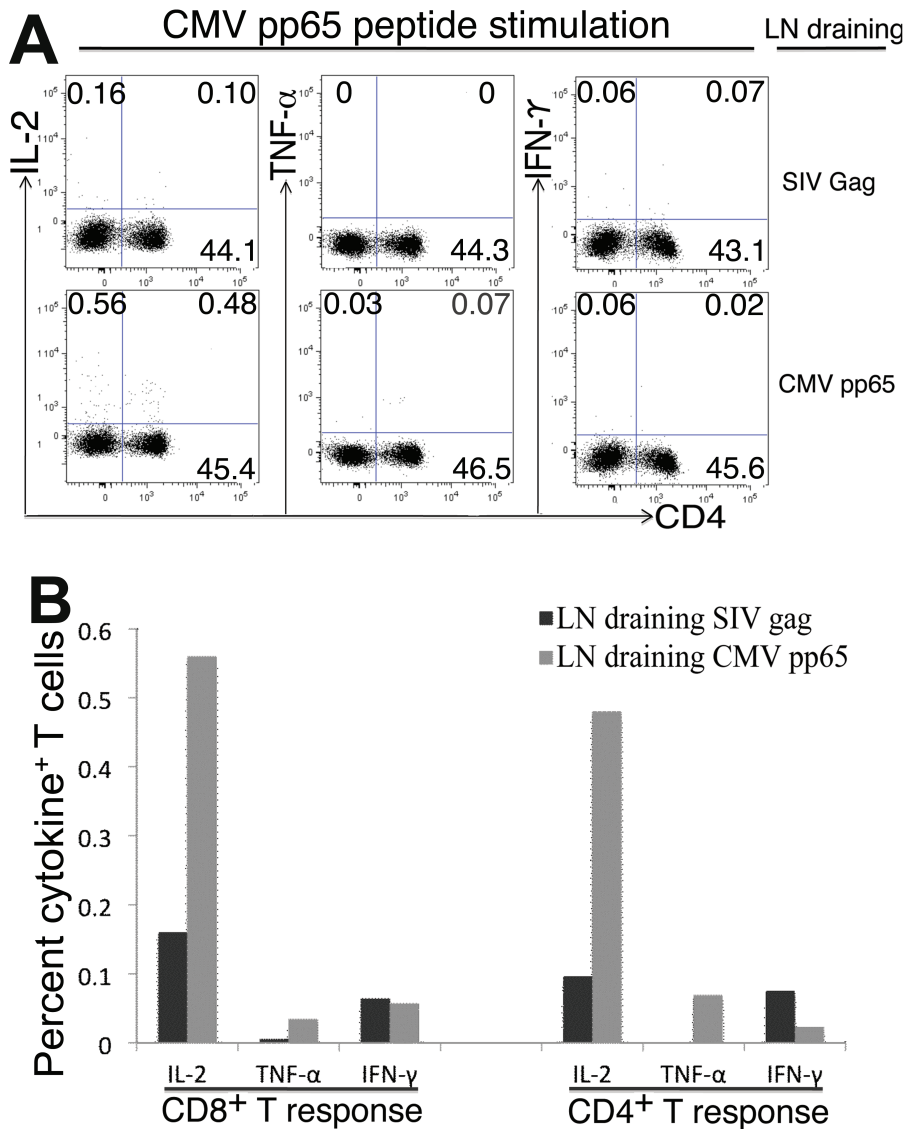


Figure 3.3.3. CMV immunization induced CMV pp65-specific T cell responses in Day 5 LN. (A) Multi-color flow cytometry data for CMV pp65-specific T cell responses in LN draining SIV-gag DNA (Top) or CMV-pp65 (Bottom) on Day 5 post

immunizations. LN mononuclear cells were stimulated with CMV-pp65 15-mer overlapping peptide pools and stained intracellularly for IL-2, TNF- α and IFN- γ . (B) The percent of individual cytokine-producing CD4⁺ and CD8⁺ T cells compared between LNs draining SIV gag and CMV pp65 stimulated with CMV pp65 peptide pool.

PART 4

SIV- and CMV-specific T cell responses in peripheral blood

4.1. Staining for the SIV gag-specific tetramer p11C in peripheral blood

After analyzing the immunization-induced local immunologic responses in draining LNs, we next sought to determine the systemic immune response in the peripheral blood of immunized macaques. PBMC were collected from macaques every two to four weeks throughout the experiment. We first determined the percent of circulating SIV-gag tetramer specific CD8⁺ T cells in PBMCs in all three macaques. PBMC were stained with SIV-gag p11c-MamuA*01 tetramer together with other lineage and viability markers. As shown, tetramer⁺ T cells were exclusively CD8⁺, demonstrating the specificity of SIV tetramer (Figure 3.4.1A). The kinetics for the number of tetramer⁺ SIV-specific CD8⁺ T cells in three animals are summarized in Figure 3.4.1B. Before SIV infection, no tetramer positive cells were detected. This was followed by potent increases after infection. The introduction of ART led to a dramatic decline of tetramer⁺ CD8⁺ T cells to lower levels through the end of experiment. During ART, some transient increases were observed for one animal (FH40), which was consistent with the transient

viral load blips seen in the same animal. After immunization, no significant increases were observed for the number of SIV-specific CD8⁺ T cells in the periphery of the macaques (Figure 3.4.1B).

Figure 3.4.1

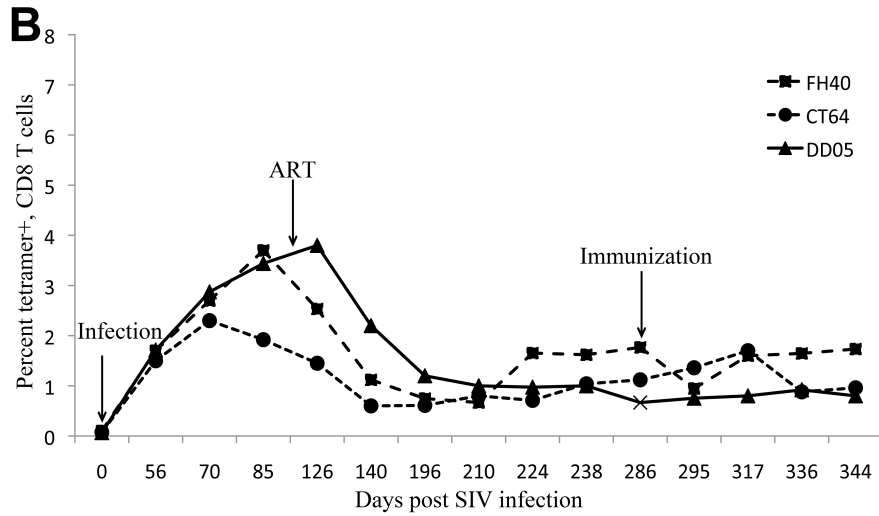
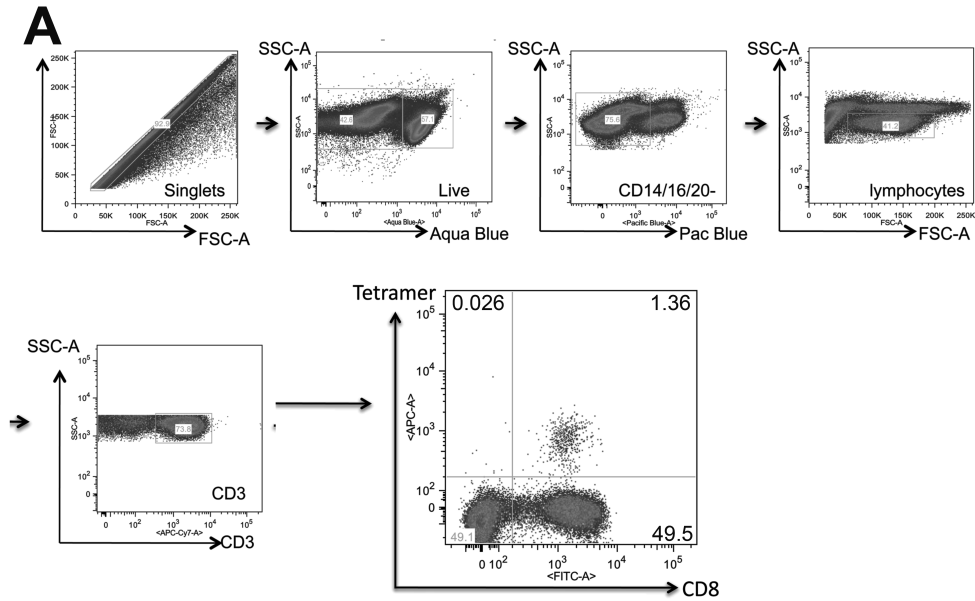


Figure 3.4.1. SIV gag-specific tetramer staining of PBMCs. (A) Representative tetramer staining and analysis for one PBMC sample from one macaque. Single, live and CD3⁺ T lymphocytes were gated and the staining for tetramer and CD8 are shown. (B) Kinetics of tetramer⁺ SIV-specific CD8⁺ T cells in the studied rhesus macaques throughout the experiment. The results are expressed as percent of tetramer⁺ CD8⁺ T cells on indicated days post SIV infection for each animal.

4.2. SIV- and CMV-specific T cell responses in PBMCs

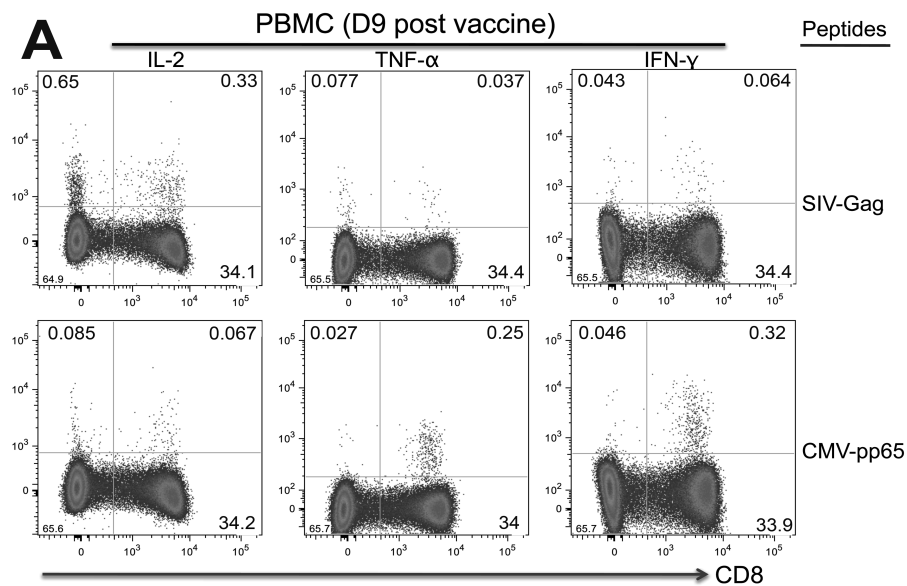
We next determined the systemic response to immunization with SIV gag and CMV pp65 by analyzing peptide stimulated T cell responses. Macaque PBMCs collected before and after immunizations were in vitro stimulated with SIV-gag or CMV-pp65 peptide pools and cytokine-producing T cells were measured by multi-color cytometry. Representative multi-color flow cytometry histograms for SIV- and CMV-specific T cell responses (IL-2, TNF- α and IFN- γ) in PBMC at day 9 post immunization are shown (Figure 3.4.2A). CMV-specific responses were primarily mediated by CD8⁺ T cells producing the effector cytokines IFN- γ and TNF- α , whereas both CD4⁺ and CD8⁺, SIV-specific T cells were induced primarily producing IL-2 with very little IFN- γ and TNF- α on day 9 (Figure 3.4.2A).

The kinetics of immunization-induced SIV- and CMV-specific CD4⁺ and CD8⁺ T cell responses are summarized in Figure 3.4.2B. For the CD4⁺ T cell response, a transient SIV gag-specific, IL-2 producing response was observed at day 9 post immunization, that was not found on day 50 post immunization, whereas no significant CMV-specific CD4⁺

T cell responses were detected at all studied time points. The gag-specific CD4⁺ T cells produced very little TNF- α and IFN- γ (Figure 3.4.2 B, Top).

Similar to SIV gag-specific CD4⁺ T cell responses, the SIV-specific CD8⁺ T cell response was transient and extinguished quickly with only an increase on day 9 post-immunization. The SIV-specific CD8⁺ T cells primarily produced IL-2 with very little TNF- α and IFN- γ productions. In contrast, CMV-specific CD8⁺ T cell responses demonstrated a long-lived, polyfunctional profile. A potent increase in CMV-specific CD8⁺ T cells was observed at day 3 post immunization with an average of 0.8% of CD8⁺ T cells able to produce TNF- α or IFN- γ . The TNF- α and IFN- γ -producing CMV-specific response was sustained until day 50 post-immunization. CD8⁺ and CD4⁺ T cell CMV-specific T cells produced very little IL-2 (Figure 3.4.2B)

Figure 3.4.2



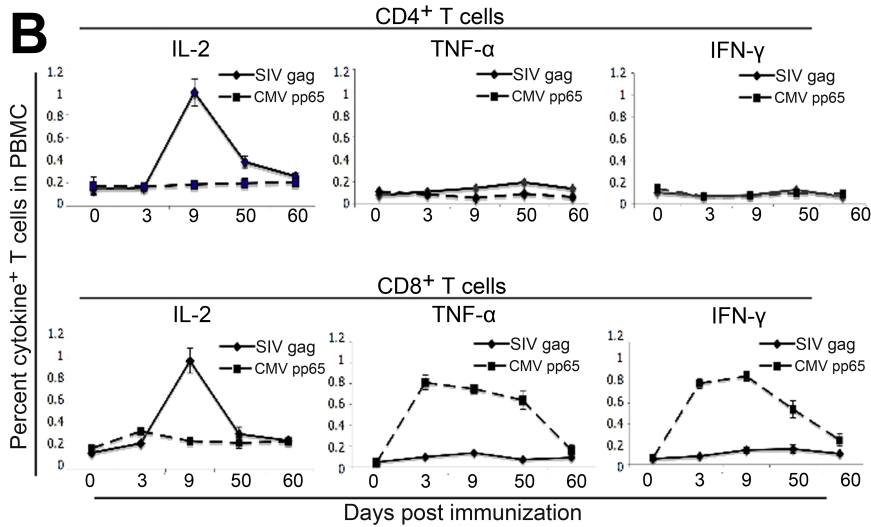


Figure 3.4.2. SIV- and CMV-specific T cell responses in peripheral blood during

immunization. (A) Representative multi-color flow cytometry data for SIV- and CMV-

specific T cell responses. PBMC before and after immunization were stimulated with SIV gag (Top) or CMV pp65 (Bottom) peptide pools and stained intracellularly for IL-2,

TNF- α , and IFN- γ . The percentages of cytokine-producing CD8⁺ T cells were determined

by multi-color flow cytometry. (B) The average of all animals for the percent of CD4⁺

and CD8⁺ T cells able to produce individual cytokines (IL-2, TNF- α or IFN- γ) is shown

and compared between SIV- and CMV-specific responses.

4.3. T cell subsets that mediate SIV- and CMV-specific T cell responses in PBMC

We also assessed the T cell subsets (naïve, central memory, and effector memory)

that mediated the antigen-specific T-cell responses in PBMC post immunization. Using

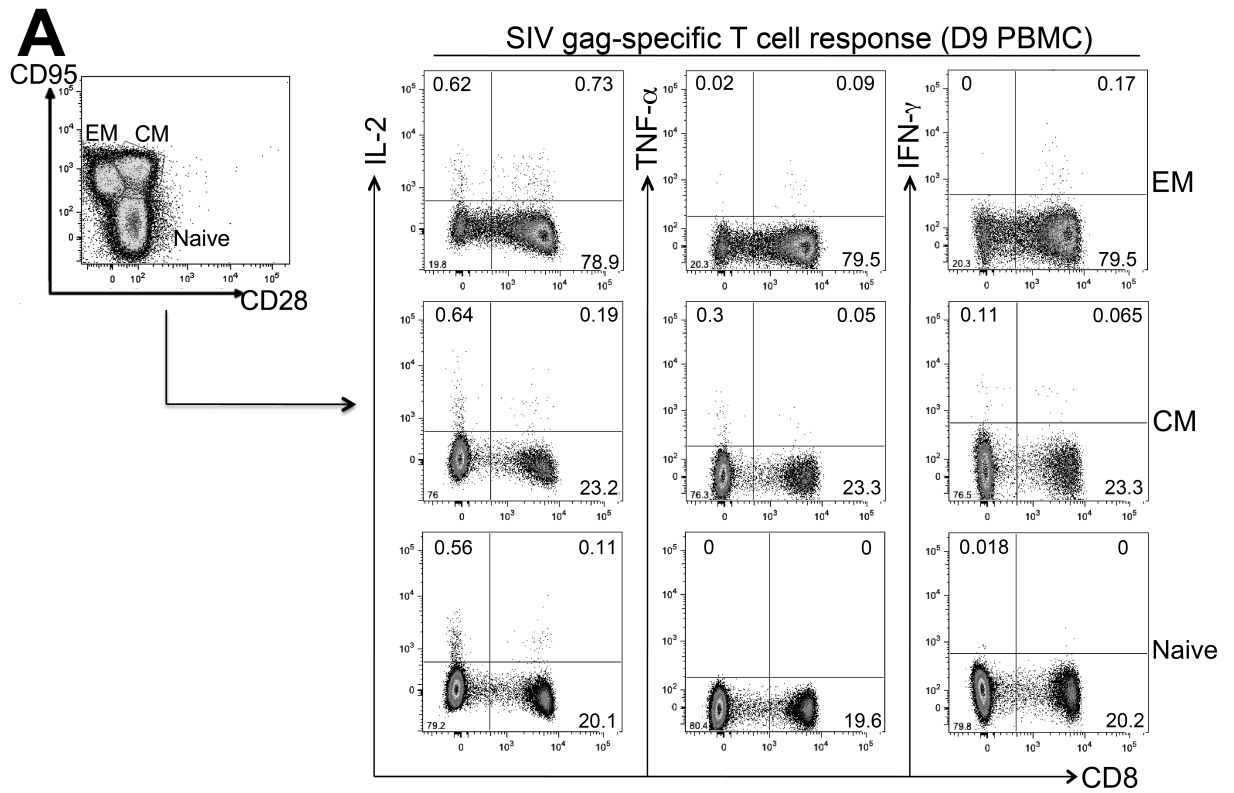
the Day 9 PBMC when both SIV- and CMV-specific T cell responses were highest, we

analyzed individual cytokine production in naïve, central memory, and effector memory

T cell subsets. The SIV-specific IL-2-producing response was mediated by effector (EM)

and central memory (CM) T cells (Figure 3.4.3A). In contrast, the CMV-specific TNF- α and IFN- γ - responses were primarily mediated by effector memory (EM) T cells (Figure 3.4.3B).

Figure 3.4.3



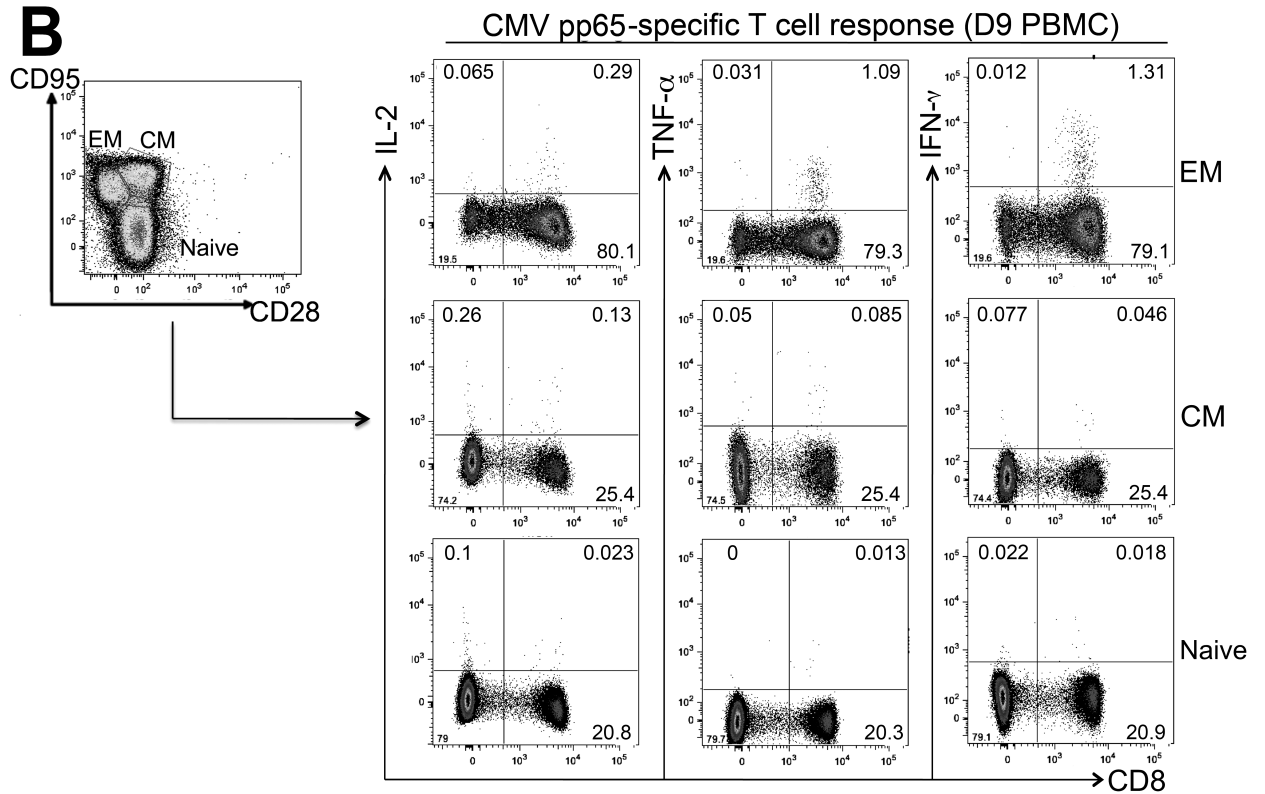


Figure 3.4.3. Analysis of T cell subsets mediating the antigen-specific T cell

responses in PBMC. Macaque PBMC were separated into naïve, central memory (CM) and effector memory (EM) cells according to CD28 and CD95 expression. The individual cytokine expression in these populations for SIV-specific (A) and CMV-specific (B) responses are shown.

4.4. Increased PD-1 expression in SIV-specific CD8⁺ T cells

T cell exhaustion often occurs and prevents optimal viral control during chronic infection¹⁷. An important role of the PD-1:PD-L1 pathway has been reported for CD8⁺ T cell exhaustion during chronic viral infection¹¹. We co-stained the PBMC for the SIV-

gag tetramer p11c and PD-1 to determine the differential expression of PD-1 on SIV-specific and total CD8⁺ T cells. As seen in a representative histogram of tetramer-PD-1 staining, PD-1 was significantly upregulated in tetramer⁺ CD8⁺ T cells compared to tetramer negative CD8⁺ T cells (Figure 3.4.4A). The PD1 expression in PBMC during the course of SIV infection and immunization was analyzed, and it was observed that PD1 expression on SIV-specific CD8⁺ T cells was significantly higher than tetramer negative CD8⁺ T cells for all time points studied, suggesting exhaustion of SIV-specific CD8⁺ T cells, which was not reversed by ART and immunization (Figure 3.4.4B).

Figure 3.4.4

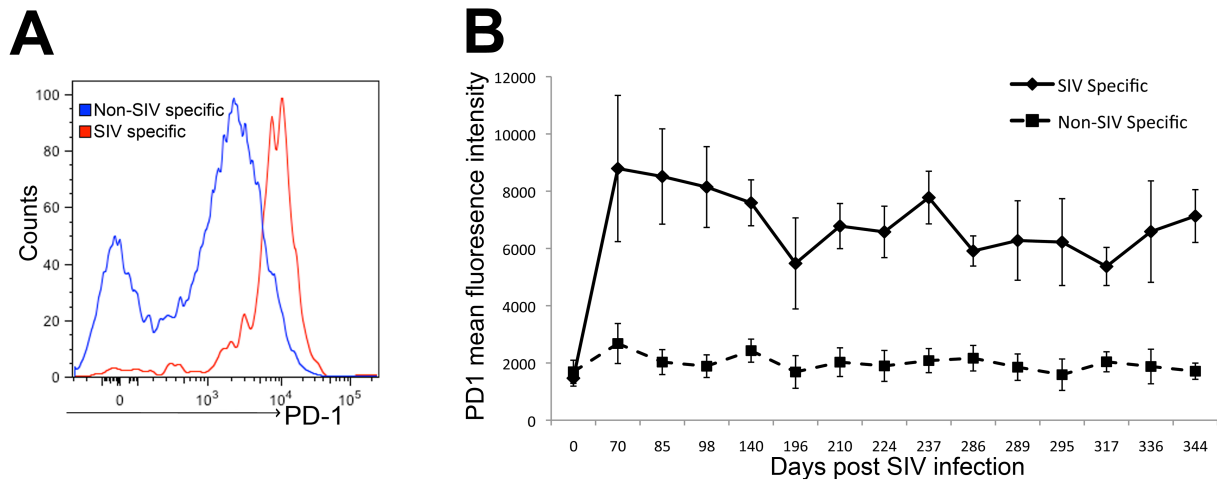


Figure 3.4.4. PD-1 expression on SIV-specific and non-SIV specific CD8⁺ T cells in macaque PBMC. (A) Representative flow cytometry histogram data for mean fluorescence intensity (MFI) of PD-1 staining on SIV-specific and tetramer negative CD8⁺ T cells. (B) PD-1 expression on SIV-specific and tetramer negative CD8⁺ T cells in

PBMC collected on indicated days post SIV infection. The results are shown as the average of mean fluorescence intensity (MFI) of PD1 staining for two monkeys.

PART 5

Immunization induced virologic responses in draining LNs

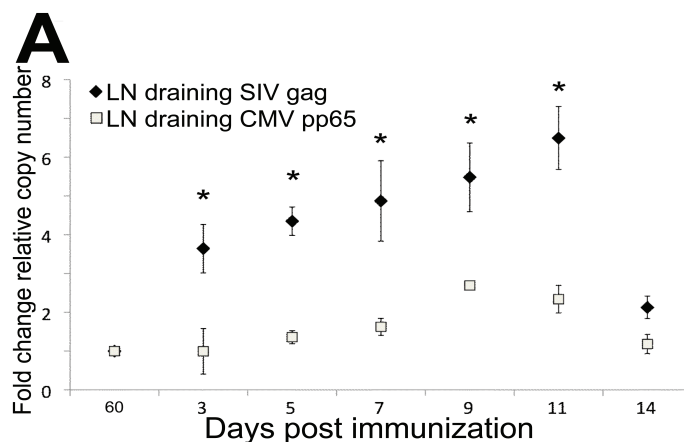
5.1. SIV immunization selectively stimulates transient activation of viral replication in draining LNs

ART can efficiently control viral replication but fails to eradicate virus due to the establishment of long-lived reservoirs¹²⁶. It has been demonstrated that a low level of viral production continues even in the setting of the most potent antiviral regimens^{43, 70, 122}, but the source of this virus is not believed to be from new infection of cells^{32, 84, 122}.

Total RNA from LNs draining either SIV gag or CMV pp65 immunizations was analyzed by real-time PCR for multiple SIV viral RNA transcripts, including unspliced, doubly spliced, and singly-spliced RNA¹¹⁹. As preimmunization control LNs could not be obtained, LN biopsies from both sides, draining either gag or pp65, from each macaque were collected. No significant differences in levels of unspliced and spliced RNA transcripts were observed comparing right to left side for each animal. Thus, this time point was chosen for comparisons. Copy numbers of SIV RNA were normalized to GAPDH RNA and the results for each time point following immunization are shown as fold change relative to RNA collected at 60 days post immunization (Figure 3.5A, B, C).

The levels of doubly spliced SIV RNA viral transcripts in LNs draining the SIV gag immunization demonstrated a 3.6-fold increase at day 3, which was sustained until day 11 post immunization when a 6.5-fold increase was observed. Minor increases in copies of doubly spliced SIV RNA were observed in LNs draining the CMV pp65 immunization (Figure 3.5A). The levels of singly spliced viral transcripts demonstrated a similar increase in RNA in LNs draining the SIV gag immunization with an 11.8-fold increase at day 11 (Figure 3.5B). Unspliced SIV RNA levels were determined and an 8.3-fold increase was observed at day 7 (Figure 3.5C). The percent of CD4⁺ T cells in LN mononuclear cells did not differ between LNs draining SIV gag and CMV pp65 immunizations, suggesting that the increase in viral replication was not due to differential CD4⁺ T cell migration to the LN draining gag antigen, although we cannot rule out that a larger fraction of cells that trafficked to the gag LN were infected and expressing viral RNA transcripts.

Figure 3.5



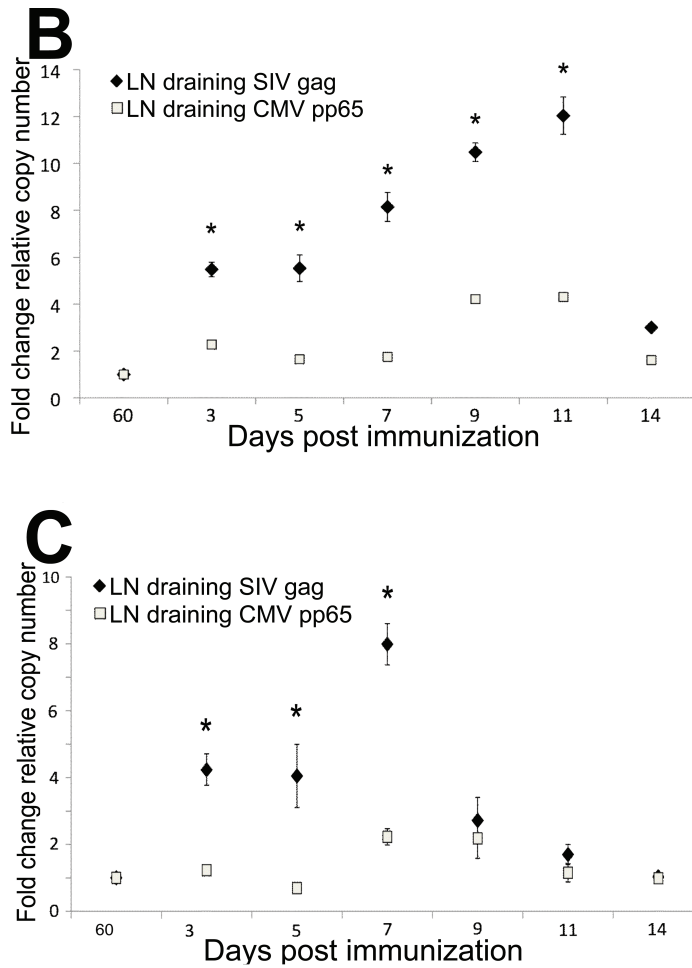


Figure 3.5 Quantitation of multiple SIV viral RNAs in LNs. Total RNA extracted from LN mononuclear cells draining either SIV gag or CMV pp65 were subject to real-time PCR with specific primers and probes for quantitation of SIV doubly spliced RNA (A), singly spliced RNA (B) and unspliced RNA (C). Copy numbers of SIV RNAs were normalized to macaque GAPDH and the results for each time point following immunization were shown as fold change relative to RNA collected on day 60 post immunization as a no vaccine control. The comparisons were made between LNs from the same animal at the same time on indicated days with a * indicating a $p < 0.05$).

5.2. No increases viral load were observed in peripheral blood post immunizations

A transient activation of SIV viral replication in LNs draining the SIV gag immunization was observed beginning at day 3 that returned to baseline at day 14-post immunization (Figure 3.5). We next sought to determine whether this transient viral replication could be seen in peripheral blood. The peripheral viral loads prior to and after antigen immunization in all animals are shown in Table 5. Except for a transient viral blip in one animal at day 31 post-immunization (270 copies/ μ l) that returned to undetectable 3 weeks later, no increase in viral loads that could be attributed to SIV gag immunization induced viral replication was detected. The data suggests that in the setting of potent anti-viral suppression, SIV antigen immunization activated viral replication was transient and restricted to draining LNs without spread to the periphery.

Table 5. Viral loads in periphery during antigen immunization

Days PI	238	290	317	336	344
Animal					
CT64	<125	<125(D0*)	<125(D27*)	<125	<125
DD05	<125	<125(D0*)	<125(D27*)	<125	<125
FH40	<125	<125(D4*)	270(D31*)	<125	<125

* Days post immunization

Discussion

A biphasic destruction of CD4⁺ T cells is observed in HIV infection, with a massive loss of CD4⁺ T cells during early infection and a subsequent progressive loss during the chronic stage of infection^{47, 125}. Retroviral suppression by ART results in an increase in peripheral CD4⁺ T cell counts and functional reconstitution of CD4⁺ T cell responses to many common antigens^{8, 83}, but HIV-specific CD4⁺ T cell responses remain deficient¹²⁹. To identify potential mechanisms for this persistent loss of HIV-specific CD4 immune function, we studied ART-treated, chronic SIV and CMV infected rhesus macaques¹³⁶, and observed 1) that both SIV and CMV antigen immunizations could induce immune activation and antigen-specific T cell responses in draining LNs. 2) In peripheral blood, the CMV-specific response induced by immunization was potent and sustained, whereas the SIV-specific response was transient and extinguished quickly. 3) I demonstrate that SIV antigen immunization induces transient activation of SIV viral replication that was not observed with CMV pp65 immunization. The data is consistent with a hypothesis that SIV antigens induce viral replication that leads to depletion or dysfunction of antigen specific cells leading to a reduction in the strength and longevity of the response.

Pathogenic SIVmac 251 infection of rhesus macaques has been well described as one of the preferred experimental models for studying HIV pathogenesis⁹². The macaques in our study were inoculated with SIVmac251 and all established primary infection with typical viral replication dynamics^{68, 114, 143}. Similar to the observations for HIV-1 infected

individuals where ART can sustain suppression of plasma viremia to undetectable levels¹²⁰, all macaques in our study demonstrated uniform responsiveness to ART (Figure 3.2A). We observed that the macaques also exhibited a sustained increase in peripheral CD4⁺ T cell numbers above 500 cells/ μ l after PMPA and D4T treatment, demonstrating the establishment of suppressed viral replication and recovered CD4⁺ T cell counts observed in HIV-infected subjects treated with ART during chronic infection.

Expression of particular MHC class I alleles are associated with different rates of disease progression in humans and macaques^{23, 24, 81, 110, 171}. In our study, all three rhesus macaques expressed Mamu A*01 to control for an MHC effect on viral immune responses and disease progression as well as to aid in the measurement of immune responses. The advantage of selecting one haplotype is that the animals studied will have similar responses to infections and immunizations, which enhances our ability to make conclusions with a small number of animals. Indeed, limited variation in viral replication dynamics were observed across monkeys after primary infection and prior to initiation of ART (Figure 3.2A).

The ability of therapeutic immunization to elicit SIV-specific immune responses in SIV-infected rhesus macaques during ART and its virologic benefits have been investigated primarily in peripheral blood using DNA immunization^{57, 150, 157, 173} and other systems^{42, 69, 96}. After release from ART, variable immunologic and virologic benefits were reported from no control¹⁷³, temporal control⁶⁹, to long-lasting control⁹⁶.

In this study, using CMV infection as a control, we investigated the immunologic and virologic consequences of immunization with SIV antigen in chronic SIV and CMV co-infected, ART treated rhesus macaques. All the comparisons between LNs at each time point were from the same animal collected at the same time, thus allowing me to use each animal as its own control and reach a statistically significant result. Also, my study focused on the local responses in draining LN where antigen-specific T cell responses and viral replication occurs, rather than systemic responses in peripheral blood. In addition, I chose a range of days post-immunization covering the early activation of memory T cells and generation of effector responses. My data show that both SIV and CMV antigen immunizations induced significant immune activation and antigen-specific T cell responses in draining LNs and peripheral blood. The CMV-specific response was sustained in PBMC with a rapid onset of cytokine producing CD8⁺ T cells 3 days post immunization, which was maintained at day 50-post immunization. However, the immunization induced SIV-specific T cell response was transient and extinguished quickly in blood. Of interest, a study that repeatedly immunized macaques with long-standing ART-treated SIVmac251 infection induced stronger SIV-specific CD4⁺ and CD8⁺ T cell responses in blood¹⁴⁸. This study used the MVA viral vector and delivered three immunizations, whereas in our study, only single gag DNA immunization was used.

In addition, SIV- and CMV-specific T cell responses in PBMC demonstrated different functional profiles. Both CD4⁺ and CD8⁺ SIV-specific T cells were transiently detected and produced mainly IL-2 with very little TNF- α and IFN- γ , whereas the CMV-

specific responses, primarily mediated by CD8⁺ T cells, exhibited high TNF- α and IFN- γ production but very little IL-2. It was previously shown that the response patterns of antigen-specific CD4⁺ T cells change as they mature. More mature cells infrequently produced IL-2 and produced effector cytokines and chemokines ²⁵. My results demonstrate that the SIV-specific IL-2-producing T cell responses were central and effector memory T cells, whereas the CMV-specific TNF- α and IFN- γ -producing responses were almost exclusively mediated by the more mature effector memory T cells. My results suggests that in peripheral blood, in addition to being present for 50 days, the CMV-specific T cells demonstrated a more polyfunctional profile and more mature phenotype, which is in agreement with a study showing that in HIV and CMV co-infected individuals, the CMV-specific CD4⁺ T cells were able to produce increased β -chemokines (MIP-1 α and MIP-1 β), which was associated with a more mature phenotype and decreased susceptibility to HIV infection, compared to HIV-specific CD4⁺ T cells ²⁶. The in vivo relevance of such IL-2 producing SIV-specific T cells is not clear in terms of protective cellular immunity and susceptibility to viral infection as potential new targets, but my data suggests that they are rapidly lost.

The progressive loss of effector function in the setting of chronic viral infections has been associated with the upregulation of programmed death 1 (PD1)¹¹. In HIV/SIV infection, increased expression of PD1 is shown to correlate with CD8⁺ T cell exhaustion and failure of immune control, which is reversible both in vitro and in vivo with PD1 blockade ^{40, 147, 152}. In our study, we monitored PD1 expression on SIV-specific and

tetramer negative CD8⁺ T cells and found that PD1 expression on SIV-specific CD8⁺ T cells were significantly higher than on tetramer negative CD8⁺ T cells throughout the entire experiment, which could not be markedly reversed by ART and immunization (Figure 3.4.4). This strongly indicated the selective exhaustion for SIV-specific CD8⁺ T cells during acute and chronic SIV infection, consistent with a study by Petrovas C et al

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HIV-specific CD4⁺ memory T cells are preferentially infected by HIV, carrying approximately 2- to 5-fold more viral DNA than total memory cells⁴⁶. It has also been shown that even during the most potent regimens of retroviral suppression, the presence of virus could be measured by some ultrasensitive assays^{43, 70, 122}. These findings raise the concern that immunizations that induced HIV/SIV-specific T cells may accelerate viral replication even in the setting of potent ART. In support of this hypothesis, a vaccine that induced poor CD8⁺ T cell and neutralizing antibody responses led to enhanced SIV replication after challenge that was associated with enhanced CD4⁺ T cell proliferation¹⁴². We hypothesized that during an HIV-specific response, activation of HIV-specific CD4⁺ T cells, which bear higher amounts of latent virus, results in activation of viral replication in the LN leading to suppression of CD4⁺ T cells and the HIV-specific responses through multiple mechanisms, whereas a non-HIV antigen-specific response activates little viral replication allowing efficient expansion of the response.

We used a naturally CMV infected, SIVmac251 infected and ART treated during chronic infection rhesus macaque model to study the effects of immune stimulation using HIV gag and CMV pp65 antigens. The studies concentrated on the early immune and viral responses in draining lymphoid organs. Both antigen immunizations were able to induce transient immune activation and generated antigen-specific T cell responses in draining LNs, but the SIV gag immunization also induced significant viral replication. The SIV response extinguished quickly in peripheral blood, while the CMV response was sustained. Our data suggests that SIV antigens, as part of a vaccine or potentially as part of the normal immune response to the virus, leads to T cell stimulation with subsequent viral replication that leads to an impairment in the immune response to the virus. It is possible that this mechanism is responsible for the observation that in progressing HIV infected subjects; CD4-specific responses to HIV antigens are lost early in infection and are difficult to restore or induce.

CHAPTER 4

Thesis conclusions and future research directions

Thesis conclusions and implications

In the first part of my thesis, using an in vitro system modeling the interaction between HIV Env and CD4⁺ T cells, I further characterize that Env protein, when delivered as a vaccine and expressed on APCs or non-APCs, present on free circulating virions, or as expressed by HIV-infected cells could induce inhibition of antigen-stimulated CD4⁺ T cell activation and proliferation, which is dependent of CD4 binding and independent of coreceptor binding. The inhibition is not characterized as apoptosis, anergy or cell death. These Env exposed CD4⁺ T cells can phosphorylate key signaling mediators including MAPK, AKT, and STAT5a in a normal fashion but fail to fully up-regulate activation markers such as CD69, HLA-DR, and CD25. Treg cells are not functionally involved in Env-induced inhibition of CD4⁺ T cell activation and exposure of PBMC to Env neither changes the frequency nor the suppressive activity of Tregs. Using this data, I established a hypothesis, which suggests a mechanism for the generation of poor responses to HIV vaccines that contain Env. This hypothesis has implications for the development of novel HIV Env vaccine candidates with improved immunogenicity.

In the second part of my thesis, I employed an in vivo model system to further analyze HIV suppression of CD4⁺ T cell antigen-specific immune responses. I demonstrate that in draining lymph nodes of chronically SIV and CMV co-infected

rhesus macaques during ART, both SIV- and CMV-immunization could induce immune activation and antigen-specific memory T cell responses. The SIV-specific T cell responses are polyfunctional with the production of multiple cytokines (IL-2, TNF- α and IFN- γ) and primarily mediated by central memory T cells. I further track the immunization-induced antigen-specific systemic responses in peripheral blood, and demonstrate that SIV-specific T cell responses were transient and extinguished quickly in blood. The SIV-specific responses in blood primarily produced IL-2 and was mediated by multiple T cell subsets, effector and central memory. In contrast, the CMV-specific T cell responses were sustained in PBMC with a rapid onset of CD8⁺ T cells able to produce high levels of TNF- α and IFN- γ with very little IL-2, and was mainly mediated by more mature effector memory cells. The SIV immunization led to a transient activation of viral replication in draining LNs, without spread to the periphery. My results may have some potentially important implications for HIV infection and immunization in humans. First, the immune activation associated with HIV immunization during ART stimulates transient viral replication in draining LNs. Second, the transient viral replication leads to impairment of the HIV-specific response by multiple mechanisms resulting in an impaired HIV-specific systemic T cell response. In applying these findings to the induction of HIV-specific immune responses in the setting of any viral replication, I propose that the generation of HIV-specific CD4⁺ T cell responses leads to elevated local viral replication with resulting loss of HIV-specific immune responses. These findings demonstrate a potential mechanism for the failure of progressing subjects to develop an HIV-specific CD4⁺ T cell response.

Ongoing experiments and future directions

In addition to the mononuclear cells prepared from the LN for immunologic and virologic analysis as described in my thesis, fixed LN tissues were also prepared for in situ analysis. We are currently working with Dr. Ashley Haase's laboratory to study these fixed LN tissues using technologies such as in situ hybridization, in situ immunohistochemistry including tetramer specific staining, and quantitative image analysis to visualize in situ viral replication induced by SIV gag immunization and the host's cellular immune responses in LN tissues draining SIV and CMV immunization. In particular, we are interested in further dissecting the mechanisms for the inhibition of HIV-specific CD4⁺ T cell immunity, including; 1) SIV- and CMV-induced apoptosis; 2) regulation of Treg cells and other potential inhibitory molecules (i.e. PD-1); and 3) viral replication associated with cell type, activation status, and antigen-specificity.

Future interesting research would include: 1) investigation of TCR or cytokine signaling cascades by Env or virus in CD4⁺ T cells to further dissect the molecular mechanisms involved in inhibition of CD4⁺ T cell activation; 2) SIV rhesus macaque studies with a larger number of monkeys. Preliminary results obtained in my thesis demonstrate that in chronic SIV infection during ART, immunization-induced systemic SIV-specific T cell responses compared to CMV-specific T cell responses differ in being more transient and have a different profile of cytokine expression, high IL-2, but very little TNF- α and IFN- γ production. Interesting questions could be asked about this difference in a larger SIV macaque study. Are these IL-2 producing CD4⁺ and CD8⁺ T

cell responses protective? Or, in contrast, do these IL-2 producing activated T cells serve as targets for viral infection or activation of latent virus? Finally, a number of questions were raised by my pilot macaque study and data worthy of investigation in a larger trial was identified. First, for time points studied, I would have multiple LN samples from different animals draining SIV and CMV immunization. This would give statistical significance for a group instead of significance for the comparison in the same animal. Second, according to our preliminary results, I focus studies on Day 3 and Day 5 when the immunization-induced local responses are highest. I would also add a time point prior to day 3 and reduce the number of time points after day 5. This will allow more LN samples for analysis without greatly increasing the number of animals needed. Moreover, if more cryopreserved LN mononuclear cells are available, cells will be analyzed for infection, apoptosis and exhaustion associated with cell type and antigen-specific immune activation. Cells could also be sorted into different subsets, such as DCs, macrophages, and CD4⁺ and CD8⁺ T cells and these cells will be analyzed for the extent of infection and viral integration and effector functions. This may allow for the identification of the cell types that are replicating virus in response to SIV immunization. I would also add studies to determine if the loss of the SIV-specific response is an active process. This could be done by analyzing apoptosis markers on tetramer positive cells. The addition of a CMV specific tetramer would allow a more complete analysis and I would work with Dr. Peter Barry at UC Davis to try and develop one. The analysis of additional LN samples will also allow a more thorough investigation of the T cell responses to antigen.

CHAPTER 5

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