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► To cite this version:

Anne-Sophie Liovat, Béatrice Jacquelin, Mickaël Ploquin, Françoise Barré-Sinoussi, Michaela Müller-Trutwin. African Non Human Primates Infected by SIV - Why Dont they Get Sick? Lessons from Studies on the Early Phase of Non-Pathogenic SIV Infection. *Current HIV Research*, 2009, 7 (1), pp.39-50. 10.2174/157016209787048546 . pasteur-01962242

HAL Id: pasteur-01962242

<https://hal-pasteur.archives-ouvertes.fr/pasteur-01962242>

Submitted on 20 Dec 2018

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African non human primates infected by SIV - why don't they get sick?
Lessons from studies on the early phase of non-pathogenic SIV infection

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Running title: Non-pathogenic SIV infection in African non human primates

Keywords: SIV, African non human primate, immune activation, acute infection, inflammation.

Abstract

African non human primates are natural hosts of SIV. The infection is generally non-pathogenic despite high steady-state levels of plasma viral RNA that in HIV-1 and SIVmac infections are associated with progression towards AIDS. The viral loads in the gut also are as high as in pathogenic HIV-1/SIVmac infections; but replication levels are lower in peripheral lymph nodes of chronically infected African green monkeys. There is a transient loss of CD4⁺ T cells in the blood in SIVagm and SIVsm infections and an early dramatic and more persistent decrease in the gut. Although SIV in natural hosts is thus cytopathic, the continuous viral replication is not associated with immunopathology. T CD4⁺ cells in blood, lymph nodes and gut manifest no or little increase of cell-death by apoptosis. The lymph node and gut architecture is not disrupted. The most striking difference between non-pathogenic SIV and pathogenic HIV-1/SIVmac infections is the lack of chronic T cell activation. Several studies are currently in progress to determine which factors are involved in the maintenance of the low activation level in the non-pathogenic SIV infections. There are two ways in which this could be achieved: (i) a lack of immune activation induction or (ii) an active downregulation of the immune activation. The arguments in favor of each of these two possible ways of immune activation control will be discussed in view of the most recent data in the literature. A particular focus is put on data on the innate immune system and the timing of induction of immunosuppressive mediators during the early phase of SIV infection.

Introduction

African non human primates (NHP) are natural hosts of the simian immunodeficiency virus (SIV). The first species identified as natural carriers of SIVs were sooty mangabeys (SMs), African green monkeys (AGMs), mandrills and chimpanzees. Today, SIV viruses from more than 30 distinct African NHP species have been reported [1]. The infection of African natural hosts (such as AGMs, SMs and mandrills) is generally non-pathogenic. It is striking that the same virus found in African NHPs can be pathogenic in other species. The African natural hosts of SIV are indeed the animal reservoirs of HIV-1 and HIV-2 [1-3]. Furthermore, SIVmac derives from SIVsm and induces the acquired immunodeficiency syndrome (AIDS) in macaques of Asian origin. Nevertheless, wild strains of SIVsm do not always replicate at high titers in macaques [4]; *in vivo* passages are often needed for adaptation to the heterologous host [5]. There are signs that HIV-1 has also adapted specifically to humans [6]. Additionally, not all macaque species show a similar susceptibility to SIV infection and AIDS. For instance, SIVagm infection of rhesus macaques does not result in persistent high viremia and AIDS, whereas pigtailed (Pt) macaques infected with a particular strain of SIVagm (ver90) or with SIV l'hoest or SIVsun succumb to AIDS [7, 8]. The pathogenicity also depends on the viral strain. For example, SIVagm.ver155 does not induce AIDS in Pt macaques in contrast to SIVagm.ver90 [9]. These are examples that illustrate that the non-pathogenic issue of the infection is determined by an interplay of both host and virus determinants.

The mechanisms of protection against AIDS have been studied most frequently in AGMs and SMs. Other less commonly studied models include mandrills, chimpanzees and l'Hoest monkeys. Animal models allow to thoroughly study the very early virus-host interactions in blood and tissues. Studies on the acute phase in tissues have been mostly addressed in AGMs, but also in SMs. Both in HIV-1/SIVmac infections, events in the early phase of infection are predictive of the further outcome [10, 11]. Such studies on the early events in African NHP models might therefore reveal important information on the mechanisms that control disease progression. Here we will review the data of the literature on the most relevant differences between non-pathogenic and pathogenic HIV/SIV infections during the chronic phase, and whenever available during the acute phase of infection.

Do natural hosts of SIV display more efficient anti-viral adaptive immune responses?

African NHPs show higher number of CD8⁺ cells in peripheral blood compared to humans and macaques [12]. It has been suggested that this could confer an advantage to the natural hosts. SIV-specific cell-mediated immune responses in the blood of SIVsm-infected SMs are indeed predominantly mediated by CD8⁺ T cells [13, 14]. There is in addition evidence of selective immune pressure on SIVsm [15]. There is, however, no evidence for stronger adaptive immune responses in natural hosts of SIV compared to those animals or individuals that progress to AIDS. Many SIV specific T cells in SMs and AGMs are not polyfunctional with respect to the production of IFN- γ , TNF- α , IL-2 and/or MIP-1 β [13, 16]. IL-2⁺ were detected in only a very small percentage of SIV-specific T cells [13, 14]. The magnitude and breadth of the SIV-specific T cell responses were either similar or weaker than in HIV-1/SIVmac infections [13, 14, 16]. Moreover, the level of viral replication correlated with neither their magnitude nor their breadth [13, 14]. SMs and AGMs display lower levels of granzyme B positive T cells than SIVmac infected macaques, suggesting an attenuated effector function [4, 16]. Only minor changes in the levels of plasma viremia are observed in *in vivo* CD8⁺ cell-depleted SMs [17]. This suggests that CD8⁺ T cells and NK cells exert limited, if any, direct immune control on SIV replication in SIV-infected SMs. The data on cell-mediated adaptive responses in natural hosts therefore indicate that the absence of AIDS is independent of a robust T cell response to the virus (Tab.1).

Seroconversion in SIV infected African NHPs generally occurs within 4 to 5 weeks, a similar time frame to SIV-infected macaques [18, 19]. However, chronically infected SMs seem to display one log lower titers of total antibodies compared to macaques [20, 21]. Surprisingly, the levels of proliferating B cells in germinal centers of peripheral LNs from AGMs and SMs is higher than in macaques during early infection [4, 22]. The biological meaning of this observation is not clear. It might reflect an early B cell dysfunction in pathogenic HIV-1/SIVmac infections [23]. Natural hosts develop neutralizing antibodies [24, 25]. It is not clear whether their titers are similar to HIV-1/SIVmac infection because they vary considerably depending on the virus and cells used for the assay [26]. They are probably not more efficient than in humans and macaques since the neutralizing activity against the autologous virus isolates seems to be weak [24]. In addition, a selective lack of anti-p27 antibody responses has been detected in many, but not all, natural hosts [19, 20, 24]. Whether this is related to the non-pathogenic outcome is unclear.

Altogether, these studies show that there is no evidence of stronger anti-viral antibody and T cell responses in natural hosts as compared to HIV-1/SIVmac infections. They are either similar or weaker. Weaker responses might protect the host against immunopathological damage on the expense of a lack of virus replication control.

Is the natural host of SIV able to restrict viral replication or does the virus target distinct cells or tissues?

Is the virus replicating less than in pathogenic HIV-1/SIVmac infections?

There is no better control of viral replication than in HIV-1/SIVmac infections. The high mutation rate of SIVagm was the first indirect demonstration of an uncontrolled replication *in vivo* [27]. This has been confirmed in SMs that display plasma viremia levels similar to macaques during the chronic phase [28]. Subsequently, it has been shown in AGMs that plasma viremia can be very high during acute infection [19]. Similar to SIVmac infection in macaques, the peak of replication in natural hosts occurs between days 6 and 14 post-infection (p.i.), with plasma RNA copy numbers between 2×10^4 and 10^9 /ml [29]. The peak levels were independent of the initial blood CD4⁺ T cell counts [30, 31]. After this early peak in viral replication, there is a sharp decline of RNA copy levels to variable set-point levels [19]. Naturally infected AGMs and SMs display a considerably wide range of viral loads (VLs) with many animals displaying high viremia levels [32-34]. Similarly high plasma VLs are observed in mandrills and chimpanzees [18, 35, 36]. These high VLs might be important for an efficient spreading of the virus among the host population.

The lack of disease progression in the context of high VL led to the conclusion that a high viremia is not sufficient to induce AIDS (Tab.1). In humans also, high viremia is not always associated with disease progression. For example, a few HIV-1 infected long term non progressors (LTNP) have remained asymptomatic while presenting high viremia [37]. Recently, a case of a patient infected with HIV-1 for 20 years who has experienced CD4⁺ T cell depletion in spite of maintaining undetectable viral loads, has been described [38]. Moreover, HIV-2 infected individuals develop AIDS in the presence of low plasma VLs [39]. Aside from high viremia, additional factors must therefore play a role in the induction of immune dysfunction.

Is the virus replicating in distinct tissues?

Differences in the outcome of the infection could be associated with a targeting to distinct tissues by the virus or by sparing specific organs such as thymus or intestine. The mucosal immune system plays a central role in both transmission of HIV infection and AIDS pathogenesis. The major site of HIV-1/SIVmac replication is indeed the gut. This is correlated with the fact that the gut associated lymphoid tissues (GALT) contain the most elevated frequency of activated CD4⁺ T cells and the highest number of viral target cells. Similar to SIVmac, however, SIVagm also replicates at high levels in the gut, both in the acute and the chronic phase of infection [40]. The infected cells are localized in the lamina propria and Peyer's patches [33, 41]. Moreover, the distribution of the virus in other tissues (e.g. thymus, cerebrospinal fluid, lungs...) is similar to HIV-1/SIVmac [32, 33, 40].

Only in LNs major differences have been reported between non-pathogenic and pathogenic HIV-1/SIVmac infections (Tab.2). While the number of productively infected cells in LNs during acute SIVagm infection is similar to SIVmac infection, this number is very low during the chronic phase (Tab.2) [9, 22, 42, 43]. Similar low numbers in LNs have also been reported for two SIVcpz-infected chimpanzees [43, 44]. For SMs, there is also no difference with SIVmac infection in the acute phase [4]. During the chronic phase, the numbers of RNA positive cells in the SM's LNs were equivalent to slow/ intermediate progressor macaques [28]. It is not excluded, that SMs show a slightly higher VL than AGMs in the chronic phase. So far, only 2 animals have been reported in the literature and more SMs need to be analyzed to answer to this question.

The proviral load in AGMs is also very low in LNs during the chronic phase of infection as compared to blood and gut [19, 32, 42]. This contrasts with humans infected with HIV-1 and macaques infected with SIVmac, which both show 5 to 10 times more extensive proviral burden in the LNs than in the peripheral blood mononuclear cells (PBMC) [45]. In line with the RNA and DNA loads, the average titer of infectious virus in AGM LNs is also low (21 TCID₅₀ per 10⁶ LN mononuclear cells) [42].

Productively infected cells in AGMs, SMs and chimpanzees are predominantly located in the T cell zone and little or no trapping of SIV by follicular dendritic cells is observed [28, 33, 42, 44]. The underlying mechanisms of this lack of virus trapping are not elucidated. It could be related to differences in the immune activation (IA) status in LNs of natural hosts (see below). The lack of virus trapping most likely results in higher levels of free virus particles in the blood. In that case, it would need less infected cells to achieve similar levels of viremia than

in pathogenic HIV-1/SIVmac infections, which could explain the lower proviral burden in AGMs as compared to macaques.

Altogether, these data demonstrate similar VL levels between non-pathogenic SIV and pathogenic HIV-1/SIVmac infections in both blood and intestine. Nevertheless, a lower viral burden, at least for AGMs, is observed in the peripheral LNs in the chronic phase. The presence of less virus in the inductive sites of immune responses might have a beneficial impact, but this remains so far an open question that needs further investigation.

Which cells are targeted?

Some hypotheses put forward that the non-pathogenic outcome of SIVagm infection in AGMs is associated with a lower number of CD4⁺ T cells in this species. CD4⁺ T cell counts vary according to age and individuals. In young AGMs (1.4 to 3.5 years) the blood CD4⁺ T cell counts (between 800-2900, mean 1400 CD4⁺ T cells/ μ l) are often not lower than in SMs and macaques, with a mean of 30% (<17-44%) of CD4⁺ T cells in PBMC. Adult AGMs (3.6-9.8 years old) have significantly reduced CD4⁺ T cell counts (<250-1300, mean 700 CD4⁺/ μ l) with 24% (<10-36%) of CD4⁺ T cells in PBMC [12, 31, 46]. Indeed, some adult AGMs display less than 300 CD4⁺ T cells/ μ l in blood [30, 31]. In humans, a lymphocyte count such as this would be associated with opportunistic infections and AIDS. Whether AGMs with low CD4⁺ T cell counts have been selected during evolution and whether this confers an advantage to the host during SIV infection is still an interesting matter of debate and needs further explorations.

Another interesting observation concerns the levels of CD4⁺CCR5⁺ T cells in natural hosts. These levels are significantly lower than in humans and non natural hosts for SIV, both in blood and tissues [47]. The frequency of CCR5⁺ plasmacytoid DCs (pDC) is also lower [48]. Like SIVmac, SIV from African NHPs use CCR5 as a major coreceptor [49-51]. There are however few exceptions (Tab.3). For instance, SIVrcm uses CCR2b instead of CCR5. The latter is characterized by a 24bp deletion in red capped mangabeys [50]. SIV viruses most of the time are not CXCR4 tropic. However, some primary SIVagm and SIVsm isolates are able to use CXCR4 in addition to CCR5 for cell entry [30, 52-54]. The ability of SIVs to replicate in human T cell lines without using CXCR4 has suggested the existence of additional simian co-receptors. Bonzo/STLR33 is preferentially used by SIVagm [51] and Bob by SIVsm/SIVmac *in vitro* [55]. Still, CCR5 is the major co-receptor for SIVsm and SIVagm *in vitro* and *in vivo* [50, 51]. A frequent mutation found in CCR5 of AGMs results in a molecule that is still functional for SIVagm but not for HIV-1. This is a sign that SIVagm is adapted

specifically to CCR5 of AGMs. Altogether, SIVs from natural hosts show a similar coreceptor usage to SIVmac.

It is a paradox that SIVsm and SIVagm, which use CCR5 as a major coreceptor, induce high VLs in the context of these low levels of CD4⁺CCR5⁺ T cells. Thus, the levels of CCR5 expression may still be sufficient to allow infection *in vivo*. In line with this, in macaques' intestine, the majority of infected cells have a CCR5⁻ phenotype [56]. The meaning of the lower frequencies of CCR5⁺ cells in the protection against pathogenesis needs to be clarified. It could play a role in distinct cell trafficking or be related to a distinct differentiation stage of target cells in natural hosts. It has been hypothesized that in natural SIV hosts, expression of CCR5 is restricted to CD4⁺ T cells that are in a more advanced differentiation stage, thereby limiting virus replication to a subset of CD4⁺ T cells that are anyway destined for activation induced cell death regardless of their SIV infection status [57]. In line with this, the bulk of SIVagm replication is sustained by very short lived cells, as suggested by mathematical modeling [58]. The exact phenotype of infected CD4⁺ T cells has not been explored so far. It is known however that, *in vivo*, most productively infected CD4⁺ T cells in the gut display a memory phenotype [41]. They rarely express Ki67 at the peak [41, 59]. SIVagm also replicates in antigen-presenting cells, such as macrophages (Fig.1) [33]. No data are yet available on the susceptibility of dendritic cells (DC) from African NHP to SIV infection. AGM Monocyte DC efficiently transmit SIVagm to T cells through DC-SIGN *in vitro* [60]. The lower VL in AGM LN is thus not related to a lack of DC-T cell transmission.

In the state of knowledge, except of the levels of CD4⁺ CCR5⁺ T cells that differ between natural hosts and macaques, there are no major differences reported between non-pathogenic and pathogenic infection regarding the target cells. However the nature of the infected CD4⁺ cell subpopulations (CD4⁺ T cells and DCs) in natural hosts is only poorly defined. More studies on the phenotype of the infected cells are urgently needed. Moreover, it is important to understand why viral replication in LNs during the chronic phase is lower in non-pathogenic SIVagm infection. It could be a consequence of less target cells. Like HIV/SIVmac, SIV isolates from natural hosts replicate preferentially in CD4⁺ T cells when they are in an activated state and generally do not grow in unstimulated PBMCs *in vitro* [61-63]. The level of viral replication *in vivo* correlates with the frequency of activated T CD4⁺ cells in naturally infected SMs [64]. It is likely that the availability of activated T CD4⁺ cells rather than immune control of viral replication is the main determinant of set-point VL during SIV infection in natural hosts [64]. Thus, the lower level of viral replication in LNs could be

related to a lower level of chronically activated CD4⁺ T cells as compared to the gut in non-pathogenic SIV infection.

Does the virus cause less immunological damage in the natural hosts?

Early depletions of peripheral CD4⁺ T Cells

In humans and macaques, sustained high VL is most often associated with CD4⁺ T cell declines. In contrast, chronic SIV infections of African NHPs are generally characterized by preservation of healthy blood CD4⁺ T cell counts despite high levels of viremia. It is of note that in acute infection, AGMs, SMs and mandrills demonstrate a significant drop in peripheral CD4⁺ T cell counts (Tab.1) [18, 30, 65, 66]. This drop is more pronounced in the animals that show a higher CD4⁺ T cell count at day 0 [30]. It is preferentially observed in AGMs displaying high peak viremia. At the end of the acute infection, the CD4⁺ T cell values in the blood rebound rapidly to set-point values that are often lower although still close to healthy levels [41, 66]. In the chronic phase, SIV infection in SMs is associated sometimes, but not always, with a moderate decline in CD4⁺ T cells [67]. Although SIV infection can thus have a moderate impact on CD4⁺ T cell counts, the latter remain generally close to normal levels in blood.

A transient decrease in CD4⁺ T cells has been occasionally observed in the LNs of AGMs and mandrills during acute infection, however not in all monkeys that display a depletion in the blood [18, 41, 65]. Again, the normal levels are readily recovered by the end of the acute phase.

Depletion of Mucosal CD4⁺ T Cells

In the GALT, the early kinetics of CD4⁺ T cell depletion are also the same than for pathogenic infections (Tab.1). Thus, an early dramatic depletion of mucosal CD4⁺ T cells is observed in SMs and AGMs [41, 68]. A rapid drastic depletion of GALT-based memory CD4⁺CCR5⁺ T cells is a characteristic feature in HIV-1-infected individuals and SIV-infected rhesus macaques [69, 70]. The cell population that was the most affected in natural hosts corresponds also to memory cells, being defined as CD3⁺CD4⁺CD28⁺CD95⁺ cells [41, 68].

However, while disease progression in macaques is associated with a further decline in mucosal CD4⁺ T cells in the chronic phase of infection, this was not the case in SMs, at least until day 175 p.i. [68]. AGMs even recovered partially their CD4⁺ cells in gut, the levels being 35-75% of baseline by day 410 p.i. [41]. A better recovery of CD4⁺ T cells in AGMs as compared to SMs might be related to their lower VL in the chronic phase of infection as shown in LNs (Tab.2) and blood. Set-point values of plasma VL are indeed 1 log lower for AGMs than SMs, mandrills and chimpanzees for the other species [29]. To test this hypothesis, comparative quantitative analysis of VL in the intestine of these two species needs to be done.

It is intriguing that the peripheral CD4⁺ T cell counts in natural hosts are preserved despite their dramatic loss in the gut. IL-7 is known for playing an important role in homeostatic proliferation of CD4⁺ memory T cells and IL-7 levels are often elevated in T cell-depleting conditions. During acute SIV infection SMs, IL-7 increases are observed in temporal association with the early declines in peripheral CD4⁺ T cell counts. This early transient IL-7 increase is followed by increases in CD4⁺ and CD8⁺ T cell proliferation [66]. Interestingly, the natural hosts, whether infected or not, display signs of higher T cell proliferation in bone marrow and an inherently high rate of peripheral CD4⁺ T cell turnover, raising the question as to whether there is a better compensation for loss of infected cells in natural hosts [71, 72]. Moreover, while bystander cell death of CD8⁺ T cells has been demonstrated in natural hosts (chimpanzees, AGMs, SMs), the latter display no increased apoptosis of CD4⁺ T cells in blood [71, 73]. In AGMs, there is also no increase of apoptosis in LN and gut during acute or chronic infection [22, 41]. In SMs, there is an increase at day 14 p.i. in LN followed by a rapid return to normal levels [4]. There is therefore no chronically increased CD4⁺ T cell apoptosis in natural hosts. It is thus probably easier to maintain homeostasis, since there is no need to compensate a loss of non-infected CD4⁺ T cells dying by bystander cell death. In addition, the nature of the depleted cells might not be the same in pathogenic and non-pathogenic HIV/SIV infections. For instance, SIVmac infection results in depletion of Th17 cells in GALT of macaques, whereas in AGMs and SMs, the Th17 cells are preserved [74] [Cervasi et al., 2008¹, Favre et al., 2008²]

¹ B. Cervasi, J. Brenchley, M. Paiardini, S. Gordon, A. Asher, I. Frank, J. Else, D. Douek, and G. Silvestri. Preferential Loss of Th17 CD4 T Cells in the Gastrointestinal Tract of HIV-infected Individuals but Not SIV-infected Sooty Mangabeys. February 3-6, 2008. Conference on Retroviruses and Opportunistic Infections, Boston, Massachusetts.

² D. Favre, S. Lederer, B. Kanwar, Z. M. Ma, S. Proll, Z. Kasakow, C. Miller, M. Katze, and J. McCune. Primary SIV Infection Causes Rapid Loss of the Balance between TH17 and T Regulatory Cell Populations in

Natural hosts are therefore characterized by early significant CD4⁺ T cell declines in acute infection that persist in the chronic phase in intestine of SMs. Moreover, very low peripheral CD4⁺ T cell counts without any sign of AIDS can be occasionally observed in AGMs and SMs [31, 33, 75]. Studies in natural hosts therefore indicate that losses of CD4⁺ T cells are necessary but not sufficient to cause AIDS. Aside from high viremia and early CD4⁺ T cell losses, additional processes are necessary. In macaques, it is not the degree of early CD4⁺ T cell decline in mucosa, but rather the degree of recovery in the post-acute and early chronic phase that are correlated with the outcome of infection [76, 77]. During antiretroviral therapy, CD4⁺ T cell restoration in GALT is best achieved when inflammation is suppressed [78]. It is crucial to understand why in AGMs and SMs, the CD4⁺ T cells in gut do not further decline in contrast to macaques that progress towards AIDS.

Lack of chronic T cell activation

The most striking difference between non-pathogenic SIV and pathogenic HIV-1/SIVmac infections is the lack of chronic T cell activation in the natural hosts despite the ongoing viral replication (Tab.1). In HIV-1/SIVmac infections, high viremia correlates with IA. IA seems to be a better prediction factor for disease progression than VL [39, 79-82]. Already before seroconversion, CD8⁺ T cell activation levels in HIV-1 infection are predictive of the rate at which CD4⁺ T cells are lost over time [11]. In natural hosts (AGMs, SMs), increases of T cell activation measured by markers such as Ki67 and MHC-II can be observed during acute infection in the blood, especially in animals displaying high viremia at the peak [4, 65]. However, the increase is only transient and there is a rapid and substantial resolution of T cell activation. As a result, during the chronic phase, the levels are close to baseline levels [65, 67, 71, 83]. There is nevertheless a marked inter-individual variability with respect to CD8⁺ T cell activation levels and some SMs display non-significant increases. The moderate decreases in CD4⁺ T cell counts in SMs are observed in those animals with higher CD8⁺ T cell activation [71]. Thus even in natural hosts, SIV infection impacts the immune system, but only to a weak extent without leading to generalized IA and impaired homeostasis.

In LNs, signs of transient T cell activation are also observed during acute infection [4, 22]. However, the levels of CD4⁺ and CD8⁺ T cell proliferation are not substantially increased during the chronic phase [22, 71]. In line with this, natural hosts do not exhibit LN follicular or paracortical hyperplasia, nor infiltrations of CD8⁺ T cells into LN germinal centers [19,

33]. There is no evidence of increased collagen deposition in post-acute and chronic phase of infection resulting in no tissue damage through fibrosis [4]. Tissular architecture in LNs is indeed conserved [19, 42].

In the gut, the IA seems to be lower in natural hosts as well. Thus, the levels of plasma lipopolysaccharide (LPS) were increased in a few SIV-infected African NHPs during acute infection, but not in all of them, and never during chronic infection, by opposition to HIV-1/SIVmac infections [41, 68, 84]. Plasma LPS is a marker of IA and is the result of translocation of microbial products to the systemic circulation [84]. The mechanisms underlying the disruption of the mucosal barrier in HIV-infected individuals and SIVmac-infected rhesus macaques are not clear. Th17 cells play a role in the protection against microbial infections [85, 86]. Consequently, depletion of Th17 cells during HIV-1/SIVmac infection contributes to dissemination of pathogenic bacteria [87]. Other mechanisms could play a role as well in the disruption of the mucosal barrier in HIV-1/SIVmac infections. Thus, the early SIVmac infection phase is associated with a massive apoptosis of intestinal epithelial cells that might represent the underlying mechanism of the regenerative enteropathy [88]. Interestingly, no apoptosis was observed in the gut of acutely infected SIVagm-infected AGMs [41]. The reasons why natural hosts display lower levels of apoptosis and whether this is simply the consequence of lower IA or due to other factors, still needs to be established.

Despite high levels of VLs, natural hosts therefore show a rapid control of T cell activation and normal or only moderate IA during chronic infection compared to humans and macaques. The studies in natural hosts have contributed to the present point of view that IA is the driving force for CD4⁺ T cell depletion and AIDS, while viral replication is only indirectly associated with disease progression, probably through increasing IA. To understand how HIV-1/SIVmac induce chronic IA and how natural hosts protect themselves against it, presents a major and the most important challenge today.

Potential mechanisms of immune activation control

Several studies are currently in progress to determine which factors are involved in the maintenance of the low activation level in the non-pathogenic models. Two ways of achieving it are possible: (i) a lack of IA induction and/or (ii) an active downregulation of it (Fig.2).

Less induction of activation?

The first sentinels that sense signals of infection are the cells of the innate immune system. Microbes and viruses are sensed through pattern-recognition receptors (PRR), such as Toll

like receptors (TLR). These receptors bind pathogen-associated-molecular-pattern molecules (PAMP) [89]. PAMPs induce the secretion of pro-inflammatory cytokines including TNF- α , IL-6, IL-12 and type I interferon (IFN-I) by binding to TLRs. IFN-I is induced through TLR3, 7, 8 and 9. TLR3 recognizes double-stranded viral RNA and is expressed on myeloid DCs (mDCs) [90, 91]. TLR7 and 8 recognize single-stranded viral RNA [92, 93], and TLR9 is stimulated by unmethylated DNA rich in cytosine-guanosine motifs from bacteria and viruses [94]. TLR7 and 9 are expressed on pDCs and TLR8 on monocytes and mDCs [91, 95]. HIV-1 signals directly through TLR7 which results in the production of TNF- α and IFN- α [96]. A signaling by HIV-1 through TLR9 is not excluded [93, 96].

The considerable inter-individual variation in humans and macaques regarding IA levels might be related in part to individual differences at the level of TLR expressions or functions. The expression of some TLRs is altered during HIV-1/SIVmac infections. TLR7 and 8 mRNA levels were significantly increased in subjects with chronic HIV-1 infection and that of TLR3, TLR7 and TLR8 in LNs of SIV-infected macaques [97-99]. On the other hand, TLR9 expression is decreased in LNs in the early phase of SIVmac infection and HIV-1 GP120 seems to diminish TLR9 signaling [98, 100, 101]. During HIV/SIVmac infections, pro-inflammatory cytokines might also be induced indirectly by exposing innate immune cells to co-infecting agents [102]. For instance, as a result of bacterial translocation, PAMPs, such as the TLR4 ligand LPS, could be involved in further driving general IA [84]. Still, there is not enough data so far to directly link differences at the level of TLR expressions and/or stimulations with IA levels.

We explored the inflammatory cytokine profiles in non pathogenic SIV infection. We measured cytokine expressions in three distinct body compartments: blood, LNs (induction site) and bronchioalveolar lavages (mucosa associated lymphoid tissue, effector site) of acutely SIVagm-infected AGMs (Tab.4). Strikingly, AGMs displayed only weak increases of IFN- γ , and no significant increases of TNF- α , IL-6, IL-12, MIP-1 α and MIP-1 β during acute infection [65, 103]. To confirm this weak inflammatory profile in SIVagm infection, we have quantified the expression of *t-bet* and *gata3* in acute infection. These transcription factors are essential for the induction of Th1 and Th2 responses, respectively. In the PBMCs of both AGMs and macaques, *gata3* expression was upregulated [104]. However, only macaques presented an upregulation of *t-bet*, whereas AGMs did not. This is in agreement to the weak induction of pro-inflammatory cytokines. In conclusion, SIVagm-infection in AGMs is characterized by a lack of inflammatory responses resulting in a Th2/anti-inflammatory

cytokine balance. A tendency towards Th2 cells has also been reported in chronically infected SMs [71, 105], although SIV-specific CD8⁺ T cell preferentially secrete Th1 cytokines [13]. A lack in early inflammatory cytokine production is likely to play an essential role in the low T cell activation profiles (Fig. 2).

Little is known about TLR expression and signaling in the non-pathogenic models. There might be an intrinsic lower response to SIV in African NHPs. Indeed, stimulation of TLR9 from pDCs of uninfected AGMs results in IFN- α production at levels significantly lower than those in healthy human donors and macaques [103, 106]. Nonetheless, under certain circumstances, TLR9 stimulation on AGM pDCs results in high IFN- α production. This is the case during acute SIVagm infection, where pDCs from blood and LNs display strong IFN- α production after *ex vivo* stimulation with a TLR9 ligand [103]. IFN- α levels in plasma are indeed increased during acute SIVagm infection. The levels were positively correlated with plasma VL [103]. However, plasma IFN- α was only detectable around the peak of viremia and the absolute levels were significantly lower than in SIVmac infection [103]. In chronic HIV-1/SIVmac infection plasma IFN- α levels and ISG expressions increase with progression towards AIDS. IFN- α does not seem to play an important role in HIV/SIV replication containment, and chronic production might even be deleterious for the immune system [107, 108]. Thus little or no chronic IFN- α production could be of benefit by inducing less immunopathological damage.

Studies in SMs also show substantially reduced levels of innate immune system activation [48]. Thus, *in vitro* stimulated pDC from non infected SMs produce markedly less IFN- α than macaque pDC. However, after stimulation with inactivated SIV or a TLR9 agonist, a similar production of TNF- α and IL-12 could be observed for SMs and macaques, suggesting that while the pathway that requires IRF7 is compromised, the NF κ B-dependent signaling pathway is intact in SM's pDC [48]. *In vivo*, the expression of inflammatory cytokines such as TNF- α , IL-6 and IL-12 mRNAs was significantly lower in chronically infected SMs than in macaques [48]. The reasons why natural hosts display a lack of pro-inflammatory cytokine production in response to SIV infection seem to be complex and far from being completely understood.

Rapid induction of immunosuppressive mechanisms?

HIV-1/SIVmac infections are associated with increases in immunosuppressive mediators, such as that of regulatory T cells (Treg) in tissues, or of TGF- β 1⁺, IDO⁺ and PD-1⁺ cells in

blood or LNs [109-112]. This is most likely a response by the host to try to control inflammation and immune-mediated pathological damage. However, these responses do not appear to be capable of limiting the massive hyperactivation [109].

Treg play a central role in controlling deleterious immune responses such as inflammation [113]. They also facilitate early protective responses to local viral infection by allowing a timely entry of immune cells into infected tissues [114]. The timing of their induction and their localization in HIV/SIV infections might be a critical determinant for the outcome of infection. Treg represent a heterogeneous fraction of T cells that include IL-10 secreting Tr1 cells, TGF- β 1 producing Th3 cells, as well as naturally CD25⁺ Treg and adaptive IL-10 and TGF- β 1 secreting CD25⁺ Treg. CD25⁺ Treg have been detected in HIV/SIVmac infection, but their implication is not clear and contradictory data have been published. In some studies, CD25⁺ Treg seem to be correlated with a lower VL, a diminished IA and a slower progression towards AIDS; while in others, they are associated with suppression of virus-specific immune responses and disease progression [109, 113, 115].

In SIVagm-infected AGMs, some animals showed early increases of IL-10, TGF- β 1 and FoxP3 expression [65]. TGF- β 1 is able to induce CD25⁺ FoxP3⁺ Treg *in vivo*. On the other hand, IL-6, which is increased in macaques but not in AGMs, can prevent CD4⁺CD25⁺ Treg functions [116]. It is thus possible that, due to the particular cytokine environment during early SIVagm infection, DCs are conditioned to preferentially induce regulatory T cells rather than Th1 cells. However, it has yet to be demonstrated if the FoxP3⁺ cells detected in AGMs correspond to Treg cells, since FoxP3 and CD25 are also upregulated upon activation [117]. One could nonetheless hypothesize that a more rapid immunosuppressive response in AGMs could abrogate immune hyperactivation, resulting in a far more benign disease outcome in comparison to that in macaques. The Treg cell response in SIV-infected macaques may be too late to counterbalance and prevent the immunopathological consequences of sustained IA, and too early and untimely with respect to immune control, as it down-regulates important effector T cell responses before immune control is achieved [65, 109]. In AGMs, the Treg cell response would also inhibit antigen-specific T cell responses, resulting in viral replication levels comparable to those in SIV-infected macaques, but they would be early enough to prevent the immunopathological effects of sustained IA.

CD25⁺ Treg have also been described in SIVsm infection in SMs [118, 119]. It is not excluded that their suppressive function is more pronounced in SMs as compared to macaques

[118]. There has been, however, no correlation detected so far between Treg levels and IA resolution [4, 118, 119].

In SMs, the resolution of early T cell activation in LNs was rather associated with a more rapid increase in PD-1 expression compared to macaques [4]. Whether this is related to differences among the natural host species, to the lack of specific Treg markers or to the lack of studies of PD-1 in acute SIVagm infection, must be further investigated. PD-1 is known to be associated with suppression of T cell responses [120, 121]. The rapid increase in PD-1 expression in SMs was observed particularly in the CD4⁺ (CD8⁺) T cell fraction. It paralleled the decreases in T cell proliferation and effector functions. Notably, there was a strong positive temporal association of PD-1 levels with IA resolution in LNs of SMs, whereas the increase in the level of IA over time was not affected by the level of PD-1 expression in macaques.

The expression of PD-1 is enhanced by TGF- β 1 *in vitro* [110]. It is not excluded that TGF- β 1 induction during the first days p.i. is stronger in natural hosts [65]. It could also be that in pathogenic HIV/SIVmac infections there is a diminished sensitivity to TGF- β 1 signaling in T cells. The TGF- β 1 major pathway is mediated by the Smad proteins. Smad3 and 4 are activators of the pathway, whereas Smad7 is one inhibitor of this pathway. We quantified *smad7* mRNA levels and detected an up-regulation of *smad7* transcripts in PBMC of acutely SIV-infected macaques. In contrast, the AGMs were characterized by a longer lasting up-regulation of *smad4* with no or a less detectable increase in *smad7* expression [104]. These results are in favor of the hypothesis of a reduced responsiveness to TGF- β 1 during primary SIVmac infection, which could result in a delay of TGF- β 1 dependent Treg and PD-1 induction and/or of T cell proliferation inhibition.

Collectively, these data support the hypothesis of a more rapid induction of immunosuppressive mediators in non-pathogenic SIV infection contributing to the rapid resolution of acute IA. Several mechanisms have been proposed for IA control in natural host. They all relate to a very early control of inflammation. They are not mutually exclusive and might even partially depend on each other. Further investigations are required to evaluate the impact of each of them on IA and AIDS protection.

Conclusion

In conclusion, non-pathogenic SIV infections are characterized by a delicate balance between high viral replication and a moderate host immune response. Natural hosts of SIV thus resemble those few LTNPs who maintain their CD4⁺ T cell counts despite high viremia levels [37]. Both have in common a very weak or absent chronic T cell activation (Fig.2). Current efforts are employed to understand the underlying mechanisms. Recent data in NHPs point toward important differences at the level of the innate immune system and the timing of induction of immunosuppressive mediators during acute infection.

In the future, more studies are needed to identify the responsible key elements in the regulation of T cell activation in the natural hosts. In particular, more detailed analysis on the phenotype of the infected CD4⁺ T cells, their localization and numbers in the intestine, and the role of the low numbers in LNs in the preservation of the host immune system will be of importance. In parallel, more studies are needed to examine the role of the mediators of the innate immune system in the weak inflammatory responses. These could be antigen-presenting cells, in particular pDC, but also, NK cells, $\gamma\delta$ T cells and neutrophils. Moreover, non immune cells such as epithelial cells, might play a role in the early cytokine profiles. Finally, phenotypical and functional studies of cells expressing molecules associated with immunosuppression (FoxP3, IDO, PD-1) should be further developed.

These studies will be crucial to delineate the mechanistic basis for the IA control during SIV infection in natural host species. They pave the way for a series of in vivo experiments that target the early inflammatory pathways. On the long term, this could contribute to the development of new prophylactic and therapeutic strategies that aim to protect against AIDS.

ACKNOWLEDGEMENTS

Our studies were supported by the French Agency for AIDS Research (ANRS), Sidaction and Institut Pasteur. Anne-Sophie Liovat was a recipient of a scholarship from the French “Ministère de l’Enseignement Supérieur et de la Recherche” and from the Denis Diderot University (Paris 7). We thank Matthew Marx for reviewing the English as well as Désirée Kunkel and Asier Saez-Cirion for helpful discussions.

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Fig 1.

Fig 1. Cellular targets of SIV in non pathogenic infection.

SIVs from African NHPs replicate predominantly in activated/memory T CD4⁺ lymphocytes. Macrophages can also be infected. No data are available about the susceptibility of African NHP DCs to SIV infection (dotted arrow). DC-SIGN expressed on AGM MDDC can efficiently transfer SIVagm to T cells and this enhances T cell infection.

Fig 2

Fig 2: Hypothetic model on the mechanisms responsible for the distinct outcomes of HIV-1/SIV infections.

Phase	Pathogenic infection		Non-pathogenic infection	
	acute	chronic	acute	chronic
Mutation rate in vivo	+	+	+	+
Viral load	+	+	+	+
Depletion of CD4⁺ T cells				
blood	+	+	+	-
peripheral LN	+	+	+	-
gut	+	+	+	+
Adaptative antiviral responses	+	+	+	+
T cell activation				
blood	+	+	+	-
peripheral LN	+	+	+	-
gut	+	+	+	-
CD4⁺ T cell apoptosis	+	+	-	-
Immune activation profile	pro-inflammatory		anti-inflammatory	

Tab. 1: Main characteristics of non-pathogenic SIV versus pathogenic HIV-1/SIV infections during the acute and chronic phase.

Parameters are presented as qualitative. The references associated with the data are cited in the text.

infection phase	outcome of the infection	species	number of animals	viral RNA positive cells #	References
primary	non pathogenic	AGM	1	25*	(42) ⁱⁱ
		SM	6	12	(22) ⁱⁱ
		macaque	5	70	(4) ⁱⁱ
	pathogenic	- slow/intermediate progressors	8	50-100	(43) ⁱⁱ
			8	15	(22) ⁱⁱ
		- rapid progressors	6	102	(22) ⁱⁱ
		SI Vagmver90/ Pt mac	8	5-80*	(9) ⁱⁱ
	non-pathogenic	AGM	19	<1	(22, 31, 42, unpublished)
		SM	2	1-5 ⁱⁱ	(28) ⁱⁱ
	chronic	macaque			
		- slow/intermediate progressors	10	1-4 ⁱⁱ	(22, 43)
		- rapid progressors	14	1-100	(22, 43)
		SI Vagmver90/ Pt mac	8	2-40*	(9) ⁱⁱ

Tab. 2: Viral load in Lymph nodes during SIV infections.

The table assembles data reported in the literature on viral load in peripheral LNs during non-pathogenic SIVagm and SIVsm infection. A few representative data in SIVmac infection are shown for comparison. A low viral load is observed in LNs during the chronic phase of non pathogenic SIVagm infection. # Data are given per mm² except those values marked with an asteriks (*) that designs numbers per analysed region.

co-receptor	HIV	SIV	
CCR5	+++	+++	SIVmac, SIVsm, SIVagm, SIVmnd, ...
CXCR4	+++	- (++)	SIVagm.sab, SIVmnd-1c, ...
Bob/GPR15	+	++	SIVmac, SIVsm, ...
Bonzo/STLR33	+	++	SIVagm, ...
CCR2b	+	+	SIVrcm
CCR3	+	+	SIVsm
CCR8	+	+	SIVmac, SIVsm
CX3CR1/V28	+	+	SIVsm, SIVst
GPR1	+	+	SIVmac
Apj	+	-	
CCR9	+	-	
CMKRL1	+	-	
ChemR23	+	-	
US28	+	-	
X	-	+	SIVagm
trans-receptor			
DC-SIGN	++	++	SIVmac, SIVagm

Tab. 3: HIV-1/SIV co-and trans-receptors.

CCR5 is the major co-receptor for SIVs from African NHP's except for SIVrcm. SIVsm and SIVagm generally are also able to use Bob and Bonzo. Some SIVagm.sab isolates have in addition a tropism for CXCR4. The other co-receptors are used only rarely. X designs non identified co-receptors. The fourth column gives examples of SIV, which use the corresponding co-receptor.

		Pathogenic infection		Non-pathogenic infection	
marker		early acute	chronic	early acute	chronic
Cytokines (blood, LN, mucosa)	Pro-inflammatory				
	IFN (alpha, gamma)	+	+	+	-
	TNF- α , IL-6, IL-12, MIP-1a, MIP-1b	+	+	-	-
	Anti-inflammatory				
	IL-10, TGF- β 1	+	+	+	-
T cell profile	t-bet	+	nd	-	nd
	gata-3	+	nd	+	nd
	foxP3	+	+	+	nd
	PD-1	-	+	+	-

Tab. 4: Inflammatory profiles during non-pathogenic and pathogenic SIV infections.

The table summarizes schematically the data of the literature. The references corresponding to the data listed in the table are cited in the text. Parameters are presented as qualitative. T-bet and Gata-3 are transcription factors essential for the induction of, respectively, Th1 and Th2 responses. FoxP3 is a transcription factor expressed by CD25⁺ Treg. Nd: non determined.