Acute depletion of activated memory B cells involves the PD-1 pathway in rapidly progressing SIV-infected macaques

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Introduction

The pathogenesis of rapid progression to AIDS following HIV-1 infection remains poorly understood, despite more than 2 decades of intensive study. Rapid disease progression has been substantially curtailed by the advent of highly active antiretroviral therapy (HAART) in developed countries, but it remains a significant problem in less developed regions of the world, where HAART use is not as widespread and where the majority of HIV-infected individuals live (1, 2). The mechanisms underlying differences in rate of disease progression are thought to involve both viral and host immunological factors (3, 4). A growing body of evidence also suggests that intense chronic systemic immune activation may be the main cause of rapid disease progression (5–7). While the timing of disease stages is shorter in rhesus macaques (RMs) compared with humans, the former presents an invaluable tool in evaluating pathogenesis of rapid disease progression. Rapid progressors develop clinical AIDS within 2–5 years of initial infection, compared with approximately 10 years in typical progressors (8); similarly, rapid progressors RMs succumb to AIDS-related illness within 6 months of infection compared with approximately 2–3 years in typical progressors (9).

Previous studies in humans and RMs suggested a role for general B cell dysfunction in rapid disease progression, with some earlier studies showing associations between depletion of circulating B cells, low Ab responses to nonviral Ags, and rapid disease progression (8, 9). Loss of total memory B cells was previously shown to be an important pathogenic mechanism in viremic HIV-infected individuals, leading to impaired HIV-specific and non–HIV-specific humoral immune responses (10–13). A recent study of acute SIV infection in RMs also described a generalized loss of total memory B cells as an important factor in B cell dysfunction (14). Although these previous studies suggest an important role for general B cell dysfunction in disease pathogenesis, the B cell compartment of rapidly progressing animals has not been thoroughly characterized. The association between specific B cell subset defects and rapid disease progression in SIV infection is also not well understood.

The mechanism of B cell depletion during HIV/SIV infection is not completely understood, although a significant role for the Fas pathway in B cell depletion during HIV infection has been demonstrated. Programmed death-1 (PD-1) has recently emerged as an important immunoreceptor involved in both SIV and HIV pathogenesis, influencing T and B cell exhaustion (15–19). Although PD-1 has been previously shown to regulate B cell survival in mice (20), very little is known about its role in B cell survival during HIV/SIV infection. Here, we sought to thoroughly characterize the B cell compartment of RMs with different rates of disease progression to determine the role, if any, of B cell dysfunction and immune activation in rapid disease progression. We also investigated the role of the PD-1 pathway in B cell dysfunction during SIV infection in vitro and in vivo. To correlate our immunological findings with clinical changes occurring in the SIV-infected animals, we also followed the incidence of non-SIV infections. In addition, we compared the B cell compartments of animals with pathogenic (RMs) and nonpathogenic (sooty mangabeys; SMs) SIV infections to further understand...
the role of B cell defects in pathogenesis and disease progression. Our results identified the loss of activated memory B (mBAct) cells as an early predictor of rapid disease progression in SIV-infected RMs and suggest an important role for the PD-1 pathway in mBAct cell depletion in rapidly progressing SIV infection.

Results
Identification of 4 distinct B cell subsets, with differential surface CD21 and CD27 expression, in RM peripheral blood. CD27 has long been used as a unique marker for human memory B cells, but with the discovery of CD27+ memory B cells (21, 22), it is becoming clearer that the memory B cell compartment is a heterogeneous population consisting of both CD27+ and CD27− subsets. We therefore sought to characterize the RM B cell compartment in a manner similar to that described for peripheral B cells of HIV-infected humans (22), using a combination of CD21 (also known as complement receptor 2; part of the BCR complex and costimulatory for B cells) and CD27. We identified 4 distinct B cell subsets in peripheral blood of healthy RMs: naive (CD21+CD27−), resting memory (CD21+CD27+), and conventional or tissue memory (CD21−CD27+) (Figure 1A and refs. 22, 23). CD20 expression (based on MFI) was higher on mBAct and tissue memory B cells than on naive and resting memory cells (P < 0.001; Figure 1B). Naive B and mBAct cells were the majority subsets, each making up approximately 38% of total B cells, followed by tissue (≤16%) and resting (≤8%) memory B cells (Figure 1C). To further confirm the identity of tissue memory B cells, we examined the expression of CD21 and CD27 on CD20+ cells in various tissues of uninfected animals and found that these were indeed the majority memory B cell subset present in the tissues (Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI43271DS1). We also examined B cell subset distribution in 15 healthy HIV− humans and found that, in contrast to RMs, resting memory B cells were the majority memory B cell subset, with the combined activated and tissue memory subsets making up less than 10% of the total B cell population (Figure 1D).

We further characterized the B cell subsets by analyzing expression of the B cell–relevant activation and differentiation markers CD40, CD80, CD95, and CD11c (Figure 1E). CD40, one of the costimulatory receptors of the TNF receptor family, is an important modulator of B cell responses. Virtually all naive and resting memory B cells — and greater than 70% of tissue memory B cells — were CD40hi, whereas the majority (>70%) of mBAct cells were CD40int. CD80 (also known as B7.1, binds to CD28 and provides costimulatory signals to T cells) expression is a marker of B cell activation. CD80 was previously shown to identify a specific subpopulation of pre-mBAct cells that responds rapidly (24), secretes particularly large amounts of class-switched IgG, and efficiently presents Ag to and activate T cells (25). Consistent with this function, the mBAct cells expressed the highest levels of CD80. In addition to their roles as activation markers, expression of CD95 (death receptor; also known as Fas) and CD11c was previously identified as part of a signature profile for a new subset of tissue-based CD27+ memory B cells (26). mBAct cells expressed the most CD95 and CD11c, closely followed by resting memory B cells. CD11c was only expressed on mBAct and tissue memory B cells, with naive B cells expressing negligible amounts of CD80, CD95, and CD11c (Figure 1E). Thus, based on their surface marker expression, mBAct cells represent an effectortype B cell subset that is activated and primed to secrete large amounts of Ab and to efficiently function as APCs.

SIV infection leads to rapid depletion of peripheral blood mBAct cells. One of the first observable changes occurring in the B cell compartment following HIV and SIV infections is a marked decrease in the number of total B cells (8, 9, 11, 27). However, which specific B cell subsets are depleted, the kinetics of depletion, and association with disease progression in SIV infection of RMs are not well defined. We sought to characterize these parameters in 2 groups of RMs with distinct rates of disease progression. We classified animals that developed AIDS and were euthanized within 6 months of infection as rapid progressors, and the rest of the animals as typical progressors. We found that as early as 2 weeks following SIV infection, total B cells in blood were severely depleted in terms of both proportion and absolute number, regardless of rate of disease progression (P < 0.001 and P < 0.0001, rapid and typical, respectively; Figure 2A and B). A rebound in total B cell numbers occurred by week 12 post infection in both rapid and typical progressors, but remained significantly different from preinfection levels (P < 0.0001). Consistent with previous reports of generalized memory B cell loss in HIV and SIV infections (11, 12, 28), we also observed decreased proportions of all memory B cell subsets, particularly in the rapid progressors (Figure 2C). Upon further analysis of the memory B cell subsets in both groups of animals, we observed a striking preferential depletion of mBAct cells by 2 weeks of infection (Figure 2C). This loss was sustained in the rapid progressors who had lost 82% of their mBAct cells by week 12 post infection; in contrast, the drop in mBAct cell proportions in typical progressors was transient and returned to preinfection levels by week 12 (Figure 2D). A significant decrease in proportions of resting memory B cells was also observed in both groups. Interestingly, we observed an apparent increase in the proportion of naive B cells by week 12 post infection in rapid progressors, but not in typical progressors. We also analyzed the absolute counts of the individual B cells subsets and found similar patterns except for a few differences (Supplemental Figure 2A). The numbers of naive B cells dipped at week 2 in both groups, but were restored to preinfection levels by week 12 in the rapid, but not typical, progressors. In addition, we observed significant decreases in the absolute number of tissue memory B cells in both groups post infection.

Early depletion of mBAct cells is associated with higher set-point viremia and rapid disease progression. The striking contrast in degree of mBAct cell depletion between rapid and typical progressors prompted us to investigate whether this depletion had any significance for set-point viremia and disease progression. Multiple factors measured during chronic infection, including plasma viral load (29), total CD4+ T cell depletion in gut (30), and total central memory CD4+ T cell (T CM cell; CD28+CD95−) depletion in blood (31, 32), have been suggested to influence disease progression in both RMs and humans. We therefore analyzed all 3 parameters in an attempt to determine their relation to rapid disease progression in RMs. Rapid and typical progressors had similar peak viral loads (week 2); however, set-point viral load (week 12) was more than 1 log greater in the rapid than in the typical progressors (P < 0.0001; Figure 3A).

By 2 weeks post SIV infection, the rapid progressors had significantly lower proportions of mBAct cells compared with the typical progressors (P < 0.0001; Figure 3C). By 12 weeks post infection, the mBAct cells were further depleted in the rapid, but not typical, progressors, resulting in even greater difference between the groups (P < 0.0001). We performed correlation analyses in order to further understand the relationship between early depletion of mBAct cells and set-point viremia, and observed moderately significant inverse
Figure 1
Identification of 4 phenotypically distinct B cell subsets in the peripheral blood of healthy RM s. Flow cytometry was used to phenotypically characterize the B cell compartment in peripheral blood of healthy RM s. (A) Representative dot plots showing gating strategy used to identify B cell subsets: naive (CD21^+CD27^-), resting memory (CD21^+CD27^+), mBAct (CD21^+CD27^-), and tissue memory (CD21^lo/–CD27^-). (B) MFI of CD20 on B cell subsets. Bars represent mean ± SEM. (C) Frequencies of B cell subsets in peripheral blood of healthy RM s, with each data point representing an individual animal (n = 45). Horizontal lines represent means. (D) Distribution of B cell subsets in healthy human adults (n = 15), with each data point representing an individual donor. Horizontal lines represent medians. (E) Activation marker profiles of B cell subsets in healthy RM s (n = 10). Shown are representative dot plots (CD20-gated cell subsets overlaid on T cell contour plots) for each B cell subset and corresponding cumulative plots showing percentages of B cell subsets expressing CD40, CD80, CD95, and CD11c. Numbers within plots indicate the frequency of marker-positive cells as a percentage of the respective B cell subset. **P < 0.01; ***P < 0.001.
correlations between both frequency and absolute number of mBAct cells by week 2 (Figure 3C and Supplemental Figure 2B), which indicated that the loss of mBAct could be an early predictor of rapid disease progression. The correlations were even stronger by week 12 post infection (r = –0.7, P < 0.0001, for both frequencies and absolute numbers). Furthermore, we observed a significant inverse correlation between the frequency of mBAct cells prior to infection and the extent of their depletion at 2 weeks post infection (Figure 3B).

Analyses of total CD4+ T cells in the colorectal tissue and total TCM cells in blood did not reveal significant differences between the groups at 2 weeks post infection (Figure 3, D and E). By 12 weeks, only the frequency of TCM cells in blood was significantly lower (P = 0.02) in rapid versus typical progressors. Similarly, the frequency of these cells at 12 weeks, but not at 2 weeks, correlated inversely with set-point viremia (Figures 3, D and E). Thus, the loss of mBAct cells is associated with, and may represent an early predictor of, rapid disease progression in RMs.

Early depletion of mBAct cells is associated with impaired SIV-specific and non–SIV-specific humoral immunity. Both HIV and SIV disease progression are characterized by overwhelming non-HIV/SIV infections that eventually lead to end-stage AIDS and death. We hypothesized that the loss of mBAct cells in rapidly progressing animals would have important consequences not only for SIV-specific humoral immune responses, but also for Ab responses to non-SIV infections. We therefore measured the serum titers of SIV Env–binding Abs in both rapid and typical progressors and found that 5 of 8 rapid progressors had undetectable SIV Env Ab titers through week 20 of infection (Figure 4A). The other 3 rapid progressors mounted a modest anti-Env Ab response by week 12, and 1 had sustained Ab titers through week 20. The typical progressors, on the other hand, developed robust anti-Env Ab responses by 12 weeks post infection, with even higher Ab titers by 20 weeks (Figure 4B).

Intestinal bacterial infections are a significant cause of morbidity in SIV-infected animals, and the causative agents of these infections are usually flagellated, with the flagella representing an important virulence factor (33). We therefore measured serum Ab titers in response to the common bacterial Ag flagellin (FliC) as a means to assess the effect of loss of mBAct cells on preexisting humoral immunity to common intestinal bacteria that frequently cause clinically relevant infections in immunocompromised hosts (34). Prior to infection, anti-FliC Ab titers were nearly 2-fold higher in the rapid progressors (median, 1,889 ng/ml) than in the typical progressors (median, 982 ng/ml), but this difference did not reach statistical significance. By 2 weeks post infection, anti-FliC Ab titers in the
Figure 3
Depletion of mBAct cells is an early predictor of disease progression. (A) Plasma SIV viral load in rapid (n = 13) and typical (n = 39) progressors at 2 and 12 weeks post infection. (B) Correlation between percentages of mBAct cells prior to infection and change in mBAct cell percentages from week 0 to week 2. (C) Percentages of mBAct cells and correlations between set-point viral load and mBAct cells at 2 or 12 weeks post SIV infection. (D) Proportions of total CD4+ T cells in colorectal tissue, and correlations between set-point viral load and CD4+ T cells in colorectal tissue at 2 or 12 weeks post SIV infection. (E) Proportion of blood TCM cells and correlations between set-point viral load and blood TCM cells at 2 or 12 weeks after SIV infection. For correlation analyses, filled and open symbols represent rapid and typical progressors, respectively. Boxes represent 25th–75th percentiles with medians, and whiskers represent 5th and 95th percentiles. *P < 0.05; *** P < 0.001.
rapid progressors were significantly lower, and further decreased as the infection progressed through week 20 ($P < 0.01$; Figure 4C). In contrast, anti-FliC Ab titers did not change significantly in typical progressors ($P = 0.9$; Figure 4C). We also analyzed the incidence of non-SIV infections in both groups over a 6-month period following initial SIV infection and found that a wide variety of other infections occurred in the animals (Supplemental Tables 1 and 2). These included bacterial (Campylobacter, Shigella, enteropathogenic E. coli), parasitic (Trichomonas, whip worms, Giardia), and yeast (Candida) infections. Importantly, rapid progressors succumbed to these infections as early as 1 month post infection, and by 3 months post infection, greater than 50% of them had developed infections compared with less than 10% of the typical progressors. This rate of infection in rapid progressors was sustained throughout the 6-month period (Figure 4D and Supplemental Tables 1 and 2).

**Figure 4**
Depletion of mBAct cells in rapidly progressing SIV infection impairs SIV-specific and non–SIV-specific humoral immune responses. Serum titers of anti-SIV Env IgG Ab in (A) rapid progressors ($n = 8$) and (B) typical progressors ($n = 10$) before and at weeks 2, 12, and 20 post SIV infection. Dashed lines denote the 200-ng/ml assay limit of detection. (C) Serum titers of anti-FliC IgG Ab in rapid ($n = 15$) and typical ($n = 7$) progressors before and at weeks 2, 12, and 20 post SIV infection. Each data point represents an individual animal, and horizontal lines represent medians. (D) Incidence of non-SIV infections in rapid ($n = 15$) and typical ($n = 39$) progressors following SIV infection. See Supplemental Tables 1 and 2 for additional information. *$P < 0.05$; **$P < 0.01$. 

Preferential depletion of PD-1+ mBAct cells following SIV infection of RMs. The coinhibitory receptor PD-1 has been shown to play an important role in T cell exhaustion and survival during chronic HIV (15–17) and SIV infections (18, 19, 35). To understand the role of PD-1 in B cell dysfunction and survival, we analyzed PD-1 expression on different B cell subsets before and after infection. Prior to infection, a substantial proportion of all 3 memory B cell subsets, but not naive B cells, expressed PD-1 (Figure 5, A and B). Moreover, the mBAct cells not only had the highest proportion of PD-1+ cells, but also expressed the highest amounts of PD-1 (based on MFI, data not shown) compared with the other subsets. Given the high expression of PD-1 on the mBAct cells, we analyzed the distribution of PD-1+ mBAct cells in rapid and typical progressors as a percentage of total B cells (Figure 5C). Interestingly, the depletion of mBAct cells in rapid progressors translated into preferential depletion of PD-1+ mBAct cells by week 2, which was sustained through week 12 post infection (Figure 5C). In contrast, the proportions of PD-1+ memory B cells in typical progressors decreased significantly by week 2, but were restored to preinfection levels by week 12 (Figure 5C). This raised the possibility that PD-1 may play a role in depletion of mBAct cells.

PD-1 regulates survival of mBAct cells in RMs. To investigate a possible role for PD-1 signaling in depletion of mBAct cells, we used the human hepatoma cell line Huh-7.5 transfected with PD-L1 (Huh-7.5-PD-L1) as a source of ligand for PD-1+ mBAct B cells (Supplemental Figure 3A). Coculture of purified B cells from SIV-infected RMs with Huh-7.5-PD-L1 significantly increased mBAct cell apoptosis ($P < 0.0001$; Figure 6, A and B). Apoptosis of naive B cells was not significantly altered in the presence of PD-L1, consistent with their lower PD-1 expression (Figure 6C). Moreover, stimulation with PD-L1 did not enhance apoptosis of mBAct cells from uninfected RMs (Supplemental Figure 3B). Thus, PD-1 signaling during SIV infection plays an important role in regulating survival of mBAct cells.
Memory B cells in HIV-infected humans are primed to undergo both spontaneous and death receptor–induced apoptosis, notably through the Fas-FasL pathway (10, 27, 36), but there is little information on what role the Fas-FasL pathway plays in B cell apoptosis during SIV infection. A recent study showed an association between Fas-induced apoptosis and high PD-1 expression on CD8+ T cells in HIV-infected individuals (37), and we wondered whether this holds true for the high PD-1–expressing mB Act cells.

In order to determine susceptibility of mB Act cells to Fas-mediated apoptosis and to identify a possible role for PD-1 in mB Act cell depletion, we cultured PBMCs from 9 SIV-infected animals with and without recombinant human FasL (rFasL) in combination with PD-1 blockade and analyzed annexin V expression on mBAct cells after 24 hours of culture (Supplemental Figure 3C). In all 9 animals, we saw a significant increase in apoptosis with the addition of rFasL to the cultures; interestingly, in 7 animals, we observed a decrease in FasL-mediated apoptosis following PD-1 blockade (Supplemental Figure 3C).

We set up an in vitro ELISpot assay to track total IgM-, IgG-, and SIV gp130 IgG–producing memory B cells, based on a previously described method (38). Following polyclonal stimulation and in the presence of anti–PD-1 blocking Ab, there was a significant increase in total IgM, IgG, and SIV gp130–specific IgG ASCs. This increased ASC generation could be the result of enhanced proliferative capacity or survival of memory B cells in the presence of anti–PD-1 blocking Ab. These results were consistent with our previous report demonstrating that in vivo blockade of PD-1 in RMs with chronic SIV infection results in increased titers of SIV Env–binding Ab (19).

In vivo PD-1 blockade enhances both non–SIV-specific and SIV-specific humoral immunity. To further investigate the restorative potential of PD-1 blockade on the B cell compartment, we assessed the effect of in vivo PD-1 blockade on non–SIV-specific and SIV-specific Ab responses (Figure 7). We measured the titers of anti-FliC and anti-SIV Gag p27 antibodies in SIV-infected RMs that received 4 doses of either anti–PD-1 Ab or control Ab at 10 weeks post infection (19). The titers of anti-FliC Ab increased 2-fold in 3 of 5 anti–PD-1 Ab–treated animals by day 21 following blockade (Figure 7A). No significant increase was observed in either of the control Ab–

Figure 5
Preferential depletion of PD-1+ mBAct cells in rapid progressors following SIV infection. (A) Representative FACS plots showing PD-1 expression on B cell subsets of a healthy RM. Dot plots of CD20-gated cell subsets overlaid on T cell contour plots. Numbers represent proportion of PD-1+ cells, expressed as a percentage of the respective B cell subset. (B) Proportions of PD-1+ B cell subsets in healthy RMs (n = 45). (C) Proportions of PD-1+ B cell subsets (expressed as a percentage of total B cells) in rapid (n = 13) and typical (n = 39) progressors before and at 2 and 12 weeks post SIV infection. **P < 0.01; ***P < 0.001.
treated animals. Similarly, we observed an increase in the titers of anti-SIV p27 Ab in 4 of 5 anti–PD-1 Ab–treated animals following blockade (Figure 7B). This increase in SIV Gag–specific Ab titers was consistent with increased titers of SIV Env–binding Ab in anti–PD-1 mAb–treated RMs (19). These results demonstrate that in vivo PD-1 blockade during early chronic SIV infection enhances humoral immunity against non-SIV as well as SIV Ags, presumably as a result of enhanced survival of memory B cells. We measured the frequency of total mBAct cells in blood following in vivo blockade and did not observe a significant increase (data not shown). However, it may be important to measure the frequency of these cells in multiple tissues to rule out the possibility of differential trafficking following blockade.

mBAct cells are maintained in SIV-infected SMs. To further understand the association between depletion of mBAct cells and progressive SIV infection, we studied a cohort of uninfected and SIV-infected SMs, which — like other natural hosts of SIV — rarely progress to AIDS (Figure 8). Healthy SMs had far fewer circulating total B cells (about 12%–15% of lymphocytes) than did healthy RMs (30%–40% of lymphocytes; see Figure 2A). Unlike in RMs, we did not observe lower frequencies of circulating total B cells in SIV+ SMs (Figure 8A). In SMs, we also identified 4 B cell subsets based on CD21 and CD27 expression, although the relative distribution of these subsets was different from that in RMs (Figure 8B). Naive B cells constituted the major peripheral blood B cell subset (>40%), and the majority memory B cell subset was the tissue memory B cells, not mBAct cells as in RMs. Like RMs, less than 10% of circulating memory B cells in the SMs were resting memory B cells; however, the percentage of SM mBAct cells was significantly lower compared with RMs. In addition, the distribution of all 4 B cell subsets was comparable in SIV+ and SIV− SMs (Figure 7B), which suggests that memory B cells are not depleted in chronically infected SIV+ SMs.

We also analyzed the expression of PD-1 on different B cell subsets in SIV+ and SIV− SMs. Although PD-1 expression was high on both SM and RM mBAct cells, on SM tissue memory B cells, unlike the RMs, PD-1 expression was equally high. In SIV+ SMs, the proportion of all 4 B cell subsets expressing PD-1 was similar to that in SIV− SMs (Figure 8C), which indicates that SIV infection did not significantly alter the proportion of PD-1–expressing mBAct cells. These results suggest that in SIV+ SMs, the B cell compartment, particularly the mBAct cell subset, is not severely perturbed. Maintenance of an intact B cell compartment may be another mechanism through which SMs avoid disease progression.
Generalized B cell dysfunction is an important feature of HIV and pathological SIV infections. An association between loss of total memory B cells and loss of CD4+ T cells was previously shown in HIV-infected individuals (12), which suggested that total memory B cell loss may be a useful marker of disease progression. While comparable B cell dysfunctions have been identified in SIV-infected RMs (8, 9, 14, 28), the kinetics, specific B cell subsets affected, mechanisms involved, and role of B cell defects in rapid disease progression are not well defined. Here, we demonstrated that mBAct cells are rapidly depleted following pathogenic SIV infection and that this depletion strongly influences disease progression. Importantly, loss of mBAct cells in rapid progressors was associated with a significantly higher rate of acquisition of various non-SIV infections. We also showed that the PD-1 pathway plays an important role in the depletion/survival of these mBAct cells. Our results highlight the rapid and profound loss of mBAct cells as an important mechanism of rapid disease progression and consequent impaired humoral immune response against SIV and non-SIV infections.

A key finding of our study was the strong association between loss of total mBAct cells and higher set-point viremia (and thus rapid disease progression). Given that strong SIV-specific Ab responses are not detectable until several weeks post infection (9), presumably, the mBAct cells being depleted during the first few weeks would be primarily non-SIV-specific and constitute an integral part of preexisting humoral immunity to non-SIV infections. The mechanisms by which these non-SIV-specific mBAct cells influence set-point viremia and disease progression are not completely clear. Our results suggest that the early drastic loss of mBAct cells in rapid disease progressors contributes to a substantially higher rate of acquisition of other infections, which in turn contributes to immune hyperactivation and thus higher set-point viremia and rapid disease progression.

Another interesting finding of our study is the significantly larger proportion of mBAct (CD20hiCD21–CD27+) cells found in RMs compared with humans. The fact that the majority of circulating B cells in RMs are CD21– suggests that RMs have a higher degree of basal immune activation compared with humans. The reasons for such a discrepancy between base levels of activation in RMs and humans remains to be elucidated, but may be related to both host and environmental factors, including housing conditions and intrinsic behavior of the animals. Confounding this theory, however, is the fact that SMs do not have similar high levels of basal activation, despite housing conditions and behavior comparable to those of RMs.

We also describe here, for the first time to our knowledge in RMs, a memory B cell subset that lacked both CD27 and CD21 expression. We likened this subset to the recently described tissue-like memory B cell subset in HIV-infected viremic individuals (22). Comparable human unconventional/tissue-like memory B cells have gained recent attention because of their expansion and accumulation in viremic HIV-infected individuals as well as their association to other disease states, including malaria and common variable immune deficiency (22, 39, 40). This is an intriguing subset that may have important consequences for disease pathogenesis and merits further study. Our ability to identify such a subset in nonhuman primates presents a unique opportunity to better understand their biology and function.

Discussion

In vivo PD-1 blockade enhances both non–SIV-specific and SIV-specific humoral immunity. PD-1 blockade was performed in 5 SIV-infected animals at 10 weeks following infection. 2 SIV-infected animals treated with an isotype Ab served as controls. (A) Serum anti-FliC IgG Ab titers. (B) Serum anti-Gag p27 IgG Ab titers.
Depletion of mBAct cells from the blood likely occurs via a combination of mechanisms, including migration to other sites/tissues, decreased survival, and/or increased apoptosis. A recent study suggested that the decrease in circulating memory B cells following SIV infection mainly resulted from trafficking to lymphoid organs, with a rebound occurring once the cells returned back to the circulation from the tissues (14). This may be the case in the typical progressors, in which we observed a rebound in memory B cells by week 12, but not in rapid progressors, in which memory B cell loss was sustained. Moreover, our results strongly suggest decreased survival/increased apoptosis as a mechanism of depletion of mBAct cells and furthermore invoke a role for PD-1.

Our results support a role for PD-1 in the deletion/survival of mBAct cells in SIV infection. We found that in vitro stimulation of mBAct cells with PD-L1 induced their deletion through apoptosis, and blockade of the PD-1 pathway in vitro increased their survival and proliferation. In addition, in vivo blockade of PD-1 led to increased titers of Ab to SIV Ags and bacterial FliC. Interestingly, Fas-mediated apoptosis of mBAct cells was also decreased following blockade of PD-1, which suggests that both Fas-mediated and PD-1–mediated pathways of apoptosis may be connected in some way. This presents the interesting possibility that Fas-mediated apoptosis could be modulated through PD-1 blockade. Given the attenuated disease course observed in SIV-infected animals treated with a FasL-blocking Ab (41), PD-1 blockade may thus be a means to enhance humoral immune responses through modulation of both the PD-1 and Fas pathways. Given the uniformly high expression of PD-1 ligand expression being a determining factor in their deletion through the PD-1 pathway cannot be excluded. While we did not
of HIV infections is highest, indeed have a higher proportion of individuals living in resource-poor nations, where the incidence progression (48). It would be interesting to determine whether (46, 47). Furthermore, concurrent infections with other patho-

spective studies in humans reported a faster rate of HIV-1 disease progression in developing compared with developed countries (46, 47). Furthermore, concurrent infections with other pathogens of bacterial, helminthic, or protozoan origin have been shown to markedly affect the severity and course of disease progression (48). It would be interesting to determine whether individuals living in resource-poor nations, where the incidence of HIV infections is highest, indeed have a higher proportion of mBAct cells prior to infection, and how this may relate to disease progression. Identification of an early predictor for rapid HIV progression will greatly enhance our ability to treat and control the HIV/AIDS epidemic worldwide.

Methods

Study subjects. 52 young adult male RMs (median age, 2.8 years; range, 2.3–4.1 years) and 34 SMs, 17 SIV+ and 17 SIV− (12 male, 7 female; median age, 14 years; range, 9–25 years) from the Yerkes breeding colony were cared for under guidelines established by the Animal Welfare Act and the NIH Guide for the Care and Use of Laboratory Animals, using protocols approved by the Emory University Institutional Animal Care and Use Committee. The rapid and typical progressor RMUs had similar median age. RMs were infected with SIVmac239 intravenously at a dose of 100 TCID50 and were followed longitudinally. N. Miller (NIH, Bethesda, Maryland, USA) provided the challenge stock. SMs were housed in colonies of 50–60 animals, and SIVsm is endemic in this population; all but 1 SM used in the study acquired infection naturally. 15 HIV− blood donors were recruited after informed consent, in accordance with the Institutional Review Board of Emory University.

Phenotypic analysis by flow cytometry. Surface lymphocyte stainings were performed using 100 μl whole blood samples using multiparameter, multicolor analysis as previously described (19). Lymphocytes were obtained from necropsy tissue as previously described (19). We used mouse anti-human Abs against CD3 (clone SP34-2; BD Biosciences), CD4 (clone L-200; BD Biosciences), CD28 (clone CD28.2; BD Biosciences), CD21 (clone B-Ly4; BD Biosciences), CD27 (clone M-T2712; BD Biosciences), CD80 (clone L307.4; BD Biosciences), CD11c (clone S-HCL-3; BD Biosciences), CD20 (clone 2H7; ebiosciences), CD40 (clone MAB89; Beckman Coulter), CD95 (clone DX2, Caltag), and PD-1 (clone EH12; ref. 19). Cells were analyzed on an LSRII flow cytometer (BD Immunocytometry Systems), and data were analyzed with Flowjo software version 8.8.2 (Treestar).

Apoptosis assays. Huh-7.5.A2, PD-L1 cells (referred to herein as Huh-7.5-PD-L1) were used to assess PD-L1−mediated apoptosis of mBAct, cells, with Huh-7.5 cells as control. Huh-7.5-PD-L1 cells expressed human PD-L1 stably under CMV promoter and Neomycin resistance. Both cell lines were seeded onto separate 24-well plates at 5 × 10⁴ cells/well and incubated at 37°C for 12–16 hours. Purified B cells (>95% purity, isolated from PBMCs using NHP-specific CD20 microbeads; Miltenyi Biotec) were then added and incubated for 24 hours at 37°C, after which the cells were stained for CD20, CD27, CD21, and annexin V and immediately analyzed on an LSRII flow cytometer.

To determine rates of spontaneous and FasL-mediated apoptosis, PBMCs from 9 SIV− animals were plated in 96-well round-bottomed tissue culture plates at 2.5 × 10⁴ cells/well in the presence or absence of 10 ng/ml FasL (R&D Systems) with or without 10 μg/ml anti-human PD-1 Ab (clone EH12 with mouse variable and human IgG1; ref. 19). Plates were incubated for 24 hours at 37°C, after which the cells were stained for CD20, CD27, CD21, and annexin V and analyzed immediately on an LSRII flow cytometer.

ELISA. The titers of anti-Env IgG (49), anti-Gag p27 IgG (50), and anti-Fc IgG (51) were determined as previously described. For anti-Env Ab, cell culture supernatants of transient transfections of 293T cells with DNA/SIV239 VLP served as the source of Ag (Env). ELISA plates (Costar, Corning Life Sciences) were coated with ConA (Vector Laboratories) by incubating 100 μl of 25 μg/ml ConA in 10 mM HEPES buffer (pH 7.5) with 0.15 M NaCl, 1 mM CaCl2, and 1 mM MnCl2 overnight at 4°C. Plates were washed 6 times with PBS containing 0.05% Tween-20 (PBS-tween), after which 100 μl of undiluted VLP supernatant was added, and incubated for 1 hour at room temperature. For anti-Fc IgG Ab titers, ELISA plates were coated with native Fc IgG derived from S. typhimurium at 100 ng/well in bicarbonate buffer (pH 9.6), and plates were incubated overnight at 4°C. For anti-p27 Ab titers, ELISA plates were coated with recombinant gag p27 SIVNL437 (Immunodiagnostics Inc.) at 100 ng/well in PBS, and plates were incubated overnight at 4°C. For all assays, plates were then washed 6 times with PBS-tween, then blocked for 1 hour at room temperature with 100 μl blocking buffer (PBS-tween with 4% whey and 5% dry milk). Test sera were diluted in PBS-tween with 4% whey and 100 μl of serial 3-fold dilutions added to duplicate wells and incubated for 1 hour at room temperature. The plates were then washed 6 times with PBS-tween, and bound Ab was detected using peroxidase-conjugated anti-mouse IgG (Accurate Chemical and Scientific Corp.) and TMB substrate (KPL). Reactions were stopped with 100 μl of 2 N H2SO4. A standard curve generated using affinity-purified goat anti-mouse IgG (capture Ab, 10 μg/ml) and known amount of rhesus IgG (both from Accurate Chemicals) was included on all plates. Standard curves were fitted and sample concentrations interpolated as ng Ab/ml serum using SOFTmax 2.3 software (Molecular Devices). The concentrations of IgG are relative to the standard curve.

Memory B cell ELISpot assay. Memory B cell ELISpot assay was performed as described previously (38), with some modifications. Briefly, PBMCs were cultured in triplicates on 24-well tissue culture plates (Costar) at 0.5 × 10⁶
cells/well in RPMI 1640 medium containing 10% FBS, 100 U/ml penicillin, 100 μg/ml streptomycin, and 50 μM β-mercaptoethanol under 3 different culture conditions: unstimulated, using medium only; mitogen-stimulated, using pokeweed mitogen (MP Biochemicals) diluted 1:1,000, fixed Staphylococcus aureus Cowan strain SAC (Sigma-Aldrich) diluted 1:10,000, and 6 μg/ml CpG ODN-2006 (Qiagen; Oropen); and mitogen-stimulated (as above) plus 10 μg/ml anti-PD-1 blocking Ab (clone EH12). Cells were cultured at 37°C with 5% CO₂ for 6 days. On day 5 of culture, 96-well ELISpot plates (Millicore) were coated overnight at 4°C with affinity-purified goat anti-monkey IgM (10 μg/ml; Rockland Immunocchemicals), IgG (10 μg/ml; Rockland Immunocchemicals), or SiVmac239 gp130 (1 μg/ml; obtained through the NIH AIDS Research and Reference Reagent Program, Division of AIDS, NIAID, NIH). On day 6, ELISpot plates were washed once with PBS-tween and 3 times with PBS, then blocked with RPMI 1640 plus 1% FBS for 2 hours at 37°C. Cultured PBMCs were washed twice with RPMI 1640 plus 1% FBS and transferred to ELISpot plates starting at concentrations of 0.5 × 10⁶ cells/well, followed by 4-2-fold serial dilutions on the plates, which were then incubated at 37°C for 6 hours. Plates were then washed 3 times with PBS and 3 times with PBS-tween and incubated overnight at 4°C with 1 μg/ml biotin-conjugated anti-monkey IgM (for detection of total IgM ASCs) or 1 μg/ml anti-mouse IgG (for detection of total IgG and anti-gp130 ASCs) diluted in PBS-tween plus 1% FBS. Plates were washed 3 times with PBS-tween plus 1% FBS and 3 times with PBS and developed using 3-amino-9-ethylcarbazole (Sigma-Aldrich). Following spot development, plates were washed with distilled H₂O and allowed to dry overnight in the dark. Spots on developed plates were counted using an ELISpot plate reader (Cellular Technology) with ImmunoSpot 3.2 software. Data are presented as number of spots (i.e., ASCs) per 10⁶ PBMCs.

In vivo Ab treatment. Cytopreserved samples from a previously described study (19) were analyzed. Briefly, RMs were infused with either partially transferred to ELISpot plates starting at concentrations of 0.5 × 10⁶ cells/well, followed by 4-2-fold serial dilutions on the plates, which were then incubated at 37°C for 6 hours. Plates were then washed 3 times with PBS and 3 times with PBS-tween and incubated overnight at 4°C with 1 μg/ml biotin-conjugated anti-monkey IgM (for detection of total IgM ASCs) or 1 μg/ml anti-mouse IgG (for detection of total IgG and anti-gp130 ASCs) diluted in PBS-tween plus 1% FBS. Plates were washed 3 times with PBS-tween plus 1% FBS and 3 times with PBS and developed using 3-amino-9-ethylcarbazole (Sigma-Aldrich). Following spot development, plates were washed with distilled H₂O and allowed to dry overnight in the dark. Spots on developed plates were counted using an ELISpot plate reader (Cellular Technology) with ImmunoSpot 3.2 software. Data are presented as number of spots (i.e., ASCs) per 10⁶ PBMCs.

Viral load measurements. Plasma viral load was determined by quantitative real-time PCR as previously described (19). All viral RNA specimens were extracted and assayed in duplicate, with mean results reported and used in the analyses.

Statistics. Paired Student’s t test was used for comparison of responses before and after SIV infection, and 2-sample Student’s t test was used to compare differences between groups. Log-transformed data were used when the data were not normal, but log-normal. For nonparametric data, Wilcoxon signed-rank test (for matched-pairs samples) or Wilcoxon rank-sum test (for unpaired samples) was used. Pearson’s product moment correlation method was used for correlations. The Bonferroni method was used to adjust the P values for multiple comparisons. Statistical analyses were performed using TIBCO Spotfire S+ 8.1. A 2-sided P value less than 0.05 was considered statistically significant.

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18. Velu V, et al. Elevated expression levels of inhibitory receptor programmed death 1 on simian immuno-


35. Petrová C, et al. SIV-specific CD8+ T cells express high levels of PD1 and cytokines but have impaired proliferative capacity in acute and chronic SIVmac251 infection. *Blood*. 2007;110(3):928–936.


