IL-10 is up-regulated in multiple cell types during viremic HIV infection and reversibly inhibits virus-specific T cells

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Murine models indicate that interleukin-10 (IL-10) can suppress viral clearance, and interventional blockade of IL-10 activity has been proposed to enhance immunity in chronic viral infections. Increased IL-10 levels have been observed during HIV infection and IL-10 blockade has been shown to enhance T-cell function in some HIV-infected subjects. However, the categories of individuals in whom the IL-10 pathway is up-regulated are poorly defined, and the cellular sources of IL-10 in these subjects remain to be determined. Here we report that blockade of the IL-10 pathway augmented in vitro proliferative capacity of HIV-specific CD4 and CD8 T cells in individuals with ongoing viral replication. IL-10 blockade also increased cytokine secretion by HIV-specific CD4 T cells. Spontaneous IL-10 expression, measured as either plasma IL-10 protein or IL-10 mRNA in peripheral blood mononuclear cells (PBMCs), correlated positively with viral load and diminished after successful antiretroviral therapy. IL-10 mRNA levels were up-regulated in multiple PBMC subsets in HIV-infected subjects compared with HIV-negative controls, particularly in T, B, and natural killer (NK) cells, whereas monocytes were a major source of IL-10 mRNA in HIV-infected and -uninfected individuals. These data indicate that multiple cell types contribute to IL-10–mediated immune suppression in the presence of uncontrolled HIV viremia. (Blood. 2009;114:346-356)

Introduction

Persistent viral infection often results in T-cell impairment due to combined effects of changes in antigen-specific lymphocytes and antigen-presenting cell (APC) populations.1-4 Recent studies have highlighted the role of inhibitory immunoregulatory pathways in T-cell exhaustion during chronic viral infection. For example, the cosignaling molecule PD-1 has been shown to mediate a reversible dysfunction of virus-specific cytotoxic T lymphocytes (CTLs) during chronic lymphocytic choriomeningitis virus (LCMV) infection in mice,5,6 as well as during HIV,7-10 hepatitis B virus (HBV),10 and hepatitis C virus (HCV)11,12 infection in humans. Besides PD-1, other surface molecules can also mediate reversible T-cell dysfunction in chronic HIV infection, as shown by recent reports of the involvement of cytotoxic T lymphocyte–associated antigen 4 (CTLA-4) in HIV-specific CD4 T-cell impairment13 and TIM-3 in HIV-specific CTL dysfunction.14 Studies in murine models suggest that other cosignaling molecules expressed on the cell surface are also involved.6

Soluble factors, such as chemokines and cytokines, also regulate T-cell function.15 Among these soluble factors, the cytokine interleukin 10 (IL-10) has been shown to play a critical role in the balance between immunopathology and protective responses in infectious diseases (reviewed in Couper et al16 and Moore et al17). IL-10 is a pleiotropic cytokine that can be produced by multiple cell populations, with diverse effects on many hematopoietic cell types. The ability of IL-10 to inhibit cytokine production by T cells and natural killer (NK) cells is thought to be largely indirect, by alteration of monocyte/macrophage function.18 IL-10 decreases major histocompatibility complex (MHC) class II and CD80/CD86 expression on monocytes and macrophages and limits production of proinflammatory cytokines. However, IL-10 can also stimulate the proliferation of B cells and enhance their maturation into plasma cells.19 Inflammatory bowel disease and other excessive inflammatory responses occurring in IL-10−/− mice indicate that IL-10 is critically involved in limiting inflammatory responses in vivo.20,21 The impact of IL-10 on host immunity in the setting of infectious diseases is complex. IL-10 knockout or blockade can enhance resistance to infection with pathogens such as Listeria monocytogenes,22 Mycobacterium avium,23 and Trypanosoma cruzi.24 IL-10 plays a key role in facilitating establishment of chronic Leishmania major infection.25 However, absence of IL-10 signaling results in severe and often fatal immunopathology in Toxoplasma gondii,26 malaria,27 and herpes simplex virus28 infections. These studies indicate that a delicate balance between proinflammatory mechanisms and dampening of excessive immune activation through IL-10 is critical for successful clearance of a pathogen without harm to the host.

Recent studies in the murine LCMV model29,30 have shed new light on the role of IL-10 in the establishment of chronic viral infection. Treating mice with an IL-10-blocking antibody
shortly after acute infection, or infecting IL-10−/− mice, resulted in enhancement of T-cell responses and rapid viral clearance of an LCMV strain that is usually associated with viral persistence.29,30 These findings in murine models suggest that manipulation of the IL-10 pathway might be used to treat chronic infections in humans. Elevated IL-10 levels (IL-10 protein and/or mRNA) have been observed for several chronic infectious diseases in humans, including visceral leishmaniasis,31 leprosy,32 and tuberculosis33 (reviewed in Couper et al16 and Moore et al17). The observation that the herpesviruses Epstein-Barr virus (EBV)34 and cytomegalovirus (CMV)35 encode for IL-10 analogues suggest that IL-10 can indeed facilitate viral persistence in humans. In chronic HIV infection, the concentration of IL-10 protein in plasma increases over time.36 Research by Clerici et al37 first indicated that mitogen-stimulated PBMCs of HIV-infected individuals produced more IL-10 than did PBMCs of HIV-negative controls. Furthermore, blockade of the IL-10 pathway with an IL-10 blocking antibody enhanced some proliferative responses to HIV Env peptides. Further studies showed that IL-10 blockade could restore Env-specific T-cell proliferative responses in subjects with relatively preserved CD4 T-cell counts, but was lost in individuals with advanced HIV infection38 and that increased IL-10 levels correlated with a more advanced disease stage and the presence of syncytium-inducing virus.39

Based upon the recent LCMV results,29,30 intervention studies to block IL-10/IL-10α have been proposed as a strategy to enhance underlying immunity in chronic viral infections, such as HIV, and as a potential adjuvant to vaccination regimens.40 However, prior to considering such trials in humans, a better understanding of the role of IL-10 in HIV infection is essential. We therefore undertook a detailed study of IL-10 activity and IL-10 expression in a cohort of 80 individuals. First, we assessed the ability of antibody-mediated blockade of IL-10α to restore virus-specific CD4 and CD8 T-cell proliferative capacity in HIV-infected persons and to augment cytokine secretion by HIV-specific CD4 T cells. To define the categories of subjects who respond best to IL-10 blockade, we associated differences with clinical markers of disease status. We also compared the impact of IL-10 blockade on T-cell responses specific for HIV and for CMV, a virus that causes lifelong infection but is usually controlled by the immune system. Second, we quantified IL-10 protein in plasma and spontaneous mRNA expression in PBMCs from HIV-infected individuals at various stages of disease in cross-sectional and longitudinal studies. Finally, we identified cell populations in PBMCs from uninfected and HIV-infected individuals that expressed IL-10 mRNA. Based on the results of these studies, we conclude that IL-10 production correlates with HIV disease progression, is up-regulated in multiple cell types, and acts as a reversible mechanism to suppress both HIV-specific CD4 and CD8 T-cell responses in individuals with uncontrolled viremia.

**Methods**

**Study subjects**

Chronically HIV-infected individuals were enrolled at the Massachusetts General Hospital (MGH) or Lemuel Shattuck Hospital, in Boston, MA. Subjects identified during acute HIV infection were enrolled at the Jessen Praxis Clinic, in Berlin, Germany. These studies were approved by the respective institutional review boards of each center. Untreated persons and individuals on antiretroviral (ART) therapy were enrolled in the study (range of viral loads: < 50 to 750,000 HIV RNA copies/mL plasma). “Elite controllers” were defined as individuals with a plasma viremia below the limit of detection of standard HIV-RNA assays (<50 or 75 copies/mL) in the absence of therapy. After written informed consent, in accordance with the Declaration of Helsinki, whole blood was collected from each subject, plasma was collected, and peripheral blood mononuclear cells (PBMCs) were isolated by density centrifugation (Ficoll-Histopaque; Sigma-Aldrich) according to standard protocols. PBMCs were used as fresh or frozen cells, as indicated in the results.

**HLA tissue typing**

DNA was extracted from peripheral blood cells using Puregene DNA Isolation Kit for blood (Genta Systems). High- and intermediate-resolution HLA class I typing was performed by sequence-specific polymerase chain reaction (PCR) according to standard procedures.

**Proliferation assays with IL-10Rα blockade**

For measurement of CD4 T-cell responses, 5- (and 6-)carboxyfluorescein diacetate succinimidyl ester (CFSE)-based proliferation assays were performed as described previously.31,32 Freshly isolated PBMCs were depleted of CD8 T cells during Ficoll purification using RosetteSep CD8 Depletion reagents (StemCell Technologies) to maximize sensitivity of the assays. Cells were labeled with 1.25 μM CFSE dye (Molecular Probes) for 7 minutes, washed, and incubated with antigen in the presence of IL-10 receptor alpha chain (IL-10Rα) blocking antibody (clone 37607; R&D Systems) or IgG1 isotype control antibody at 10 μg/mL. Conditions of stimulation were tested against antigen, phytohemagglutinin (PHA; 5 μg/mL; Sigma-Aldrich), recombinant HIV p24 protein (10 μg/mL; Protein Sciences), or CMV lysate (1 μg/mL; Advanced Biotechnologies). Cells were incubated for 7 days in RPMI 1640 medium (Invitrogen) supplemented with 10% human AB serum (Gemini Bioproducts) at 37°C, 5% CO2, stained with antibodies (CD3-phycocerythrin [PE]-Cy7, CD8–Alexa 700, and CD4-allophycocyanin [APC]; BD Biosciences), and analyzed by flow cytometry (LSRII flow cytometer; BD Biosciences).

**CD8 T-cell proliferation was determined as described previously.**7 PBMCs were left undepleted, and peptides corresponding to human leukocyte antigen (HLA)–matched optimal CD8 T-cell epitopes were used as HIV antigens (0.2 μg/mL; MGH Peptide Synthesis Core Facility). No antigen and PHA were included as controls. At the end of the 7-day incubation period, cells were washed with the relevant PE- or APC-labeled HLA class I tetramer (Beckman Coulter) or pentamer (ProImmune) refolded with the respective CD8 T-cell epitopes (listed in supplemental Table 1, available on the Blood website; see the Supplemental Materials link at the top of the online article). After 30 minutes of incubation at 37°C, cells were washed and stained with surface antibodies (CD3-PE-Cy7, CD8-APC-H7, and CD4–Alexa 700; BD Biosciences) before acquisition (LSRII; BD Biosciences).

**Cytokine secretion assays with IL-10Rα blockade**

For measurement of CD4 T-cell responses, freshly isolated PBMCs were depleted of CD8 T cells during Ficoll preparation as for the proliferation assays. Cells (105) were then incubated with antigen in the presence of IL-10Rα blocking antibody (clone 37607; R&D Systems) or IgG1 isotype control antibody at 10 μg/mL. After 48 hours of incubation at 37°C, 5% CO2, supernatants were collected and IFN-γ and IL-2 levels were detected using a multiplex immunoassay (Milliplex beads; Millipore) on a Bio-Plex 200 array system (Bio-Rad Laboratories).

**Plasma IL-10 detection**

Plasma IL-10 cytokine concentration was determined by luminex assay using the Bio-Plex Precision Pro Human Cytokine 10-Plex Panel (Bio-Rad Laboratories) according to the manufacturer’s instructions on a Bio-Plex 200 array system (Bio-Rad Laboratories).

**IL-10 mRNA detection**

PBMCs were resuspended in RNA lysis solution (RLT buffer; QIAGEN) containing 1% β-mercaptoethanol (Sigma-Aldrich). Total RNA was extracted using the RNeasy Mini kit (QIAGEN). Complementary DNA
(cDNA) was generated from 0.5 µg total RNA using 0.5 µM oligo dT primer and the MultiScribe reverse transcriptase enzyme and reagents (Applied Biosystems) for 60 minutes at 42°C. A 1:10 dilution of cDNA was quantitated by quantitative PCR using Brilliant II SYBR green enzyme mix (Stratagene) on an Mx3000P real-time PCR instrument (Stratagene; 40 cycles at 95°C for 15 seconds, 60°C for 15 seconds, and 72°C for 15 seconds). To minimize detection of genomic DNA, gene–specific primers were designed to recognize spliced transcripts for human IL-10 gene (IL10/NM_000572) (F: gctgggacgagctgaag; R: gagcagctgccattg; 195 nt) and the control gene human hypoxanthine guanine phosphoribosyltransferase 1 (HPRT/NM_000104) (F: gacccagcaATG; R: gaggatgcaggtcctta; 209 nt). Cycle threshold (Ct) values for each reaction were compared with a standard curve containing a known copy number of plasmid DNA encoding each product. Results are shown as ratio of IL-10 copies per HPRT copies for each sample.

Isolation of PBMC subsets with cell sorter
Two panels of antibodies were used to isolate live cell subsets according to combinations of surface markers. In the first panel, PBMCs were stained with antibodies CD3-PE-Cy7, CD4-PerCP-Cy5.5, CD8–fluorescein isothiocyanate (FITC), CD14-PacBlue, CD19-APC-Cy7, and CD56-APC (BD Biosciences). A second panel was used to isolate dendritic cell (DC) subsets. PBMCs were stained with a lineage exclusion antibody cocktail: FITC, CD11c-PE, CD123-PE-Cy5, and HLA-DR-APC-Cy7 (BD Biosciences). T-cell subsets, B cells, NK cells, monocytes, and DC subsets were sorted as outlined in supplemental Figure 3 on a FACS Aria (BD Biosciences) equipped for biohazardous material. The FACS Aria was operated at 70 pounds per square inch with a 70-µm nozzle. For all populations, 20,000 cells were collected directly into RLT lysis buffer. IL-10 mRNA expression was assessed by reverse-transcription (RT)–PCR as described for IL-10 mRNA detection in unseparated PBMCs.

Statistical analysis
Flow cytometry data were analyzed with FlowJo software version 8.7 for Apple Macintosh (TreeStar). All statistical analyses were nonparametric and were performed using Prism 4.0 (GraphPad Software). Comparisons of cellular proliferation responses between groups were made using the Wilcoxon matched pairs test. Pairwise comparisons were verified using the Dunn posttest. IL-10 expression in cell subsets from 3 or more groups was done using the Kruskal-Wallis test. All statistical analyses were nonparametric and were performed using Prism 4.0 (GraphPad Software). Comparisons of cellular proliferation responses between groups were made using the Wilcoxon matched pairs test. Pairwise comparisons were verified using the Dunn posttest. IL-10 expression in cell subsets from 3 or more groups was done using the Kruskall-Wallis test. All statistical analyses were nonparametric and were performed using Prism 4.0 (GraphPad Software). Comparisons of cellular proliferation responses between groups were made using the Wilcoxon matched pairs test. Pairwise comparisons were verified using the Dunn posttest. IL-10 expression in cell subsets from 3 or more groups was done using the Kruskal-Wallis test. All statistical analyses were nonparametric and were performed using Prism 4.0 (GraphPad Software). Comparisons of cellular proliferation responses between groups were made using the Wilcoxon matched pairs test. Pairwise comparisons were verified using the Dunn posttest. IL-10 expression in cell subsets from 3 or more groups was done using the Kruskal-Wallis test. All statistical analyses were nonparametric and were performed using Prism 4.0 (GraphPad Software). Comparisons of cellular proliferation responses between groups were made using the Wilcoxon matched pairs test. Pairwise comparisons were verified using the Dunn posttest. IL-10 expression in cell subsets from 3 or more groups was done using the Kruskal-Wallis test. All statistical analyses were nonparametric and were performed using Prism 4.0 (GraphPad Software). Comparisons of cellular proliferation responses between groups were made using the Wilcoxon matched pairs test. Pairwise comparisons were verified using the Dunn posttest. IL-10 expression in cell subsets from 3 or more groups was done using the Kruskal-Wallis test. All statistical analyses were nonparametric and were performed using Prism 4.0 (GraphPad Software). Comparisons of cellular proliferation responses between groups were made using the Wilcoxon matched pairs test. Pairwise comparisons were verified using the Dunn posttest. IL-10 expression in cell subsets from 3 or more groups was done using the Kruskal-Wallis test. All statistical analyses were nonparametric and were performed using Prism 4.0 (GraphPad Software).

Results
Blockade of the IL-10/IL-10Rα pathway restores proliferative and effector CD4 T-cell functions in individuals with uncontrolled viremia
To define the role of IL-10 in virus-specific CD4 T-cell dysfunction, we first examined the impact of inhibition of the IL-10 pathway on HIV-specific T-cell proliferation in a cross-sectional study of 45 HIV-infected individuals. These included elite controllers who spontaneously suppressed HIV replication and chronically infected subjects—either viremic off therapy or aviremic on ARV therapy. PBMCs were labeled with CFSE and stimulated with either recombinant HIV p24 protein or CMV lysate for 7 days in the presence of IL-10Rα blocking antibody or an isotype control (Figure 1A). Blockade of IL-10Rα significantly enhanced CD4 T-cell proliferation to HIV p24 in subjects with ongoing viral replication (P < .001; Wilcoxon; Figure 1B), consistent with previous results obtained using HIV Env peptides in a smaller number of subjects.

Enhancement of HIV-specific CD4 T-cell proliferative response by IL-10Rα blockade was restricted to viremic individuals and was not observed in aviremic subjects, including elite controllers and subjects on successful ARV therapy (P = .001, Kruskall-Wallis; Figure 1C). A significant positive correlation was observed between viral load at the time of the proliferation assay and fold increase of HIV p24–specific proliferation in the presence of blocking antibody (P < .001, Spearman; Figure 1D). No significant association was seen between absolute CD4 T-cell count and p24–specific proliferation after IL-10Rα blockade (P > .05, Spearman; Figure 1E). To determine whether the impact of IL-10Rα blockade was restricted to HIV-specific T-cell responses, we assessed CMV-specific CD4 T-cell responses in a subgroup of HIV+ subjects. IL-10Rα blockade enhanced CMV-specific CD4 T-cell proliferation in 17 of 30 individuals (supplemental Figure 1B); however, this increase was not statistically significant for the entire group. Furthermore, we observed no significant difference in enhancement of CMV-specific CD4 T-cell responses between HIV viremic and aviremic individuals (P > .05; Mann-Whitney) and no correlation between the impact of IL-10Rα blockade and HIV viral load (P > .05; Spearman). In addition to assessing the restoration of proliferative capacity, we also examined the effect of IL-10Rα blockade on cytokine secretion by HIV-specific CD4 T cells. PBMCs were depleted of CD8 T cells and then cultured in the absence of antigen or stimulated with an HIV Gag peptide pool in the presence of an isotype control antibody or IL-10Rα blocking antibody. After 48 hours, supernatants were collected and the secreted IFN-γ and IL-2 levels were measured (Figure 2A–C). Overall, there was a significant increase in the amounts of both IFN-γ (Figure 2D; P = .004, Wilcoxon) and IL-2 (Figure 2E; P = .026, Wilcoxon) that were produced after IL-10Rα blockade. These data demonstrated that blockade of the IL-10 pathway not only restored proliferative responses, but also augmented direct effector functions of HIV-specific CD4 T cells.

Because of the link between CTL function and control of viremia in HIV infection (reviewed in Deeks and Walker), we next investigated the effect of IL-10Rα blockade on HIV-specific CTL responses. We stimulated PBMCs with optimal HIV epitopes in the presence of anti–IL-10Rα antibody or isotype control. After a 7-day incubation, cells were stained with the relevant HIV class I tetramer or pentamer to identify the frequency of epitope-specific CTLs. Two representative examples are shown in Figure 3A. Similar to HIV-specific CD4 T-cell responses, the fold increase of HIV-specific CTLs in the presence of IL-10Rα blocking antibody versus isotype control was significantly higher in viremic individuals than in aviremic persons (P = .009; Mann-Whitney; Figure 3B). Furthermore, we observed a significant correlation between viral load and increase of CTL responses in untreated subjects after IL-10 blockade (P = .013, Spearman; Figure 3C). Because these assays were done after removal of patient serum that may contain high levels of IL-10 protein, our data show that in vitro expression of IL-10 by PBMCs directly impacts the proliferative capacity of T cells.

IL-10 protein and mRNA levels correlate with HIV viremia and response to IL-10Rα blockade
Since antibody blockade experiments demonstrated that the negative impact of IL-10 was most apparent in individuals displaying...
higher HIV viral load, we assessed cytokine expression in HIV-infected subjects presenting with different levels of plasma viremia. IL-10 protein was measured in plasma samples collected from 35 untreated HIV-infected individuals. We observed that IL-10 concentration correlated strongly with HIV viral load ($R = 0.6017; P < .001$, Spearman; Figure 4A).

To assess cytokine gene expression, we measured IL-10 mRNA in PBMCs using quantitative RT-PCR. IL-10 mRNA expression in PBMCs correlated significantly with HIV plasma viral load (Figure 4B; $P = .001$, Spearman). In a subset of 20 untreated HIV-infected individuals, IL-10 mRNA expression in PBMCs correlated well with IL-10 protein levels in serum ($P = .001$, Spearman; data not shown). Our results for IL-10 protein and mRNA expression are consistent with previous studies. However, the correlation between IL-10 expression and response to IL-10 blockade has not been defined. We observed that plasma IL-10 protein levels (Figure 4C) and IL-10 mRNA expression in PBMCs (Figure 4D) both significantly correlated with the CD4 T-cell proliferative response after IL-10Rα blockade ($R = 0.5234$ and $R = 0.6951$, respectively; Spearman).
Relationships between IL-10 mRNA expression and disease stage

To further examine the impact of HIV infection and treatment status on IL-10 expression, we measured IL-10 mRNA in PBMCs collected from 10 healthy HIV-negative subjects and 4 cohorts of chronically HIV-infected individuals (Figure 5): 15 elite controllers; 20 viremic, untreated subjects; 12 persons with uncontrolled viremia on ARV therapy; and 11 individuals with successful ARV treatment. IL-10 mRNA levels differed among these groups (P = .002; Kruskal-Wallis), with a significant elevation of IL-10 mRNA in individuals with ongoing viremia, either untreated or on ARV, compared with elite controllers and uninfected controls (P < .01 for each, comparison Dunn posttest; Figure 5). In contrast, IL-10 mRNA expression was similar in HIV controllers and uninfected individuals. Intermediate levels of IL-10 mRNA were observed in subjects with viral loads lower than 50 copies/mL on ARV therapy, compared with the other 4 groups. Taken together, these results suggest that the reduced IL-10 expression observed upon successful ARV therapy is a consequence of decreased HIV viral load and does not result from a direct effect of ARV drugs on IL-10 expression.

Variations in IL-10 expression during periods of rapid viral load changes

Previous studies have indicated that plasma IL-10 protein levels are elevated during acute HIV infection. To better understand the link between HIV viremia and IL-10 expression during non–steady-state disease, we first examined IL-10 protein concentration in longitudinal samples collected from 3 individuals, beginning in acute infection (Figure 6A-C). Substantial levels of plasma IL-10 protein were detected at the earliest time point available; they declined rapidly during the first 6 months after infection in the 2 untreated subjects (Figure 6A-B) as well as in the treated individual (Figure 6C), suggesting that resolution of acute infection was associated with a decline in plasma IL-10, even in the absence of viral control. This analysis was extended to a total of 10 individuals for whom acute/early (within 30 days after infection) and chronic (6-12 months after infection) samples were available, and we saw a significant decline in IL-10 protein concentration at the later time point (P = .004, Wilcoxon; Figure 6D). Next, we analyzed IL-10 mRNA and protein levels in longitudinal samples collected from 2 individuals who discontinued ARV therapy (Figure 6E-F). In the absence of treatment, we observed a rapid rise in HIV viremia, but only a gradual increase of plasma IL-10 protein and mRNA expression in PBMCs for both individuals. Reinitiation of ARV treatment resulted in a marked reduction of both IL-10 protein and mRNA, suggesting that the pathogenic process responsible for IL-10 induction is at least partially reversible.

IL-10 is up-regulated by multiple PBMC subsets in HIV-infected individuals

Monocytes have been reported to be major producers of IL-10 in HIV-infected individuals, but numerous cell types can express this cytokine. To better identify IL-10–producing cell populations, we first used antibody-conjugated magnetic beads to isolate cell subsets in PBMCs from HIV-infected subjects and healthy uninfected controls. We measured IL-10 mRNA expression using quantitative RT-PCR and observed IL-10 mRNA up-regulation in all cellular...
Figure 3. IL-10Rα blockade restores HIV-specific CTL function in individuals with uncontrolled viremia. (A) Flow cytometry was used to assess proliferation of HIV-specific HLA class I tetramer–labeled CD8 T cells from 2 subjects incubated with the cognate HIV epitope, plus isotype control antibody or IL-10Rα blocking antibody. Representative data from 2 subjects is shown. Numbers in upper right quadrants indicate the percentage of CD3+CD8+Tetramer+ cells for each condition. (B) Statistical comparison of impact of IL-10Rα blockade on HIV p24–specific Tet+ CD8 T-cell proliferation in chronically HIV-infected subjects with undetectable viral loads (AVIR; n = 7), and chronically infected individuals with viral loads of greater than 1000 RNA copies/mm³ (VIR; n = 13) demonstrated a significant difference between these groups (P = .009; Mann-Whitney U test). The vertical axis (IL-10 proliferation index) corresponds to the ratio of the fraction of Tet+ cells (%CD3+CD8+Tet+) to isotype control. Short horizontal bars indicate median proliferation index. (C) Statistical analysis of the data on subjects in panel B indicated a significant correlation between IL-10Rα blocking antibody versus isotype control. The level of IL-10 protein in plasma IL-10 (pg/mL) for each condition. (D) Correlation coefficients suggest that IL-10 mRNA expression in PBMCs is a better predictor of response to IL-10 blockade than plasma IL-10 protein concentration. All statistical analyses used the Spearman rank sum test.

Figure 4. IL-10 expression correlates with HIV viremia and response to IL-10Rα blockade. The level of IL-10 protein in plasma and mRNA expression in bulk PBMCs were examined in 35 untreated HIV+ individuals. IL-10 protein concentration (A) and IL-10 mRNA expression values (B) correlated significantly with plasma viral load measurements (R = .6017 and R = .6754, respectively). Next, the impact of IL-10Rα blockade on HIV p24–specific CD4 T cells was correlated with IL-10 protein in plasma and mRNA results. The IL-10 proliferation index, defined as the ratio of the fraction of proliferating (%CD3+CD4+CFSE+low) cells in the presence of the IL-10Rα blocking antibody versus isotype control, indicated a significant correlation between an individual’s responsiveness to IL-10 pathway blockade and both plasma IL-10 protein (R = .5324, P = .0338) (C) or PBMC IL-10 mRNA (R = .6951, P < .001). (D) Correlation coefficients suggest that IL-10 mRNA expression in PBMCs is a better predictor of response to IL-10 blockade than plasma IL-10 protein concentration. All statistical analyses used the Spearman rank sum test.

As the ratio of the fraction of proliferating (%CD3+ protein or mRNA results. The IL-10 proliferation index, defined on HIV p24–specific CD4 T cells was correlated with IL-10 significantly with plasma viral load measurements (R (A) and IL-10 mRNA expression values (B) correlated significantly with plasma viral load measurements (R (A)).
Chronic differences in the frequencies of the cell populations sorted (range: gating strategies, see supplemental Figure 3). Despite major differences in the frequencies of the cell populations sorted (range: < 0.1% for DC subsets to > 10% for T-cell subsets), purities of these sorted populations were comparable and consistently higher than 98%. IL-10 mRNA expression was analyzed in PBMCs and in each purified subset using quantitative RT-PCR (Figure 7). IL-10 mRNA levels in the total PBMCs from these HIV-infected individuals were significantly elevated relative to healthy controls (P < .001; Mann Whitney). In addition, we observed a significant elevation of IL-10 in CD14+ monocytes (P = .035), CD4+ T cells (P = .001), CD8+ T cells (P = .002), CD19+ B cells (P < .001), and CD56+ NK cells (P = .004) in HIV+ individuals (Figure 7D-H; all statistics: Mann Whitney). In contrast, we did not see a significant difference in IL-10 expression for the CD123+ pDC subset (P = .161; Figure 7B) in HIV-infected individuals compared with HIV-negative controls, and observed a decline in IL-10 expression in CD11c+ mDCs (P = .028). The difference between IL-10 mRNA levels observed in CD11c+ isolated by beads and CD11c+ DCs sorted by FACS is likely due to the mixed populations in bead-sorted cells that contain a majority of CD11c-expressing monocytes and a minority of CD11c+ DCs. These data demonstrate that IL-10 expression is induced in multiple PBMC subsets in HIV-infected individuals and identify potential candidate suppressor cell populations.

**Discussion**

In this study, we examined IL-10 expression and the ability of this cytokine to suppress HIV-specific T-cell function in HIV-infected individuals. Our results indicate that IL-10 mRNA expression is increased in the setting of chronic uncontrolled HIV infection and that IL-10 mRNA expression levels correlate with plasma viremia in infected persons. Increased IL-10 mRNA levels were identified in multiple different hematopoietic cells, and IL-10Rα blockade restored not only HIV-specific CD4 cell proliferation but also antigen-specific CD8 T-cell proliferation, although these effects were seen only in viremic subjects, as well as cytokine secretion by HIV-specific CD4 T cells. Both IL-10 mRNA expression and IL-10 concentration were significantly elevated relative to healthy controls (P < .001; Mann Whitney). In addition, we observed a significant elevation of IL-10 in CD14+ monocytes (P = .035), CD4+ T cells (P = .001), CD8+ T cells (P = .002), CD19+ B cells (P < .001), and CD56+ NK cells (P = .004) in HIV+ individuals (Figure 7D-H; all statistics: Mann Whitney). In contrast, we did not see a significant difference in IL-10 expression for the CD123+ pDC subset (P = .161; Figure 7B) in HIV-infected individuals compared with HIV-negative controls, and observed a decline in IL-10 expression in CD11c+ mDCs (P = .028). The difference between IL-10 mRNA levels observed in CD11c+ isolated by beads and CD11c+ DCs sorted by FACS is likely due to the mixed populations in bead-sorted cells that contain a majority of CD11c-expressing monocytes and a minority of CD11c+ DCs. These data demonstrate that IL-10 expression is induced in multiple PBMC subsets in HIV-infected individuals and identify potential candidate suppressor cell populations.

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plasma levels were reduced through successful ARV treatment, indicating a direct effect of viral antigen load on IL-10 production. These data show that IL-10 contributes to a reversible T-cell dysfunction in HIV infected persons, and that viral antigen is a major driver of the increased levels of IL-10 observed.

Blockade of the IL-10 pathway significantly increased HIV-specific CD4 and CD8 T-cell proliferation, and also augmented effector T-cell functions, as shown by the enhanced secretion of the cytokines IFN-γ and IL-2 by antigen-specific CD4 T cells upon IL-10R blockade. Although less consistent, IL-10Rs blockade also increased CMV-specific CD4 T proliferation in HIV-infected individuals. These data show that IL-10 contributes to a broad reversible T-cell dysfunction. Significant positive correlations were observed between the fold increase in HIV p24--specific proliferation after blockade and plasma viral load or IL-10 mRNA. The variable enhancement of T-cell responses after IL-10R blockade observed in individuals with similar HIV viremia is consistent with a significant heterogeneity in the degree of T-cell exhaustion and activity of different inhibitory pathways among HIV-infected subjects. It is notable that we observed enhancement of HIV-specific T-cell proliferative responses upon IL-10R blockade only in individuals with uncontrolled viral replication. Elite controllers and individuals who suppressed viral replication while on successful ARV treatment failed to respond to IL-10R blockade. An association between IL-10 and viremia has been observed previously, and this may help to explain inconsistent results of studies where individuals were not stratified on the basis of viral load.

Consistent with results of IL-10R blockade, we observed that both plasma IL-10 protein levels and IL-10 mRNA expression in PBMCs were elevated in chronically HIV-infected individuals. A significant positive correlation between IL-10 protein and IL-10 mRNA suggests that IL-10 expression by PBMCs is representative of IL-10 production in cellular compartments that contribute to plasma IL-10. IL-10 protein and mRNA levels were significantly correlated with plasma viremia. We performed longitudinal analyses to further address the relationship between viral load and IL-10 up-regulation. Our data from 10 individuals confirm previous work demonstrating elevated plasma IL-10 levels at the time of acute infection, concurrent with very high HIV viral load, which declined significantly after resolution of primary infection, regardless of continued viremia. Conversely, our results from 2 additional patients illustrated that interruption of ARV therapy resulted in elevation of HIV viral loads.
and subsequent increases in IL-10 protein and mRNA levels. Reinstitution of ARV treatment led to a substantial decline in both measures of IL-10. In each of these cases, changes in IL-10 were delayed relative to changes in plasma viremia. Additional studies will be necessary to define whether direct stimulation of the IL-10 pathway in immune cells by viral products, indirect feedback mechanisms due systemic immune hyperactivation, or both, lead to up-regulation of IL-10 in viremic individuals.

These data raise important questions about the cellular sources of IL-10 in viremic individuals. In the murine LCMV model, recent studies by Brooks et al.29 and Ejrnaes et al.30 reach different conclusions, either favoring the hypothesis that DCs from chronically infected mice induce IL-10 up-regulation by CD4 T cells, or concluding that regulatory T cells are the major source of IL-10 during chronic LCMV infection, respectively. A similar controversy exists for HIV infection in humans as well. Early studies assessing cytokines in culture supernatants of PBMCs stimulated with PHA suggested an up-regulation of IL-10 and a switch from a Th1 to a Th2 profile in T cells.50 This hypothesis was subsequently challenged by other reports.49 Trabatoni et al.51 showed that monocytes, T cells, and B cells from HIV-infected individuals produce more IL-10 than HIV-uninfected controls, but IL-10 production by other subsets has been poorly characterized.

Our results provide a detailed assessment of IL-10 mRNA expression by several PBMC subsets. Consistent with previous studies,51 our assays suggest that CD14⁺ monocytes are a major source of IL-10 in both HIV-infected and HIV-uninfected individuals. However, analysis of highly purified cellular subsets obtained by FACS indicated that CD14⁺ monocytes showed a less dramatic up-regulation of IL-10 mRNA in HIV-infected subjects than some other subsets examined in our study, including CD19⁺ B cells and T cells. In addition, CD11c⁺ myeloid DCs, which are known to produce significant amounts of IL-10 under certain physiological conditions, appeared to produce less IL-10 in chronic HIV-infected individuals compared with uninfected controls. This result is in contrast with data obtained at the time of acute viral infection in the LCMV model29 and suggests that the IL-10-producing cell populations may differ among viral diseases, or perhaps vary in tissue location or change over time after acute infection. Such differences in IL-10-producing subsets may contribute to the broad spectrum of outcomes observed upon blockade of the IL-10 pathway in several murine models of infectious diseases. Previous reports have indicated that up-regulation of IL-10 by monocytes may be due to a direct effect of HIV Env52 and that the Env of different HIV strains could induce various degrees of IL-10 secretion by monocyte-derived dendritic cells (MDDCs).53 However, we found no significant increase of IL-10 mRNA in pDCs and mDCs. The HIV Tat protein has also been implicated in up-regulation of IL-10,54 and soluble Tat may correlate with HIV viral load in untreated infection.55 Additional studies will be necessary to determine whether IL-10–secreting cell populations differ in acute and chronic HIV infections and to define the cell types that are critical for IL-10–mediated T-cell inhibition. In particular, the role of regulatory T cells remains to be better defined. Altogether, our data demonstrate that IL-10 expression by several diverse cell populations, rather than induction of IL-10 by a single cell type, may be a key factor leading to the increase in IL-10 levels observed in HIV-infected subjects.

Interventional studies to block IL-10 activity have been proposed to enhance immunity in chronic human viral infections. Besides its impact on T-cell responses and viral load in established LCMV infection,29,30 blockade of the IL-10 pathway also allowed an otherwise ineffective therapeutic DNA vaccine to stimulate antiviral immunity and enhance viral clearance.40 IL-10 blockade may therefore be a promising adjuvant therapy to boost preventive or therapeutic vaccines. However, caution is needed with such approaches in HIV infection, since the relationship between IL-10 and HIV disease progression is complex. First, IL-10 blockade does not enhance HIV-specific T-cell function in our assays for subjects optimally treated with ARV therapy. If these findings in vitro correlate with activity of the IL-10 pathway in vivo, this may suggest that the potential impact of IL-10 blockade as a therapy would be limited in an era when optimal suppression of viral replication is the goal of patient care. Second, whereas our study, and others, show that IL-10 levels increase with disease progression and can mediate T-cell dysfunction, a genetic polymorphism in the IL-10 promoter that leads to a decrease in IL-10 expression (−592C > A) has been associated with more rapid disease progression in late stages of HIV.56 A second IL-10 promoter polymorphism (−1082A > G) that may increase IL-10 expression has been associated with slower CD4 decline and longer survival in HIV+ individuals.57 Given the crucial role of immune activation in HIV disease progression,58 it is possible that despite its inhibitory effect on T cells, IL-10 has a net beneficial impact during later stages of HIV infection by limiting systemic immune hyperactivation and CD4 T cell loss. Third, the direct effects of IL-10 on virus production are complicated, as IL-10 has been reported to inhibit HIV replication in monocytes,52,59 but also to induce expression of CCR5 in CD4 T cells,60 which may hasten infection. A placebo-controlled trial investigating the tolerance and impact of recombinant IL-10 therapy on parameters of disease progression in HIV-infected individuals did not show significant changes in either viral load or CD4 T-cell counts during 4 weeks of treatment.61 Additional studies are required to better understand the overall impact of the IL-10 pathway in viremic individuals, which likely combines the detrimental effect of IL-10 on HIV-specific CD4 and CD8 T-cell immunity with a beneficial down-regulation of systemic immune activation. Our data showing IL-10 mRNA up-regulation in multiple cell types of HIV-infected subjects suggest that, beyond T-cell responses, IL-10 could broadly alter innate and adaptive immune functions in HIV infection. It will therefore be important to develop new technical approaches selectively targeting this pathway in specific cellular subsets. Such tools will provide additional insight into HIV pathogenesis and may pave the path toward novel therapeutic interventions to enhance virus-specific immunity.

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Authorship

Contribution: D.E.K. was responsible for the overall design and conduct and provided supervision; M.A.S., D.G.K., B.D.W., and D.E.K provided intellectual input and contributed to the experimental design; M.A.B., D.S.K., D.P.T., D.F.P., P.C.R., J.S., F.P., S.L.G., and D.E.K. performed experiments; S.L.G., K.M., H.J., and F.P. provided clinical samples; M.T.W. provided technical assistance for flow cytometry and cell sorting; and M.A.B., D.S.K., and D.E.K. wrote the paper.

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