The apparent failure to maintain or restore human immunodeficiency virus type 1 (HIV-1)-specific memory responses in highly active antiretroviral therapy-treated patients is a major problem and may prevent long-term control of viremia. Untreated clinical nonprogressors control HIV-1 infection over long periods of time, with no diminution in CD4+ T-cell numbers, significantly delaying progression to AIDS. This control of viremia has been attributed to strong HIV-1-specific immune responses (2, 18, 19, 28, 40, 41). Longitudinal studies indicate that a number of apparent clinical nonprogressors show eventual immunological abnormalities and subsequently progress to disease (7, 31).

It has been suggested that genomic mutations and/or deletions found in both the host and the virus could account for the nonprogressor status, but work observing progression to AIDS in patients infected with a nef-deletion strain of HIV-1 stresses the complexity of this issue (14). In contrast to clinical nonprogressors, immunologically discordant progressors manage to control viral replication despite diminution of CD4 T-cell numbers and thus remain an enigma (these patients have also been described as long-term survivors) (38). However, the vast majority of chronically HIV-1-infected individuals fail to control HIV-1 replication, so that high-level viremia results, causing CD4+ T-cell decline despite maintenance of large numbers of HIV-1-specific CD8 T cells (12).

We have shown that an important dysfunction seen in the majority of HIV-1-infected patients from very early in disease is the inability of HIV-1-specific CD4 memory T cells to proliferate and produce interleukin-2 (IL-2) in response to HIV-1 antigens rather than an absolute loss of circulating virus-specific CD4 T cells capable of producing antiviral cytokines (50). Although HIV-1-specific CD8 cytotoxic T lymphocytes (CTL) appear to be essential in limiting viral replication, the mechanisms by which they exert their effector function remain controversial (1, 6, 12, 13). It has, however, become apparent that, as in other viral infections, the presence of functional HIV-1 Gag p24-specific CD4 helper T-lymphocytes (HTL) is crucial in driving the activation of both virus-specific CTL and other HTL as well as in maintaining their effector functions (3, 25, 26, 36, 51).

Thus, CD4 HTL dysfunction in HIV-1 infection, which becomes evident at the earliest stages of the decline in absolute numbers of CD4 T cells and is marked by an early loss of HIV-1-specific responses in terms of both proliferation and type 1 cytokine production (5, 9, 10, 30, 48), might be responsible for a later decline in CTL responses, allowing disease progression (26). Highly active antiretroviral therapy, although able to restore anti-HIV-1 CD4 HTL if given early during primary infection, is unable to rescue these responses in chronically infected individuals (11, 27, 34, 41).

In this study, we performed detailed analyses of HTL responses to a number of HIV-1 antigens and peptides (focusing on p24, the major viral capsid protein, in which multiple highly conserved HTL and CTL epitopes have been defined), in chronically infected clinical progressors, clinical nonprogressor...
sors, and immunologically discordant progressors. We assessed the magnitude, breadth, and kinetics of both proliferation and type 1/type 2 cytokine secretion in order to decipher differences in HTL responses which may determine these distinct clinical outcomes. Our data provide evidence that peripheral blood mononuclear cells (PBMC) from clinical nonprogressors exhibit strong and broad responses to many HIV-1 antigens associated with secretion of both type 1 and type 2 cytokines (IL-2 and IL-4, respectively), expressing a normal naïve/memory phenotype. We propose a role for balanced cytokine production in maintaining immune system homeostasis and keeping viral replication in check throughout the course of HIV-1 infection. This most likely represents a complex cross-regulation of the cytokine network, which balances the immune response.

MATERIALS AND METHODS

Study subjects and samples. Blood samples were taken from 20 uninfected control donors from 10 drug-naive clinical nonprogressors who had been infected with HIV-1 for more than 5 years and had an undetectable viral load and a stable CD4 T-cell count (median, 824 cells/µl; range, 480 to 1,130 cells/µl). We also studied three immunologically discordant progressors who had been infected with HIV-1 for >10 years, who were defined as showing a decline in CD4 T-cell numbers and yet demonstrating control of their viral load in the absence of antiretroviral treatment (for more detail, see below). In addition, we studied 70 patients with chronic progressive HIV-1 disease (clinical progressors) who were therapy naive, had a decreasing CD4 T-cell count (median, 285 cells/µl; range, 86 to 420 cells/µl) and high plasma viral load (median, 165,220 copies/ml; range, 5-5,000,000 copies/ml). The HLA haplotypes of patients and uninfected donors were assessed by PCR using sequence-specific primers (PCR-SSP) (8). The patients’ informed consent and Ethics Committee approval were obtained for the studies described.

Plasma viral RNA assay. Viral load in patient plasma samples was measured at each time of sample collection with the Bayer HIV-1 RNA 3.0 assay (bDNA) (detection limit, <50 copies/ml; dynamic range, 50 to 500,000 copies/ml; Bayer Diagnostics, Newbury, United Kingdom).

Isolation of PBMC and culture conditions. PBMC were isolated by density gradient centrifugation and cultured in supplemented RPMI 1640 medium (22, 23). All cell lines were cultured in 10% fetal calf serum for the studies described.

Antibodies, flow cytometry, and lymphocyte subset quantification. Murine antibodies anti-human CD3, CD4, and CD8 monoclonal antibodies were used (TetraOne; Beckman Coulter, High Wycombe, United Kingdom). The EpiX XL-MCL (Beckman Coulter) was used for flow cytometric analyses of total CD3, CD4, and CD8+ lymphocytes in whole blood.

Detailed phenotypic evaluation was carried out by three-color flow cytometry on a Becton Dickinson FACScalibur flow cytometer. Quantification of cell surface receptor expression on CD4+ T cells was performed on PBMC samples by multiparameter flow cytometry with commercially available antibodies (Beckman Coulter, High Wycombe, United Kingdom, and PharMingen, Oxford, United Kingdom). Cells were labeled with a cocktail of monoclonal antibodies: anti-CD4 conjugated with phycoerythrin (PE) or with peridinin chlorophyll protein (PerCP); anti-CD3RA-PE or anti-CD45RO-PE; and anti-CD26 conjugated with fluorescein isothiocyanate (FITC), anti-CD3-PE-FITC, or anti-HLA-DR-FITC for 30 min at 4°C. After staining, cells were washed in phosphate-buffered saline (PBS) and fixed in 2% paraformaldehyde solution in PBS. On acquisition, a gate was set around the lymphocyte population on a forward versus side scatter dot plot, and 10,000 gated events were collected for each sample. Data analysis was performed with CellQuest software, and data are presented as mean percent ± standard error of the mean (SEM). Appropriate isotype-matched negative controls were run in parallel.

Gag p24 peptides and HIV-1 antigens. Peptides and HIV-1 recombinant antigens were obtained from the Medical Research Council (MRC) Virus Resources Facility for AIDS Reagents (National Institute for Biological Standards and Controls, Potters Bar, United Kingdom). The 22 Gag p24 peptides were 20-mers with a 10-amino-acid (aa) overlap covering p24 Gag (aa 133 to 363 of HIV-1 SF2; ARP7881-22). The HIV-1 recombinant antigens used were described previously (22). Cells were cultured with antigen or peptide at a final concentration of 10 µg/ml. In addition, the inactivated gp120-depleted HIV-1 immunogen (Remune) and its constituent Gag p24 antigen (native p24; clade A) (both from Immune Biological Corp., San Diego, Calif.) were used at 3 µg/ml.

Proliferation assays. PBMC (10⁴/well) were cultured with antigen, mitogen, or cytokine in round-bottomed microtiter plates (Greiner) in 10% AB plasma (200 µl; Sigma) in RPMI. The antigens, mitogens, and cytokines used were described previously (22). On day 5, 100 µl of supernatant was collected and stored at −20°C for subsequent cytokine measurement. Each well was then pulsed with 1 µCi of [methyl-3H]thymidine (Amersham International, Amersham, United Kingdom), and 16 h later cells were harvested onto filter filters (Nunc, Oy, Turku, Finland). Proliferation as measured by [methyl-3H]thymidine incorporation was evaluated by liquid scintillation spectroscopy with a 1205 Betaplate counter (Wallac) (22, 23). Results are expressed as the stimulation index and as the mean counts per minute (cpm) for triplicate cultures, with an error of the mean of <15%. A positive response is defined as a stimulation index value of 5 or more. Control wells for calculation of background activity contained PBMC only. Control wells for calculation of background activity contained indicator cells only (plus 16 U of mouse recombinant IL-4 per ml for D36 cells).

RNA preparation and first-strand cDNA synthesis. Stimulated PBMC from HIV-1-infected individuals were obtained at different time points before and after stimulation in bulk cultures, and 10⁵ PBMC were taken for RNA isolation. Total RNA was extracted with a single-step RNA STAT-60 method (AMS Biotechnology, Abingdon, United Kingdom). Samples were then treated with RNAase-free DNase I (Boehringer-Mannheim, Germany) and 0.1 µg of random hexamers (Pharmacia Biotech, St. Albans, United Kingdom). After 1 h, the reaction was stopped by incubation at 90°C for 5 min, and samples were cooled on ice.

Amplification of gamma interferon (IFN-γ), IL-2, IL-4, and IL-10 gene-specific cDNA was performed as described previously (20). Briefly, primers were for IFN-γ (upstream 5′ primer, 5′GCATGTTTGGGT CTCTTTGCTGTGTACTGC3′, and downstream 3′ primer, 5′CTCCTTTTCTC GCTCCCTGTTTTAAGCTGTGCTG3′), IL-2 (upstream 5′ primer, 5′CATGGCTCAA GAGATCTGCTGCTGCTGCTG3′, and downstream 3′ primer, 5′GTTTATCCATGCTGGTGTTTCTTG3′), IL-4 (upstream 5′ primer, 5′GGCAGACATT GACCGAGGACACAAGTGCGATA3′, and downstream 3′ primer, 5′AAGCTCTGGTGTTGCTTCCCAAGCACAGAC3′), and IL-10 (upstream 5′ primer, 5′AAGCTGAGAACCAAGACACAGACACTAAGGGC3′, and downstream 3′ primer, 5′AAGCTATCCAGGAGACAGGATGCGGCTG3′) (all from Clontech, Basingstoke, United Kingdom), and the standard protocol from Clontech was used. Samples, including appropriate controls (Clontech positive controls; sterile H₂O, no cDNA negative controls), were subjected to 35 cycles of amplification using denaturing conditions for 45 s, an annealing phase at 60°C for 45 s, and an extension at 72°C for 2 min. After amplification, 15 µl of each PCR mixture was resolved on a 1.8% agarose gel (Gibco-BRL Life Technologies, Paisley, United Kingdom), visualized under UV light after ethidium bromide staining, and photographed. A 100-bp DNA ladder (Gibco-BRL) was included to confirm the size of the PCR products.
IFN-γ and IL-4 Elispot assays. Detection of single-cell IFN-γ and IL-4 release by Elispot was carried out as previously described (22, 29). Briefly, 10^5 PBMC/well were cultured in 10% AB plasma–RPMI (200 µl; Sigma) in 96-well polyvinylidene difluoride-backed plates (Millipore, Watford, United Kingdom) coated with anti-IFN-γ or anti-IL-4 monoclonal antibodies (Mabtech, Stockholm, Sweden). Cells were stimulated with appropriate peptides at 10 µg/ml or phytohemagglutinin at 10 µg/ml in 10% AB plasma–RPMI (200 µl). Negative controls comprised cells cultured in the presence or absence of an irrelevant peptide. Plates were incubated overnight at 37°C for 16 and for 40 h for detection of IFN-γ and IL-4, respectively. Prior to the development and detection step, supernatants were collected from the IFN-γ and IL-4 plates and stored at −110°C for subsequent measurement of IL-4 and IL-2, respectively, as described above. Spot-forming cells (SFC) were then detected according to the manufacturer’s instructions (Mabtech). A positive result is defined as a score of 5 or more SFC above background.

Statistical analysis. Computer software (Statview 5; Abacus, Berkeley, Calif.) was used for all statistical calculations. Responses to the antigens, mitogens, and cytokines in patients and controls were compared with a Mann-Whitney U test.

RESULTS

HIV-1-specific proliferative responses to HIV-1 antigens in clinical nonprogressors are associated with production of both IL-2 and IL-4. We assessed the magnitude and breadth of both proliferation and cytokine (IL-2 and IL-4) production in response to a wide range of HIV-1 antigens and peptides. All 10 nonprogressors showed strong proliferation in response to all HIV-1 antigens tested (Fig. 1A). Although the most vigorous responses were those directed at recombinant p24 (mean stimulation index ± standard deviation, 60 ± 41), recombinant gp120 and Nef also elicited large responses (stimulation index, 30 ± 28 and 30 ± 29, respectively). In addition, strong responses to whole HIV-1 immunogen (Remune) and native clade G p24 were seen (stimulation index, 26 ± 20 and 25 ± 20, respectively).

Strong anti-p24 responses were accompanied by a wide breadth of responses across overlapping p24 peptides (Fig. 1B, Table 1). Proliferative responses to all HIV-1 antigens tested were absent both in uninfected controls and in 93% of clinical progressors. The responses of nonprogressors to the majority of other viral and recall antigens were comparable to those seen in uninfected controls (Fig. 1C and D). Nonprogressors showed significantly (P < 0.001) increased responses to herpes simplex virus and cytomegalovirus antigens compared to uninfected controls, which concurs with our previous findings (21).

In clinical nonprogressors, IL-2 production was readily detectable in response to all HIV-1 antigens and peptides tested where proliferative responses were seen (Fig. 2A and B, re-
respectively; Table 1). Surprisingly, and in contrast to previous findings, in the majority of nonprogressors (7 of 10), proliferation to all HIV-1 antigens was associated with notable production of IL-4 (Fig. 2C). The p24-induced IL-4 production was due to the ability of the cells to secrete IL-4 in response to p24 mRNA expression as assessed during 4 days of culture with RT-PCR revealed that specific IL-4 mRNA expression peaked at day 3 in culture and had declined by day 4 (Fig. 2E). A representative control experiment showing kinetics of IL-4-specific mRNA expression when PBMC were stimulated with phytohemagglutinin is presented in Fig. 2F. The data shown are typical of over five RT-PCRs performed and are consistent with the bioassay results.

Phenotypic analysis of peripheral blood CD4+ T cells from clinical nonprogressors revealed normal percentages of subpopulations of naïve (CD4CD45RA) and memory/effector (CD4CD45RO) CD4+ T cells (mean ± SEM, 36.6% ± 3.3% and 64.9% ± 3.5%, respectively) comparable to those seen in uninfected or seronegative controls (17) (Fig. 3). The majority of CD4+ T cells expressed the costimulatory molecule CD28 (32) (mean ± SEM = 33.8% ± 3.9%), while the majority of CD4+ T cells did not express HLA-DR (mean ± SEM = 6.2% ± 1.8%), suggesting that these were resting CD4+ T cells (17) (Fig. 3).

Presence or absence of IL-4 does not affect strength or breadth of HIV-1-specific responses in clinical nonprogressors. Two drug-naïve nonprogressor patients who had stable or increasing CD4 T-cell counts (480 and 1,052 cells/µl) and undetectable viral loads (<50 copies/ml) and had been infected with HIV-1 for >8 years were selected from this cohort based on in vitro CD4 HTL responses and studied in more detail. Both individuals showed comparable proliferative responses and broad IL-2-producing capacity in response to all antigens and peptides tested (Fig. 4A, B, D, and E). Strong p24-, gp120-, and Nef-specific responses (mean stimulation index = 79; stimulation index range, 65 to 99) as well as strong responses to whole HIV-1 immunogen and native clade G p24 (mean stimulation index = 57; stimulation index range, 44 to 67) were observed in both clinical nonprogressors.

The responses of nonprogressors to recall antigens, mitogens, and IL-2 were comparable to those seen in uninfected controls. For recall antigens, the mean stimulation index was 81, and the stimulation index range was 13 to 190 (Fig. 4A), and, as previously described, for mitogens the mean stimulation index was 319 and the stimulation index range was 55 to 559 and for IL-2 the mean stimulation index was 335 and the stimulation index range was 305 to 373 (21). However, IL-4-producing capacity was limited, and differential IL-4-inducing capacity in response to recombinant p24, clade G p24, and overlapping Gag peptides as well as gp120, influenza virus, herpes simplex virus, purified protein derivative, and Candida antigens was observed (Fig. 4C and F). In PBMC from clinical nonprogressor patient 1, Gag p24 induced significant IL-4 production mainly in response to peptides 14 and 6, whereas in clinical nonprogressor patient 2 notable amounts of IL-4 were observed only in response to recombinant Nef and Remune, while no IL-4 was produced in response to either recombinant p24, clade G p24, or Gag peptides (Fig. 4C and F). This response to Remune is likely to be directed at HIV-1 antigens other than p24 contained in the immunogen (R. Moss, personal communication).

HIV-1-specific responses in immunologically discordant progressors are mainly restricted to p24. Three drug-naïve immunologically discordant progressor patients were studied who, despite declining CD4+ T-cell counts (240, 450, and 346 cells/µl), maintained control of HIV-1 replication (for two patients, <50 copies/ml, and for patient 1, with the lowest CD4+ T-cell count, 1,236 copies/ml). Strong proliferative responses to HIV-1 Gag p24 recombinant antigen were seen in all three patients (mean stimulation index = 104; stimulation index range, 92 to 122; Fig. 5A). However, gp120- and Nef-specific responses (although significant) were lower than those to p24 (mean stimulation index = 9.5; stimulation index range, 4 to 21), in contrast to the results obtained with the clinical nonprogressors. Two of the three patients responded to both native p24 and Remune (mean stimulation index = 17; stimulation index range, 10 to 30; Fig. 5A).

Strong p24-specific lymphoproliferative responses were reflected in the breadth of the responses to p24 peptides (Fig. 5B). Proliferation was in all cases associated with IL-2 production, while no IL-4 was detected in cultures (data not shown). The responses of immunologically discordant progressors to recall antigens, mitogens, and IL-2 were lower than those seen in nonprogressor patients and uninfected controls. For recall antigens, the mean stimulation index was 51 and the stimulation index range was 2 to 143 (Fig. 5A); for mitogens, the mean stimulation index was 80 and the stimulation index range was 21 to 142; and for IL-2, the mean stimulation index was 189

### Table 1. Percentage of patients responsive to each peptide

<table>
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<tr>
<th>Peptide</th>
<th>Clinical nonprogressors (n = 10)</th>
<th>Clinical progressors (n = 10)</th>
<th>Uninfected controls (n = 10)</th>
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*LPR, lymphoproliferative response; *, borderline positivity (stimulation index = 5). Cutoff for IL-2 assay, 250 cpm; detection threshold, 0.05 U/ml; cutoff for IL-4 assay, 265 cpm; detection threshold, 0.025 U/ml.*
and the stimulation index range was 116 to 235 (data not shown).

Kinetics of p24-specific responses in immunologically discordant progressor 2: proliferation and cytokine profiles. Since immunologically discordant progressor patient 2 showed the strongest anti-p24 responses in the immunologically discordant progressor group, we chose to investigate these responses in more detail. The kinetics of p24-specific proliferative responses were consistent and revealed maximum proliferation between days 5 and 6, followed by a gradual decline (Fig. 6A). Aliquots of culture supernatants were collected from bulk cultures at different time points to assess the time course of cytokine production following activation with recombinant p24. Production of IL-2 was found to peak at day 3 in culture,
after which it declined gradually and disappeared after day 7 (Fig. 6B). Lack of IL-4 detection (Fig. 6C) did not reflect its consumption or utilization, since IL-4 was not detected throughout the 12 days of culture, which concurs with the RT-PCR analysis (see below). Secretion of IL-10 was found to peak within the first 24 h of stimulation, at day 1 of culture, followed by a dramatic fall (Fig. 6D).

In order to demonstrate the sensitivity of our bioassays, the responses of these cells (CTLL-2, CT.h4S, and D36) to a range of concentrations of recombinant IL-2 (rIL-2), rIL-4, and rIL-10, respectively, were included in each experiment (Fig. 6E to G). These cells consistently proliferated to human recombinant cytokines in a dose-dependent manner, with significant proliferation above background control levels occurring for rIL-2 at a concentration of 0.05 U/ml with CTLL-2 cells, for rIL-4 at a concentration of 0.025 U/ml with CT.h4S cells, and for rIL-10 at a concentration of 0.01 U/ml with D36 cells. Bioassays are in general more sensitive than enzyme-linked immunosorbent assay (ELISA) (23, 43), and others have shown that similar levels and kinetics are obtained when supernatants are assessed in parallel with bioassays and ELISA (24). However, the best correlations are obtained when bioassays are compared with RT-PCR analysis (Fig. 6D).

**RT-PCR analysis of cytokine-specific mRNA in immunologically discordant progressor 2.** RT-PCR analysis was used to confirm changes in expression of type 1 and type 2 cytokine-specific mRNAs occurring during 4 days of culture (Fig. 7). Specific IFN-γ, IL-2, IL-4, and IL-10 mRNA expression was determined by RT-PCR, which showed strong expression of specific IFN-γ mRNA after 24 h in culture that increased dramatically on days 2 and 3 and then diminished by day 4. IL-2 mRNA expression, although at lower levels, occurred early in culture (day 1) and persisted throughout. In contrast to the clinical nonprogressors, in whom IL-4 was detected at days 3 to 4 in culture, IL-4-specific mRNA remained undetectable throughout p24 stimulation for the immunologically discordant progressors. IL-10-specific mRNA expression was visible at 24 h, after which it disappeared. Thus, RT-PCR analysis concurs with the data obtained in our bioassays.

**During clinical progression, both CD4+ and CD8+ HIV-1-specific T-cell responses are associated with increased IL-4 production, decreased IFN-γ production, and absence of IL-2.** Follow-up of immunologically discordant progressor patient 1 revealed that clinical progression to symptomatic disease was associated with a drop in CD4 T-cell count from 240 to 201 cells/µl and an increase in viral load from 1,236 to 6,483 copies/ml in 6 months. Although these changes in CD4 T-cell number and viral load were moderate, they were associated with a decline in both proliferation and IL-2 production in response to all antigens tested (Fig. 8A and B). However, a dramatic increase in the secretion of IL-4 in response to all the antigens tested was apparent (Fig. 8C).
FIG. 4. Lymphoproliferative responses (LPR) and IL-2 and IL-4 production in response to HIV-1 antigens, other viral and recall antigens, and Gag p24 peptides in clinical nonprogressor patients 1 and 2 (CNP-1 and CNP-2). PBMC from two clinical nonprogressor patients were cultured in the presence of various antigens (A) or peptides (D) in triplicate for 6 days, and [3H]thymidine incorporation was measured as described in Materials and Methods. Results are expressed as stimulation indices. IL-2 production in response to various antigens (B) or peptides (E) was measured in culture supernatants by proliferation of the indicator cell line CTLL-2, and IL-4 production in response to various antigens (C) or peptides (F) was measured in culture supernatants by proliferation of the indicator cell line CT.H4S. Results are expressed as the mean cpm for triplicate cultures, with an error of the mean of <15%. See Fig. 1 legend for other abbreviations.
peptides. Before progression, activation with relevant peptide induced significant levels of IFN-γ SFC, accompanied by secretion of substantial amounts of IL-2 in culture (Fig. 9A and B). However, upon progression to disease, there was a decrease in IFN-γ SFC, an absence of IL-2 in culture, and an increase in IL-4 SFC, which was confirmed by measurable levels of IL-4 in culture supernatants with the CT.h4S bioassay (Fig. 9C and D). Additional follow-up analysis on this patient...

![Graphs](image-url)
at two subsequent time points 6 months apart showed that lack of overall responses remained a reproducible finding and was associated with low CD4 count and a persistent viral load of \( \sim 10,000 \) copies/ml, confirming that this patient had indeed progressed.

**DISCUSSION**

Design of successful immunotherapeutic interventions for chronic HIV-1 infection during highly active antiretroviral therapy is dependent on our understanding of HIV-1-specific immune responses observed in clinical nonprogressors. Such interventions should guarantee protection from HIV-1 disease by controlling viremia and preventing the onset of AIDS. In this study, we examined the magnitude, breadth, and kinetics of the CD4 T-cell component of the anti-HIV-1 immune responses in patients with undetectable or low viral loads who maintained strong proliferative responses, particularly to Gag p24. We present data concerning the quantity and quality of anti-HIV-1 T-cell responses observed in clinical nonprogressors and immunologically discordant progressors that lead to effective control of viremia.

Although the strong anti-p24 responses found in nonprogressors have been linked to control of viremia (41), our data show that this is also true in the immunologically discordant progressors. In contrast to previous reports, clinical nonprogressors, who were characterized by normal and stable CD4 T-cell counts and undetectable viral loads, displayed robust anti-HIV-1 responses to all HIV-1 antigens tested (including clade B p24, gp120, and Nef as well as clade G p24 and the whole HIV-1 immunogen Remune), showing CD4 T-cell cross-reactivity to both clade B and G. Anti-p24 responses specific for the majority of Gag peptides were present, indicating that perhaps MHC class II restriction is more promiscuous than class I.

Proliferation was always accompanied by significant production rather than loss of circulating virus-specific T cells capable of producing the antiviral cytokines tumor necrosis factor alpha and IFN-\( \gamma \) was observed in chronic progressors (50). Indeed, it is the proliferative dysfunction of CD4 T cells seen early during infection and before any decline in CD4 T-cell numbers that has been suggested to be responsible for their inability to support virus-specific CD8 T-cell effector function (26).

Although the strong anti-p24 responses found in nonprogressors have been linked to control of viremia (41), our data show that this is also true in the immunologically discordant progressors. In contrast to previous reports, clinical nonprogressors, who were characterized by normal and stable CD4 T-cell counts and undetectable viral loads, displayed robust anti-HIV-1 responses to all HIV-1 antigens tested (including clade B p24, gp120, and Nef as well as clade G p24 and the whole HIV-1 immunogen Remune), showing CD4 T-cell cross-reactivity to both clade B and G. Anti-p24 responses specific for the majority of Gag peptides were present, indicating that perhaps MHC class II restriction is more promiscuous than class I.

Proliferation was always accompanied by significant production of IL-2. Furthermore, substantial secretion of IL-4 was detected in cultures of 7 of 10 clinical nonprogressors and across the majority of HIV-1 antigens and peptides tested. Immune responses to other recall antigens, mitogens, and IL-2 were comparable to those observed in HIV-1-seronegative controls. The increased responses to herpes simplex virus and cytomegalovirus antigens most likely reflect the higher prevalence of these infections within the HIV-1-infected population.
In a representative patient, clinical nonprogressor 1, IL-4 production was seen in response to all antigens tested except varicella-zoster virus, cytomegalovirus, and tetanus toxoid. While IL-4 was produced in response to all p24 antigens tested, when individual p24 peptides were assessed for the ability to induce secretion of IL-4, it was revealed that this IL-4-inducing capacity was restricted to peptides 14 and 6. The strength of p24-specific responses was not affected by the differential ability of PBMC to release IL-4 in response to p24 antigen and/or peptides, since in this case the presence of IL-4 in the culture did not have a detrimental effect on either proliferation or the IL-2-producing capacity of the anti-HIV-1 HTL response. This was also the case for clinical nonprogressor patient 2, in whom IL-4 production was seen only in response to Nef and Remune.

These data might reflect cross-modulation and cross-regulation between type 1 and type 2 cytokines (particularly their characteristic products, IFN-γ and IL-4, known to often oppose each other’s action), which may be crucial in maintaining a consistent cellular response in HIV-1-infected nonprogressors, or may reflect the intrinsic ability of individuals to pro-
duce type 2 cytokines due to a previous anamnestic encounter with other pathogens (44, 46). Alternatively, restricted numbers of p24 peptides might act as altered peptide ligands for particular CD4 T-cell subsets or clones, resulting in a qualitatively different pattern of signaling and subsequently in different functional capacity (45). Such a phenomenon may well be responsible for shaping the T-cell-mediated immune responses to HIV-1.

Immunologically discordant progressors, characterized by decreasing CD4 T-cell counts and undetectable or low viral loads, displayed equally vigorous anti-HIV-1 responses, which were directed mainly towards recombinant clade B p24. These anti-p24 responses were approximately 10-fold higher than those seen towards the other HIV-1 antigens tested, and the strength of p24 antigen-specific response was reflected in equally broad responses to the panel of overlapping p24 peptides. Responses to other recall antigens, mitogens, and IL-2 were approximately twofold lower than those seen for clinical nonprogressors and seronegative controls. IL-2 but not IL-4 was readily detected in cultures where significant proliferation was seen. This concurs with previous findings and with the rationale of successful prophylactic vaccination, which is aimed at developing strong type 1 cell-mediated responses, which should not be affected by type 2 responses (39, 41).

Our findings raise two important questions. First, are the anti-p24 responses seen in immunologically discordant progressors protective? It is possible that the p24-specific responses are of primary importance and that clinical nonprogressors make good responses to other antigens only because those responses remain intact, as they have not been depleted by disease progression. The second important question is why do CD4 T-cell numbers decline in immunologically discordant progressors despite controlled viremia? The mechanism behind this phenomenon is difficult to explain and may involve increased sensitivity to one or more factors causative of indirect virus-mediated cytopathic effects, such as activation-induced cell death or multinucleated cell syncytium formation (47). Clonal deletion and/or anergy may result from a variety of indirect mechanisms of HIV-1 infection. These include hyperactivation, activation-induced cell death of uninfected T cells, and damage or destruction of regenerative compartments (thymus and/or bone marrow) (45). Different levels of genetic susceptibility of the host to these mechanisms may explain CD4 T-cell depletion in the presence of low viremia.

The kinetics of p24-specific response in an immunologically discordant progressor reflected a predominant presence of type 1 cytokine responses, with the immunosuppressive type 2 cytokine IL-10 seen only transiently at the beginning of culture. A strong type 1 cytokine (IFN-γ and IL-2) profile in response to p24 suggests suppression of virus replication; however, in addition, it could also suggest that these anti-p24 responses are inappropriately skewed. Such a polarized immune response may be indicative of a hyperactivated state. However, our data do not support this hypothesis, as the breadth of response to the Gag p24 peptides was comparable in immunologically discordant progressors and in nonprogressors.

A very strong type 1 response might be detrimental in some HIV-1-infected patients in whom viremia is controlled but the indirect bystander effects of the immune activation (possibly of a type 1) may be important. This would concur with work by Zinkernagel et al. and implicates anti-HIV-1 T-cell responses in immune-mediated pathology (52). We suggest that a balanced type 1/type 2 T-cell profile is indicative of a nonpolarized immune response which has not been skewed by hyperactivation. In addition, the kinetics of cytokine production at both the protein and mRNA level suggest strong type 1 p24-induced responses, with only transient detection of the immunosuppressive cytokine IL-10. A sharp but very transient peak of IL-10 production at day 1 is likely to be due to nonspecific or innate immune cells such as monocytes and may not represent type 2 polarization.

Our data suggest that in individuals who control viremia through maintenance of p24-specific responses, there is a minimal requirement for cross-regulation of immune responses or these type 1 responses are cross-regulated efficiently by brief transient type 2 cytokine (IL-10) production. The data may indicate reduced immune activation in immunologically discordant progressors compared with that observed in chronic progressive HIV-1 infection, with no need for the IL-10-secreting anergic suppressor T-cell subset (33, 49). Commitment to IFN-γ or IL-4 production does not necessarily require proliferation and cell division (4), and thus, loss of a particular virus-specific cytokine-secreting subset might simply reflect preferential infection with HIV-1 (42).

Clinical progression in one immunologically discordant progressor patient, although accompanied by only moderate changes in CD4 T-cell count and viral load, resulted in dramatic immunological changes and was associated with a diminution of type 1 responses and a substantial release of the type 2 cytokine IL-4 by HIV-1-specific CD4 and CD8 T cells. Reduced proliferation in response to all antigens and peptides tested was associated with an absence of IL-2 production except for the Candida antigen, with which proliferation was minimal. This absence of IL-2 production was accompanied by increased production of IL-4 in response to all antigens tested and was maximal for all three HIV-1 antigens tested (p24, gp120, and NeF) and Candida antigen. This increased production of IL-4 also extended to HIV-1 peptide-specific CD8 T cells and was accompanied by an overall reduction in expression of CD28 (data not shown), which is known to be down-modulated by IL-4 and which correlates with virus replication and dissemination (16, 35).

Our findings confirm that HIV-1-specific CD4 T cells are present but have been rendered anergic directly by HIV-1 or indirectly by clonal suppression. The dramatic increase in IL-4 secretion suggests that cells are responsive to antigenic challenge, but because of the immunosuppressive microenvironment, the cells are only capable of responding with secretion of IL-4. Induction of type 2 immunity may not necessarily diminish the quantity and quality of cell-mediated immune responses required to be elicited when a vaccine antigen is administered, but such data must be taken into consideration when designing prime-boost strategies (39).

Conclusions regarding correlations between the secretion of immunoregulatory cytokines such as IL-4 and IL-10 and disease progression should be considered carefully, since although IL-4 secretion was enhanced with disease progression in one immunologically discordant progressor, in the clinical nonprogressor patients secretion of IL-4 did not affect virus-specific T-cell responses. Our data raise questions for prophy-
lactic approaches where the rationale may be to induce a protective immune response that controls HIV-1 but does not entirely eliminate HIV-1 and also for therapeutic approaches where the aim is to induce and maintain protective responses in an immunosuppressed microenvironment.

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REFERENCES


