# Tissue Infiltration in a CD8 Lymphocytosis Syndrome Associated with Human Immunodeficiency Virus-1 Infection Has the Phenotypic Appearance of an Antigenically Driven Response

S. Itescu, J. Dalton, H.-Z. Zhang, and R. Winchester

Division of Autoimmune and Molecular Diseases, Department of Pediatrics, Columbia University, College of Physicians and Surgeons, New York 10032

## Abstract

HIV-1 infection may initiate to an HLA-associated response designated diffuse infiltrative lymphocytosis syndrome, characterized by increased numbers of circulating CD8 T cells that infiltrate salivary glands, lungs, gastrointestinal tract, and kidneys. Since this response could either be an antigenically driven process induced by HIV-1 or a lymphoproliferation of cells with neoplastic or unusual features, we sought to define the phenotype of the cellular populations, the nature of tissue derangement, and the tissue localization of virus in diffuse infiltrative lymphocytosis syndrome. Circulating CD8 T cells were greatly increased while CD4 T cell numbers remained in the range found in asymptomatic seropositive persons. The majority of CD8 and CD4 T cells in both blood and tissues had the memory phenotype of CD29<sup>+</sup> ( $\beta_1$  integrin) and CD11a<sup>+</sup>/ CD18 ( $\beta_2$  integrin) expression, but lacked markers of recent activation. A proportion of the circulating CD8 T cells also expressed CD57 (Leu 7) but not other markers of natural killer cells. HIV-encoded proteins were identified in tissue macrophages located in periacinar areas of the salivary glands. CD54 (intercellular adhesion molecule-1), a ligand for the CD11a integrin, was strongly expressed on postcapillary venule endothelium within lymphoid foci, and HLA-DR molecules were found on limited regions of ductular epithelium adjacent to lymphoid aggregates. These findings suggest that (a) the visceral lymphocytic infiltration in diffuse infiltrative lymphocytosis syndrome is an antigen-driven, and MHC-determined, host immune response to an element associated with HIV-1 infection, and (b) that the specific adhesive molecule interactions mediating the cellular influx, as well as the subsequent tissue damage, reflect altered patterns of gene expression in tissues undergoing an immune response. (J. Clin. Invest. 1993. 91:2216-2225.) Key words: diffuse infiltrative lymphocytosis syndrome • CD8 T cells • tissue infiltration • antigen driven • adhesion molecules

### Introduction

Progressive CD4 T cell depletion and incapacitation occurring at various rates in different persons are hallmarks of infection

J. Clin. Invest. © The American Society for Clinical Investigation, Inc. 0021-9738/93/05/2216/10 \$2.00 Volume 91, May 1993, 2216–2225 with HIV-1, however the mechanisms responsible for this impairment are currently not well understood (1). In contrast, T cells of the CD8 lineage are generally not infected with HIV-1 and have been implicated in both potentially protective anti-HIV responses (2, 3) as well as in mechanisms causing the depletion of autologous CD4 T cells (4). CD8 lineage T cells are often moderately elevated early in the course of HIV-1 infection, however, in most cases, this is usually a transient phenomenon and CD8 T cell numbers tend to subsequently decline in parallel with CD4 T cell depletion (5).

Certain adults respond to HIV-1 infection by developing a syndrome characterized by persistent circulating CD8 lymphocytosis, and visceral lymphocytic infiltration, predominantly of the salivary glands and lungs (6). The more prominent clinical features of this diffuse infiltrative lymphocytosis syndrome (DILS)<sup>1</sup> have some similarities to those of classic Sjögren's syndrome and include sicca symptoms, parotid enlargement that is often massive, and, in some individuals, progressive lymphocytic interstitial pneumonitis and renal infiltrative involvement that is manifest as a type IV renal tubular acidosis. Less frequently, other tissues are also infiltrated, leading to manifestations such as periportal inflammatory disease of the liver, dysmotility syndromes of the small and large intestines, and bilateral mastitis. Individuals with DILS have a low rate of progression to frank immune deficiency, with an annual rate of developing an opportunistic infection of only 0.9% (7). Moreover, most of these persons maintain relatively preserved absolute numbers of CD4 T cells, at levels of asymptomatic seropositive individuals, suggesting that their host response may in some way prevent the T cytopathic events characteristic of progressive HIV-1 infection. However, despite persistence of high CD4 T cell numbers, individuals with DILS have a greatly increased risk of developing salivary or lacrimal gland B cell lymphoma (7).

Susceptibility to developing DILS is associated with the presence of particular MHC class II and I alleles. The increase in frequencies of the structurally related MHC class II alleles HLA-DR5 (DRB1\*1102) and HLA-DR6 (DRB1\*1301) (8) supports the notion of DILS being a distinct syndrome and, more importantly, an immunogenetically determined host response to HIV infection. Directing further attention to possible T cell recognition events in DILS that may involve CD8 T cells, there is also a significantly decreased frequency in this disorder of the MHC class I determinant HLA-B35, a specificity that, when present in other HIV-infected persons, has been

Address correspondence to Silviu Itescu, M.D., Columbia University, College of Physicians and Surgeons, Department of Pediatrics, 630 W. 168th Street, BHS 116, New York, NY 10032.

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<sup>1.</sup> *Abbreviations used in this paper:* ARC, AIDS-related complex; DILS, diffuse infiltrative lymphocytosis syndrome; ICAM, intercellular adhesion molecule; LFA, lymphocyte function-associated antigen; NK, natural killer; TCR, T cell receptor.

associated with an increased rate of progression to AIDS-related opportunistic infections (7).

CD8 or CD4 lineage T lymphocytes recognize antigen in the context of MHC class I or II determinants, respectively (9). This interaction and its consequences (10-12), the binding of T cells to and their migration across endothelial surfaces, as well as their homing to specific tissues undergoing immunemediated inflammation (13) depend on adhesive interactions between cell surface structures such as integrins and a variety of ligands including intercellular adhesion molecules (ICAMs) and extracellular matrix components (14, 15). Acquisition of these surface structures, for example, CD29, the  $\beta$  chain common to members of the  $\beta_1$  integrin family, the  $\beta_2$  integrin CD11a/CD18, and the tyrosine phosphatase CD45RO(16), is regulated in a relatively coordinated fashion (17) and leads to a permanent phenotype that distinguishes memory from naive T cells. This sustained phenotype contrasts with the transient expression of markers of recent activation, such as HLA-DR or interleukin-2 receptors. These memory cells demonstrate rapid anamnestic response to recall antigens (11, 16, 18), contain cytotoxic T cell precursors (19), and are capable of enhanced  $\gamma$ -interferon production (17). In particular, CD8 T cells use CD11a/CD18 (lymphocyte function-associated antigen [LFA-1]) molecules to facilitate target adhesion and cytotoxicity (11, 14). The entry of these memory T cells to specific tissues undergoing an immune response is regulated by the cytokine-mediated induction of ICAMs and other adhesion molecules on vascular endothelium (14).

In this study we sought to determine whether the circulating and infiltrative lymphocytosis occurring in DILS has the characteristics of an antigenically driven host response, where these lymphocytes would display the phenotype of memory T cells, possibly with evidence of recent activation. Alternatively, the lymphocytosis could reflect a nonantigenically driven process of T cell lymphoproliferation or the accumulation of cells with unusual phenotypes. Moreover, since HIV-1 was the principal candidate for providing antigen stimulation, we analyzed involved tissues to determine expression and cellular localization of virally encoded gp120 molecules.

#### Methods

Patients. 10 individuals with DILS were studied. All were HIV-1 positive by both ELISA and Western blot analysis. At the time of study none met criteria for AIDS, and eight met criteria for AIDS-related complex (ARC), or stage 3B (20), with fevers, generalized adenopathy, and weight loss. All patients had persistent bilateral parotid gland enlargement for > 6 mo, which was massive in seven. Lymphocytic interstitial pneumonitis was present in four, lymphocytic interstitial nephritis in two, and lymphocytic hepatitis in one, all documented by tissue biopsy. Detailed clinical data describing these patients have been previously reported (21). None had received corticosteroid or other immunosuppressive therapy before this analysis. Controls consisted of 34 individuals, 25 of whom were HIV positive (8 healthy asymptomatic, 8 with ARC, 9 with AIDS) and 9 who were HIV negative. After diagnostic minor salivary gland, pulmonary, or renal biopsy, four of the patients received zidovudine treatment, at dosages 100-200 mg every 6-8 h. This therapy caused > 50% diminution in salivary gland size in each of the four patients with DILS, as early as 5 d after initiating therapy.

Lymphocyte phenotypic analysis. Mononuclear cells from heparinized peripheral blood were separated by centrifugation on Ficoll-Hypaque gradients (Pharmacia LKB Biotechnology, Inc., Piscataway, NJ). Monoclonal antibodies OKT-11 (CD2), 89b1 (CD3), 91d6 (CD4), OKT8 (CD8), LFA-1 (CD11a), MN41 (CD11b), BW209 (CD16), SG157 (HLA-DR), and Tac (CD25) were used in immunofluorescence after conjugation with fluorescein isothiocyanate or fluorescein succinimidyl biotin as described (22). CD29, CD14 (LeuM3), and CD20 monoclonal antibodies, graciously provided by Becton Dickinson Immunocytometry Systems, Mountain View, CA, were directly conjugated with phycoerythrin, while monoclonal antibodies CD45R, CD57 (Leu7), both gifts of Becton Dickinson, and  $\gamma\delta$  T cell receptor (TCR), purchased from T Cell Sciences, Inc., Cambridge, MA, were directly conjugated with fluorescein isothiocyanate. Avidin conjugated with allophycoyanin or phycoerythrin was used in two or three color formats. The percentages of double- and triple-labeled lymphocytes obtained from either peripheral blood or bronchoalveolar lavage were determined by flow cytometry using an instrument equipped with argon and helium neon ion lasers, five parameter electronics, and a logarithmic amplifier (Becton Dickinson). Gating on combinations of low angle forward and 90 degree side scatter defined for each individual by CD3 and CD14 reagents enabled positive selection for lymphocytes and exclusion of monocytes in the analysis. The statistical significance of differences in the medians between groups were calculated using the Mann-Whitney two-tailed test.

*Histopathologic studies*. Minor salivary gland biopsies obtained from all 10 patients, pulmonary parenchymal tissue from two, and one renal biopsy specimen, were fixed, embedded in paraffin, sectioned serially, and stained with hematoxylin-eosin. Minor salivary gland biopsy specimens were graded according to the criteria of Chisholm and Mason (23). All of the tissues were stained and cultured for acidfast bacilli and fungi.

Biotin-avidin immunoperoxidase staining. Four minor salivary gland biopsies of the lower lip and one renal biopsy were placed in embedding medium (O.C.T. compound; Miles Laboratories Inc., Naperville, IL) in airtight plastic capsules, snap frozen in liquid N<sub>2</sub>, and stored at  $-70^{\circ}$ ; 5-µm sections were prepared, fixed in acetone, and stained with CD2, CD3, CD4, CD8, CD14, CD29, CD11a, CD54 (ICAM-1), and HLA-DR monoclonal antibodies, as described (24). Sections were also stained with the monoclonal antibody 91-35 specific for HIV-1 gp120, kindly provided by Dr. F. Celada, New York University Medical Center. The mAb 91-35 does not inhibit CD4-gp120 binding or syncytium formation (25). Appropriate dilutions of monoclonal antibody, usually from ascites samples, or control myeloma protein were incubated with the tissue sections. After rinsing, the sections were treated with biotin-conjugated F(ab')<sub>2</sub> anti-mouse Ig followed by avidin-conjugated horseradish peroxidase (Vector Laboratories, Inc., Burlingame, CA) and substrate 3'-, 3'diaminobenzidine (Sigma Chemical Co., St. Louis, MO). The slides were then counterstained with methylene blue.

# Results

# Immunophenotypic analysis of circulating lymphocyte populations

Enumeration of CD4 and CD8 T-cells. In two-color fluorescence experiments designed to determine the proportion of T cells expressing CD4 and/or CD8 molecules, the median numbers of circulating CD8 cells were found to be significantly higher in those with DILS than in healthy HIV-positive controls, those with ARC, AIDS, or HIV-negative controls, Table I. Median CD4 T cell numbers in individuals with DILS were not different from levels seen in healthy HIV seropositive controls, but were significantly higher than in those with ARC or AIDS. Less than 1% of cells stained for both markers, data not illustrated. Other two-color experiments (not illustrated) showed that at least 90–95% of the expanded CD8 lymphocyte population in individuals with DILS expressed brightly the pan–T cell marker CD3, and < 5% stained positively for  $\gamma \delta$  T

Table I. Increased Circulating CD8 Cells and Relative
Preservation of CD4 Cells in Individuals with DILS

	CD4	P*	CD8	P*
	median no /mm³±SE		median no/ mm³±SE	
DILS				
(n = 10)	370±37		1660±148	
HIV seropositive				
(n = 8)	410±26	NS	721±129	0.003
ARC				
(n = 8)	152±18	0.0013	377±106	0.0007
AIDS				
(n = 10)	66±13	0.0003	389±127	0.0005
HIV seronegative				
(n = 9)	792±62	0.0006	589±55	0.0004

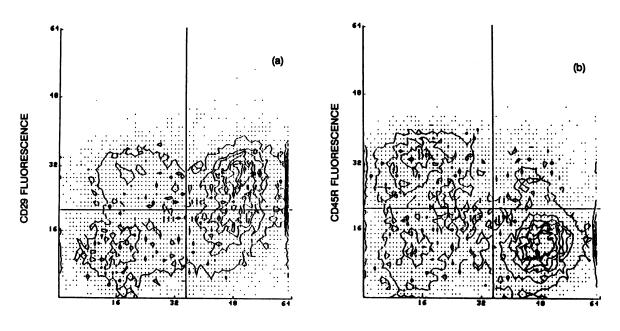
\* Mann-Whitney two-tailed test comparison to individuals with DILS.

cell receptors, implying that the majority of the circulating CD8 cells in individuals with DILS are T lymphocytes using  $\alpha\beta$  receptors.

Delineation of memory phenotype. In triple-staining experiments designed to examine the surface expression of the  $\beta_1$ integrin CD29 and the T200 leukocyte common antigen structure CD45R on circulating CD8 T cells, the major proportion of CD8<sup>+</sup> cells in 9 out of 10 individuals with DILS cells stained moderately to very brightly with CD29 mAbs and dimly to negatively with CD45R mAbs, as shown in Fig. 1. The CD8<sup>+</sup> CD29<sup>bright</sup> CD45R<sup>-</sup> subset accounted for an average of 67% of the total increase in CD8 T cell numbers in these persons. The ratio of CD8<sup>+</sup>CD29<sup>bright</sup> to CD8<sup>+</sup>CD45R<sup>bright</sup> cells, as well as the total numbers of circulating CD8<sup>+</sup>CD29<sup>bright</sup> cells, were significantly higher in those with DILS than in either HIV-negative or HIV-positive controls, Table II.

In other triple-staining experiments, the CD8<sup>+</sup>CD29<sup>bright</sup> cells in all cases studied were shown also to express brightly the  $\beta_2$  integrin molecule CD11a/CD18 (LFA-1), as detected by a CD11a mAb (Table III). A subset of the expanded CD8 population among those with DILS also expressed CD57 (Leu 7) molecules, Table III. Triple staining experiments demonstrated that 41–97% of CD8<sup>+</sup>CD57<sup>+</sup> cells also expressed CD29 molecules brightly (Fig. 2, *inset*), but not natural killer (NK)– associated markers such as CD11b/CD18 and CD16, data not shown. In addition, these cells were of the same size and morphology as other circulating lymphocytes, suggesting that they are not large granular lymphocytes which have been reported to be CD57<sup>+</sup> and to possess NK function (26).

No detectable increase in expression of markers of recent activation. Dual staining experiments demonstrated that, in contrast to markers of memory acquisition, the expanded CD8 lymphocyte population in DILS did not express markers of recent activation, such as MHC class II molecules, Table II, or interleukin-2 receptors, data not illustrated. Similarly, individuals with DILS had significantly lower proportions of CD4 cells expressing CD25 (interleukin-2 receptors) than asymptomatic HIV-positive controls, those with ARC, or those with AIDS, Table IV. Surface expression of HLA-DR molecules by the CD4 T cell population in DILS was also lower than in all other categories of HIV-infected individuals, most significantly in comparison to those with AIDS. In contrast, HLA-DR expression by B cells was preserved in DILS, as evidenced by the CD4<sup>-</sup>DR<sup>+</sup> population in Fig. 3. Indeed, the total numbers of circulating B lymphocytes, defined by CD20 expression, did not differ between individuals with DILS and normal controls. data not shown.



#### CD8 FLUORESCENCE

Figure 1. Three-color immunofluorescence analysis of circulating lymphocytes from an individual with DILS stained with phycoerythrin-, fluorescein-, or biotin-conjugated CD29, CD45R, and CD8 monoclonal antibodies, respectively, illustrating a predominance of CD8 T cells with a memory phenotype. While 59% of the CD8-positive cells expressed CD29 molecules (a), only 11% expressed CD45R molecules (b).

	CD8 <sup>+</sup> CD29 <sup>bright</sup>	P*	CD8 <sup>+</sup> CD45R <sup>bright</sup>	P*	CD29:CD45R	CD8 <sup>+</sup> DR <sup>+</sup>
	median no/mm³±SE (%)		median no/mm <sup>3</sup> ±SE (%)			median %±SE
DILS						
(n = 10)	903±85 (54)		408±57 (25)		1.94	7±1
HIV <sup>+</sup>						
(n = 8)	344±61 (45)	0.004	363±57 (50)	NS	0.92	8±6
ARC						
(n = 8)	175±42 (39)	0.0007	149±51 (40)	0.0083	0.97	16±4
AIDS						
(n = 10)	195±48 (50)	0.0004	120±84 (31)	0.014	1.46	15±4
HIV-						
(n = 9)	322±42 (55)	0.007	327±40 (56)	NS	0.93	7±1

Table II. The Expanded CD8 Population in those with DILS Has a Preponderant Phenotype of 29 bright 45-
but Does Not Express Markers of Recent Activation such as HLA-DR Molecules

\* Mann-Whitney two-tailed test comparison of all numbers to individuals with DILS. () Values in parentheses are percentages of gated lymphocytes.

# Immunohistologic studies

Tissue morphology. Histological examination demonstrated each of the minor salivary gland biopsy samples to have two or more discrete foci of at least 50 lymphocytes per 4 mm<sup>2</sup> of tissue, or grade 4 according to the Chisholm and Mason criteria for diagnosis of Sjögren's syndrome (23). The infiltrating mononuclear cells were pleomorphic in size and shape and did not have any morphologic characteristics of neoplastic lymphoproliferation. In most individuals, the integrity of parenchymal structures remained relatively preserved despite dense periacinar and perivascular focal infiltrates. However, the degree of salivary gland destruction tended to reflect the duration of lymphocytic infiltration, and ranged from complete preservation of glandular architecture to atrophic duct epithelium, canal dilatation, and interstitial fibrosis. Pulmonary parenchymal tissue from patients with lung involvement revealed both focal lymphoid aggregates and diffuse interstitial lymphocytic infiltration. Renal tissue from the two patients with progressive

Table III. The Expanded CD8 Population in those with DILS Primarily Expresses CD11a/CD18(LFA-1) and a Subset Expresses CD57 (LEU7)

	CD8+CD11a/CD18+	P*	CD8+CD57+	P*
	no/mm³(%) median±SE		no/mm³ (%) median±SE	
DILS				
(n = 10)	915±193 (55)		438±112 (26)	
HIV <sup>+</sup>				
(n = 8)	481±92 (66)	0.042	187±77 (26)	NS
ARC				
(n = 8)	232±87 (62)	0.005	112±40 (30)	0.003
AIDS				
(n = 10)	237±157 (61)	0.005	127±46 (33)	0.010
HIV-				
(n = 9)	273±71 (46)	0.002	116±31 (20)	0.005

\* Mann-Whitney two-tailed test comparison to individuals with DILS.

renal insufficiency and type IV renal tubular acidosis revealed interstitial and tubular lymphocytic infiltration without any glomerular involvement.

Lineage of cells infiltrating salivary gland tissue. Lymphocytes accounted for 80-90% of the mononuclear cells infiltrating salivary tissues, while cells of the monocyte/macrophage lineage, defined by staining with a CD14 mAb, accounted for 10-20%. Of the lymphocytes infiltrating the salivary tissues in each of five samples, 80-90% expressed the pan-T cell markers CD2 and CD3, and, in three of three samples, 10-20% of lymphocytes, scattered within the foci, expressed the B cell marker CD20. The predominant population of infiltrating lymphocytes was CD8 positive, accounting for 60-80% of all lymphocytes. 10-15% of the lymphocytes were CD4 positive, as were most of the monocyte/macrophage lineage cells which could be differentiated by their abundance of cytoplasm, characteristic periacinar location, and CD14 expression in consecutive sections. The CD8-positive cells were usually distributed diffusely throughout salivary tissue, within lymphoid aggregates, and surrounding and infiltrating acini. In contrast, the smaller numbers of CD4-positive T cells were primarily located within lymphoid foci.

Memory phenotype of infiltrating lymphocytes. From 75-90% of the infiltrating lymphocytes stained strongly positive with CD29 (not illustrated) and CD11a mAbs (Fig. 4 A). These molecules were expressed most brightly on cells surrounding postcapillary venules and salivary gland acini, while lymphocytes intermediate in location between these sites stained either less intensely or negatively. In four of four biopsy specimens studied, as illustrated in fig. 4 B, CD54 (ICAM-1) expression was observed on endothelial cells of postcapillary venules within lymphoid aggregates, but not in vessels of uninvolved salivary tissue. While this molecule was also strongly expressed by the periacinar monocyte/macrophage lineage cells, CD54 expression was entirely absent on salivary ducts and acini. Staining with mAbs specific for HLA-DR, Fig. 4 C, revealed acini that were DR positive within and adjacent to lymphoid foci, but not at distances away from these aggregates. A small proportion of the lymphocytes within the foci were DR positive, and these were accounted for numerically by CD20-positive B cells in consecutive sections.

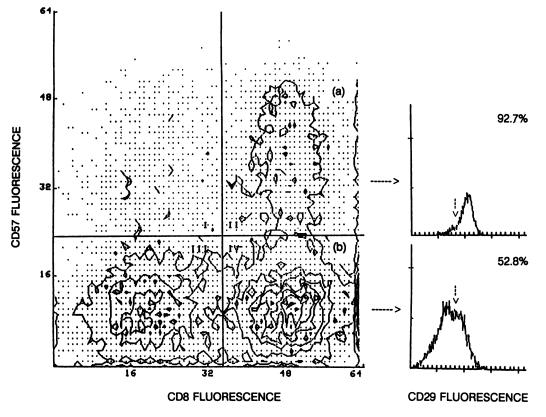


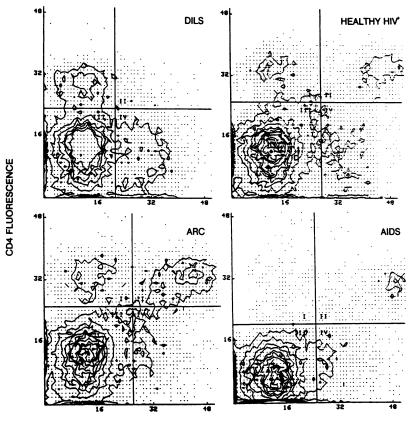
Figure 2. A representative three-color immunofluorescence analysis of circulating lymphocytes from a person with DILS stained with phycoerythrin-, fluorescein-, or biotin-conjugated CD29, CD57, and CD8 monoclonal antibodies, respectively, demonstrating a prominent CD8<sup>+</sup> CD57<sup>+</sup> subset. Whereas 24% of the CD8-positive cells expressed CD57, 76% of the CD57 population was CD8 positive. The CD8<sup>+</sup> CD57<sup>+</sup> population was enriched for expression of CD29, which was present on 92.7% of these cells compared with 52.8% of the CD8<sup>+</sup> CD57 population (inset).

HIV-1 gp120 antigens localized to salivary gland macrophages. The cells which stained positively with CD14 mAbs, and morphologically appeared to be of monocyte/macrophage lineage, were located in periacinar and perivascular areas, as well as adjacent to lymphoid foci, and strongly expressed HLA-DR molecules (Fig. 4 C). In consecutive sections, in three of three cases studied, from 5 to 10% of these monocyte lineage cells demonstrated positive cytoplasmic and cell-surface staining with mAbs specific for HIV-1 gp120 (Fig. 4 D). HIV-1-related antigens were more frequently observed within periacinar monocyte lineage cells at sites distant from lymphoid aggregates. Scanning multiple sections in each tissue sample studied, expression of HIV-1 proteins was very rarely observed in occasional lymphocytes, and no positive staining of acini, ducts, or venules was demonstrable with the gp120 mAb.

Characterization of cellular infiltration in other tissues. Immunohistologic analysis of renal tissue from one individual with interstitial nephritis demonstrated that > 90% of the lymphocytes in the interstitium, primarily surrounding and infiltrating the renal tubules, expressed CD2 and CD3 pan-T cell markers. Over 70% of these cells were CD8<sup>+</sup> and 20% were CD4<sup>+</sup>. More than 75% of the lymphocytes were strongly stained by CD29 and CD11a reagents, while the glomerular endothelium, but not the tubules, expressed CD54 (ICAM-1).

CD4	CD29:CD45R	CD4 <sup>+</sup> CD25 <sup>+</sup>	P*	CD4 <sup>+</sup> DR <sup>+</sup>	P*
no/mm <sup>3</sup> mm <sup>3</sup> ±SE	median±SE	median%±SE		median%±SE	
370±37	4.35	9±2		27±4	
410±26	2.27	36±5	0.006	29±6	NS
152±18	2.31	39±8	0.0082	45±6	NS
66±13	1.82	36±6	0.003	64±9	0.021
792±62	1.08	8±1	NS	14±2	0.011
	no/mm <sup>3</sup> mm <sup>3</sup> ±SE 370±37 410±26 152±18 66±13	no/mm³ median±SE   370±37 4.35   410±26 2.27   152±18 2.31   66±13 1.82	no/mm³ mm³±SEmedian±SEmedian%±SE $370\pm37$ $4.35$ $9\pm2$ $410\pm26$ $2.27$ $36\pm5$ $152\pm18$ $2.31$ $39\pm8$ $66\pm13$ $1.82$ $36\pm6$	no/mm³ mm³±SEmedian±SEmedian%±SE $370\pm37$ $4.35$ $9\pm2$ $410\pm26$ $2.27$ $36\pm5$ $0.006$ $152\pm18$ $2.31$ $39\pm8$ $0.0082$ $66\pm13$ $1.82$ $36\pm6$ $0.003$	no/mm³ mm²±SEmedian±SEmedian%±SEmedian%±SE $370\pm37$ $4.35$ $9\pm2$ $27\pm4$ $410\pm26$ $2.27$ $36\pm5$ $0.006$ $29\pm6$ $152\pm18$ $2.31$ $39\pm8$ $0.0082$ $45\pm6$ $66\pm13$ $1.82$ $36\pm6$ $0.003$ $64\pm9$

\* Mann-Whitney two-tailed test comparison to individuals with DILS.



HLA-DR FLUORESCENCE

Figure 3. Two-color immunofluorescence analysis of circulating lymphocytes from four HIV-infected individuals stained with fluorescein- or biotin-conjugated CD4 or HLA-DR monoclonal antibodies, respectively, indicating the low level of CD4 T cell activation. While only 9% of the CD4 cells in the individual with DILS expressed HLA-DR molecules, the proportion of activated CD4 cells was 52% in the asymptomatic HIV-positive individual, 63% in the one with ARC, and 74% in the individual with AIDS.

In another individual with DILS and biopsy-proven lymphocytic interstitial pneumonitis, 56% of the lymphocytes obtained by bronchoalveolar lavage were T cells, of which > 96% were CD8<sup>+</sup> and bore  $\alpha\beta$  T cell receptors. 73% of these cells expressed brightly CD29 and CD11a/CD18 structures, while only 10% were positive with CD45R mAbs. Only 11% of the CD8 cells in double label experiments were activated as defined by the presence of interleukin-2 receptors (CD25) and MHC class II molecules, data not illustrated.

#### Discussion

These studies suggest that both the circulating and infiltrative CD8 lymphocytosis in individuals with DILS are components of an antigenically driven, and immunogenetically determined, systemic host response to HIV-1 infection. These conclusions are based on the following major points: (a) The expanded circulating lymphocyte population in persons with DILS primarily consisted of CD8 T cells with a memory phenotype, characterized by increased surface expression of CD11a/ CD18 (LFA-1), CD29 (very late arising-4 [VLA]  $\beta_1$  chain), and, to a lesser extent, CD57 (Leu 7) molecules. These cells did not express markers of recent activation such as interleukin-2 receptors or HLA-DR molecules. (b) The predominant lymphocytic population that infiltrated salivary, pulmonary, or renal tissues was accounted for by CD8 T cells with a similar memory phenotype to that found in the blood. (c) Strong expression of CD54 (ICAM-1) molecules was observed on postcapillary venule endothelium within the lymphoid aggregates, but not on endothelium at sites distant from the foci. This suggests that the entry of lymphocytes into tissues involves interactions between specific adhesion molecule receptor-ligand pairs such as LFA-1/ICAM-1. (d) Expression of HIV-encoded proteins was demonstrable in salivary gland monocyte/macrophage lineage cells which were usually located in periacinar and perivascular locations in close proximity to lymphoid aggregates, and strongly expressed MHC class II molecules.

CD8 T cells bearing the same memory phenotype as identified in this report have been shown to suppress HIV-1 replication in humans by mechanisms requiring cell-cell interaction and MHC-restriction (27). Accordingly, it may be that individuals with DILS selectively expand a circulating population of CD8 T cells capable of suppressing HIV replication in vivo. In support of this possibility are the relatively preserved levels of CD4 T cells in this group, the low rate of progression to opportunistic infections (7), and the low expression of HIV-1 proteins in DILS peripheral blood, as determined by circulating p24 antigenemia (21). These observations imply that events in HIV-1 infection which normally result in CD4 T cell depletion and progression to opportunistic infections are attenuated or retarded in DILS. The low frequency of activated CD4 T cells in these individuals, as defined by surface expression of interleukin-2 receptors and MHC class II molecules, is of interest in view of the association between T cell activation and the transactivation of HIV-1 gene expression (28).

Because a substantial portion of the expanded CD8 population in patients with DILS expressed CD57 (Leu 7) molecules in addition to the  $\beta$ -integrins, this prominent cell type could be an important element of these individuals' cellular response to

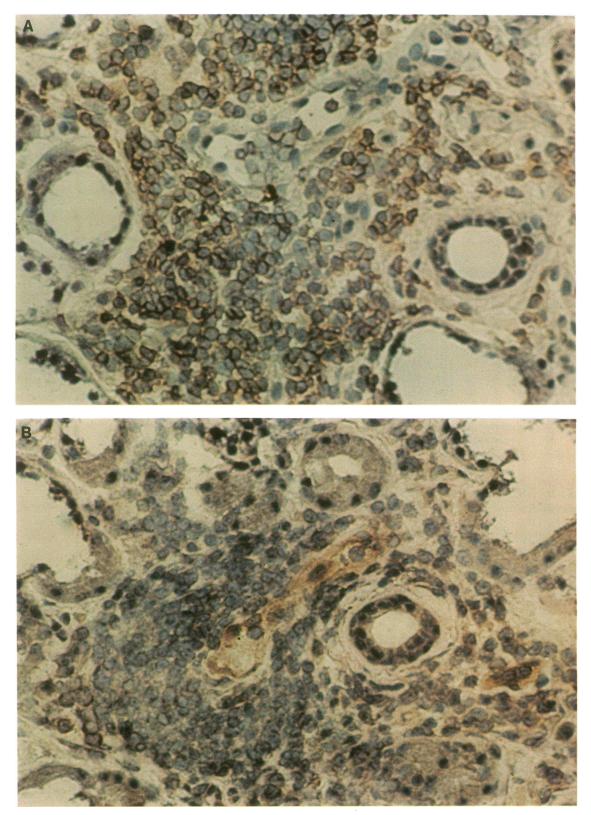


Figure 4. Immunohistochemical studies of minor salivary gland specimen from an HIV-infected individual with DILS demonstrating expression of the  $\beta_2$ -integrin CD11a/CD18 (LFA-1) by infiltrating lymphocytes (A), CD54 (ICAM-1) expression by postcapillary venule endothelial cells within lymphoid focus (B), and HLA-DR expression by acini, but not ducts (C), in the vicinity of monocyte/macrophage lineage cells that also expressed HLA-DR (C) and HIV-1 gp120 molecules (D).

HIV. Increased numbers of CD8<sup>+</sup>CD57<sup>+</sup> cells have previously been described in HIV-infected individuals at various stages of disease (29, 30), however their function remains unknown. In vitro, CD8<sup>+</sup>CD57<sup>+</sup> cells have been demonstrated to be capable of suppressing pokeweed mitogen-induced B cell antibody production (26), and of mediating lectin-induced cytotoxicity

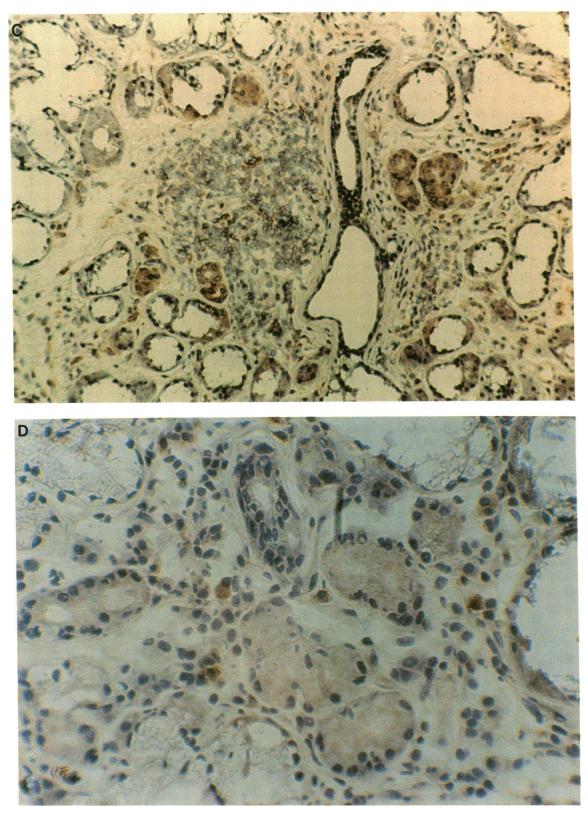


Figure 4 (Continued)

(31). Although CD57 is a marker present on certain NK cells, the CD8<sup>+</sup> CD57<sup>+</sup> cells present in DILS patients are unlikely to be NK cells because they possess the CD3 T cell receptor complex, lack other NK surface markers such as CD16 (Fc receptors) and CD11b/CD18 (CR3), and cells with this phenotype

in other HIV-infected individuals have been shown to have little NK function (30).

In support of an antigen-driven basis to the CD8 lymphocytosis in DILS, analysis of the T cell receptor  $\alpha$  and  $\beta$  chain repertoire in lymphocytes infiltrating the minor salivary glands has revealed selective V and J gene segment usage and a preference for specific somatically encoded amino acid residues in the third complementarity determining region (32). These findings suggest that the infiltrative T cell population is significantly restricted by virtue of antigen-driven clonal selection. While the exact nature of the inciting antigen in DILS remains to be determined, the present demonstration of HIV-1 proteins in salivary gland monocyte lineage cells, as well as the marked diminution in parotid gland enlargement and degree of lymphocytic tissue infiltration after therapy with the reverse transcriptase antagonist zidovudine, likely implicates HIV-1 as the causal agent. However, we cannot exclude the less likely possibilities that the infiltrating lymphocytes may alternatively be reacting to endogenous or exogenous structures whose expression is regulated by the local presence of HIV-1, or even to other retroviruses present at these sites.

The results presented in this report suggest the following hypothetical model for the HIV-host interactions occurring in DILS. Monocytes infected with HIV-1 leave the circulation and localize at various mucosal sites where they differentiate into resident tissue macrophages and induce clinical complications such as lymphocytic interstitial pneumonitis or central nervous system disease (33). The anatomic localization of these monocytes to salivary and pulmonary tissues, gastrointestinal tract, and renal tubules, which occurs for unknown reasons, may account for the particular pattern of organ involvement that is characteristic of DILS. Viral persistence in these cells is associated with the secretion of a number of cytokines, including TNF- $\alpha$ , which is a very potent inducer of ICAM-1 expression on endothelial cells (11, 14). Alone or in combination with other cytokines, TNF- $\alpha$  may induce the enhanced local expression of adhesion molecule ligands by postcapillary venule endothelium in the salivary glands and other tissues.

In those immunogenetically distinct individuals who are capable of responding to HIV-1 infection by developing a reactive circulating CD8 lymphocytosis, memory CD8 T cells are first primed by exposure to HIV-1 within lymphatic tissues. These lymphocytes express selective homing receptors and are sequestered into salivary glands, lungs, and other organs by interactions between accessory molecules on lymphocytes such as the integrins and those on endothelial cells such as ICAM-1. It is possible that immune recognition events occurring between TCR structures on these memory CD8 T cells and HIV peptide/MHC complexes on the tissue macrophages may lead, by as yet undefined mechanisms, to the suppression of HIV-1 replication in the infected cells. The elaboration of cytokines such as  $\gamma$ -interferon during this immune interaction would, in conjunction with factors released by the infected macrophages, lead to additional secondary parenchymal damage.

Similar mechanisms to those occurring in HIV-1 infection, involving abnormal tissue expression of adhesion molecules, may be responsible for the tissue infiltration and damage that occur in other disorders associated with the sicca syndrome, including HTLV-1 infection, chronic graft-versus-host disease, and idiopathic Sjögren's syndrome. Although each of these disorders differs from DILS and from each other in many respects, the present results suggest that the characteristic pattern of T cell recruitment into the same organs in each disorder may reflect a common final pathway. A further factor contributing to the tissue specificity of the infiltrative process in these disorders might be the presence of regulatory cofactors at involved sites. In DILS, for example, HIV-1 replication in salivary tissues may be enhanced by human herpes virus 6 and epidermal growth factor, two cofactors that are principally located in the salivary glands (34, 35), and that are, respectively, capable of transactivating HIV-1 (36) and increasing HIV-1 replication in infected cell lines (37).

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