Immunopathology and Infectious Diseases

Host Defense Mechanisms in Secondary Syphilitic Lesions

A Role for IFN- γ /IL-17-Producing CD8⁺ T Cells?

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Cell-mediated immunity is thought to be of critical importance in antisyphilitic host defense, but the exact mechanisms are still unknown. This fact is particularly true for HIV-infected persons with a deficit in CD4⁺ T-cell number. We therefore obtained lesional skin samples from HIV⁺ and HIV[−] patients with secondary syphilis at different time points of lesional age to search both for causative microorganisms and to characterize the inflammatory infiltrate. By doing so, we detected *Treponema pallidum* spirochetes with a much greater abundance in late lesions of HIV⁺ individuals compared with the HIV[−] cohort. The dominating inflammatory cells were T cells, macrophages, and neutrophils at all stages and plasma cells in older lesions. In HIV[−] persons, T cells consisted of equal numbers of CD4⁺ and CD8⁺ T-cells, whereas in HIV⁺ patients, the majority of T cells belonged to the CD8 lineage and produced both IFN- γ and IL-17. Regulatory T cells and Langerhans cells were reduced in these patients compared with their HIV[−] counterparts. Because of our observations, we propose that T cells of both the CD4 and CD8 lineage are needed for an at least partial protective antisyphilitic immunity. Compensation mechanisms in HIV⁺ individuals, such as an increase of Tc1/17 cells as well as a reduction in immunoregulatory Langerhans cells and T cells, apparently do not overcome the deficiencies in these patients to eliminate the spirochete. (Am J Pathol 2010, 177:2421–2432; DOI: 10.2353/ajpath.2010.100277)

Syphilis is a multistage sexually transmitted disease that continues to be a public health problem. Natural infection with *Treponema pallidum* subsp. *pallidum* (*T. pallidum*) results in the formation of a primary lesion at the entry site of the bacteria and, after dissemination of the microorganisms to different tissues, in secondary manifestations. A vigorous immune response develops early during infection and causes local clearance of the majority of treponemas. In less than one third of untreated syphilitic patients, the infection can progress after a latency phase to tertiary syphilis with skin, bone, cardiovascular, and/or neurological involvement.¹

The inability to cultivate *T. pallidum in vitro*, coupled with difficulties to establish suitable animal models for immunological studies, has greatly hindered our understanding of the basic aspects of syphilis. Evidence for the importance of a strong cellular immunity evolved first from studies in a rabbit model of syphilis, as i) *T. pallidum* can induce an antigen-specific proliferation of T-cells a few days to months after infection,^{2,3} ii) activated macrophages and the infiltration of mononuclear cells at the site of inoculation of the bacteria correlate with *T. pallidum*-reactive T-cells,^{4,5} and iii) high levels of IL-2 and IFN- γ can be detected after incubation of splenocytes with particles of *T. pallidum*.³ In human skin samples of primary and secondary syphilis, the dominance of T-cells and macrophages and IL-2, IFN- γ , and IL-12 as classical Th1 cytokines has been demonstrated.^{6,7} However, the exact cellular subset responsible for pathogen clearance has yet to be found. CD4⁺ T-cells are believed to mediate the eradication of treponema primarily through the production of cytokines, such as IFN- γ , thereby activating

The work for this article was performed at the Division of Immunology, Allergy and Infectious Diseases of the Department of Dermatology, Medical University of Vienna, Vienna, Austria without additional funding.

Accepted for publication July 1, 2010.

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macrophages,⁸ which presumably then kill opsonized spirochetes.^{9,10} Other studies suggest that activated CD8⁺ T-cells participate in the local pathogen clearance.^{6,11} Recently published data of suction blisters of syphilitic skin lesions and peripheral blood mononuclear cells show that *T. pallidum* simultaneously elicits also innate immune responses,^{12,13} as evidenced by the direct activation of macrophages by *T. pallidum*-derived TLR2 ligands.¹⁴

In this study we aimed at carefully characterizing the tissue-based inflammation during secondary syphilis and at defining candidate effector cells in pathogen clearance.

Materials and Methods

Patients and Tissue Material

Twenty-five patients with serologically confirmed secondary syphilis and cutaneous involvement (2 females, 23 males; mean age 49.7 ± 16.2) were enrolled in the study. All patients were recruited in the STD and HIV outpatient center of the Department of Dermatology, Medical University of Vienna. Diagnoses were made by trained physicians and with laboratory confirmation. All patients were serologically positive for the *Treponema pallidum* particle-agglutination assay (TPPA), fluorescent treponemal antibody absorbed (FTA-ABS), and Venereal Disease Research Laboratory (VDRL) tests. Routinely, the patients' history with the time of lesional persistence and the HIV-1 status were assessed and documented. Thirteen of 25 patients were HIV-infected (12 men, 1 woman; mean age 44 ± 9 years, range 36–62 years; mean CD4⁺ cell count: 465.9 ± 195.4 cells/ μ l, range 228–808 cells/ μ l; mean viral load: $182,975 \pm 275,459$ copies/ml, range 50–724,436 copies/ml). Three HIV⁺ individuals received antiretroviral therapy resulting in a viral load of less than 50 RNA copies/ml plasma. According to the CDC classification of HIV infection two patients belonged to category A1, six to A2, four to A3, and one to B2 at the time point of diagnosis. Syphilitic skin manifestations and locations were classified as localized papules (eg, palmoplantar, genital, facial papules, condylomata lata, plaques muqueuses) and/or as generalized rash on trunk and limbs. Skin biopsies were taken from the trunk ($n = 14$), from upper extremities ($n = 7$), or from lower extremities ($n = 4$) before antibiotic therapy. From each patient a punch biopsy was taken under local anesthesia and divided into two or three pieces for formalin and trizol fixation and, in 14 cases, for embedding in optimal cutting tissue (OCT) compound (Tissue-Tek) and subsequent deep-freezing in liquid nitrogen. Trizol-fixed and deep-frozen tissue specimens were stored at -20°C and -80°C , respectively, until further processing. Normal skin samples from healthy individuals ($n = 10$) were used as the control group.

The study was performed according to the Declaration of Helsinki. All study subjects participated voluntarily and gave informed consent; the study was approved by the Ethics Committee of the Medical University of Vienna (EK 721/2008).

Histopathological Evaluation

A routine hematoxylin and eosin (HE) staining was performed on all formalin-fixed biopsy specimens ($n = 25$) and evaluated blindly by one of the co-authors (P.P.) with a scoring system for the following panel of histopathological features: epidermal acanthosis (0 denotes $<0.15 \mu\text{m}$, 1 denotes $0.15\text{--}0.3 \mu\text{m}$, 2 denotes $>0.3 \mu\text{m}$); dyskeratosis of keratinocytes (0, absent; 1, present); occurrence of plasma cells, macrophages, and neutrophils (0, absent; 1, single; 2, $<20\%$ of inflammatory cells; 3, $>20\%$ of inflammatory cells); nodular infiltrate (0, absent; 1, present); vasculitis (occlusion of vessels by fibrin and erythrocytes: 0, absent; 1, present); endothelial cell proliferation (0, absent; 1, present); extravasation of red blood cells (0, absent; 1, present).

Quantitative RT-PCR

Trizol-fixed frozen biopsies ($n = 25$) were thawed on ice and then homogenized with a FastPrep-24 homogenizer (MP Biomedicals; Solon, OH) for RNA isolation according to the manufacturer's instructions. The RNA concentration was determined with a NanoDrop ND-100 spectrophotometer (Peqlab Biotechnology; Erlangen, Germany) followed by the removal of DNA by TURBO DNase (Applied Biosystems; Foster City, CA) treatment according to the manufacturer's guidelines. The reverse transcription of RNA to cDNA was performed with a High-Capacity cDNA Reverse Transcription Kit (Applied Biosystems) according to the standard protocol for a $20 \mu\text{l}$ reaction on a Thermocycler Perkin Elmer 480 (Perkin Elmer; Waltham, MA). The reactions were amplified and quantified by a TaqMan Universal PCR Master Mix (Applied Biosystems) on an Applied Biosystems ABI PRISM 7700 using the following thermal cycler conditions: 2 minutes at 50°C ; 10 minutes at 95°C ; and 40 cycles of 15 seconds at 95°C followed by 60 seconds at 60°C . The human β -2-microglobulin (β 2m) gene (Hs00187842_m1, Applied Biosystems), a housekeeping gene, was used to normalize each sample and each gene. For the primers and probes for the TaqMan RT-PCR assays we used TaqMan Gene Expression Assays (all Applied Biosystems): IL-12p35, Hs01073447_m1; IL-12/IL-23p40, Hs01011518_m1; IL-23p19, Hs00413259_m1; IFN- γ , Hs00174143_m1; IL-4, Hs00929862_m1; IL-13, Hs99999038_m1; IL-18, Hs99999040_m1; IL-17, Hs00174383_m1; IL-22, Hs01574154_m1; IL-10, Hs00961622_m1; and TGF- β , Hs00171257_m1. The data were analyzed and samples were quantified by software provided with the Applied Biosystems ABI PRISM 7700 (Sequence Detection Systems, ver. 1.7) and calculated using the comparative C_T method with β 2m as reference gene relative to normal skin. For quantitative analysis of *T. pallidum* amplification of the 16S-rDNA *T. pallidum* gene was performed (accession number of the GenBank NCBI: AE000520.1).

Table 1. List of Antibodies Used in this Study

Antibody specificity	Clone (source of antibodies)	Isotype
BDCA-2 (pur.)	124B3.13 (Dendritics, Lyon, France)	mIgG1
CD1a (FITC)	HI149 (BD Bioscience, Schwenchat, Austria)	mIgG1
CD1c (pur.)	M241 (Ansell, Bayport, MN)	mIgG1
CD3 (FITC)	SK-7 (BD Bioscience)	mIgG1
CD4 (pur.)	SK-3 (BD Bioscience)	mIgG1
CD8 (pur.)	C8/144B (DakoCytomation, Glostrup, Denmark)	mIgG1
CD8 (APC)	B9.11 (Immunotech Coulter, Brea, CA)	mIgG1
CD11b (FITC)	BEAR1 (Beckman Coulter, Brea, CA)	mIgG1
CD14 (pur.)	MΦP9 (BD Bioscience)	mIgG2b
CD19 (pur.)	SJ25C1 (BD Bioscience)	mIgG1
CD25 (pur.)	2A3 (BD Bioscience)	mIgG1
CD56 (biot.)	B158 (BD Bioscience)	mIgG1
CD123 (biot.)	9F5 (BD Bioscience)	mIgG1
CD138 (pur.)	MI15 (BD Bioscience)	mIgG1
CD207 (pur.)	DCGM4 (Beckman Coulter)	mIgG1
FoxP3 (biot.)	236A/E7 (eBioscience, San Diego, CA)	mIgG1
HLA-DR (FITC, APC)	L243 (BD Bioscience)	mIgG2a
IFN- γ (biot.)	B27 (Abcam, Cambridge, MA)	mIgG1
IL-17A (pur.)	eBio64DEC17 (eBioscience)	mIgG1
<i>T. pallidum</i> (pur.)	Rabbit polyclonal Ig to <i>T. pallidum</i> (Abcam)	Rabbit Ig polyclonal

biot., biotinylated; pur., purified; APC, allophycocyanin.

Immunofluorescence and Immunohistological Staining

Single and multicolor immunofluorescence staining procedures were performed on frozen sections of patients with secondary syphilis ($n = 14$) and healthy controls ($n = 10$) as previously described.¹⁵ The mAbs used in this study and their sources are shown in Table 1. Cytokine stainings for IFN- γ and IL-17 were performed using a tyramide signal amplification (TSA) system (PerkinElmer, Waltham, MA) according to the manufacturer's suggestions. Negative controls were obtained in all staining experiments by substituting isotype-matched IgG or IgM (purified, APC-, or FITC-labeled) for the primary antibody. For the evaluation of immunofluorescence results biopsy specimens were read in a blinded fashion by two independent investigators with a mean observer coefficient of $<10\%$. Labeled cells were enumerated per visual field and expressed as the number of epidermal cells per millimeter basement membrane as well as of cells per millimeter² dermis $\pm$ SEM.

Paraffin sections of all 25 patients were used to visualize *T. pallidum* by immunohistological stainings after paraffin dewaxing and antigen retrieval. To determine the distribution and the number of *T. pallidum* in lesional skin, single immunostainings were performed with a polyclonal rabbit-anti-*T. pallidum* antibody (Abcam). Substitution of the antibody with concentration-matched rabbit immuno-

globulins (DAKO) served as negative control. To prevent nonspecific protein binding, antibodies were mixed with normal horse serum and 2% BSA in PBS (Gibco, Carlsbad, CA) before being applied to the sections at 4°C overnight. After incubation of the slides with biotinylated horse anti-rabbit IgG (Vector Laboratories, Burlingame, CA) at room temperature for 45 minutes, the sections were rinsed and incubated with avidin-biotin peroxidase complex (Vector Laboratories) at room temperature for 45 minutes. For visualization of the cells, 0.01% 3-amino-9-ethyl-carbazol in 50 mmol/L acetate buffer (pH 5, 0.015% H₂O₂, 5% dimethylformamide) was used as chromogen. Sections were counterstained with hematoxylin (Merck, Darmstadt, Germany) and mounted in Aquamount (BDH Laboratory Supplies, Poole, UK). Treponemas were enumerated as spiral or punctual structures per visual field and expressed as the number treponemas per millimeter basement membrane as well as of treponemas per millimeter² dermis $\pm$ SEM.

Statistical Analysis

The significance of the differences in cellular and treponema numbers between HIV⁺ and HIV⁻ patients was verified by the nonparametric Mann-Whitney *U* test. The levels of significance of data from quantitative RT-PCR analysis were assessed by the parametric Students *t*-test. A *P* value of <0.05 was considered as statistically significant.

Results

Clinical Characterization of Secondary Syphilitic Lesions According to the Duration of Their Persistence

Twenty-five patients with secondary syphilis, 13 thereof being HIV-positive, were classified into four categories according to the age of their lesions: six patients with an early rash being present for not longer than 10 days; six individuals with a rash of 11–20 days duration; seven persons with skin symptoms persisting for 21–30 days, and, finally, six patients with papules having appeared 30 days ago or even longer. Differences in the clinical presentation resulted from the duration of lesional persistence: a slightly erythematous maculo-papular rash without apparent scaling was typical for early lesions of less than 10 days duration (Figure 1A); when persisting longer, the skin manifestations became more infiltrated and exhibited a violaceous color and showed signs of epidermal involvement (Figure 1B). With an age of >20 days, the individual papules had often coalesced to scaling plaques that often displayed a psoriasiform appearance and configuration (Figure 1, C and D). In general, the extent of skin involvement was more pronounced in HIV⁺ patients (table 2).

Figure 1. Correlation of clinical symptoms and lesional age of secondary syphilitic eruptions. Patients were arbitrarily divided into four groups according to the length of persistent lesions [(A) 10 days or less, $n = 6$; (B) 11 to 20 days, $n = 6$; (C) 21 to 30 days, $n = 7$; (D) more than 30 days, $n = 6$]; type as well as severity of these lesions were recorded. The clinical pictures are representative of the indicated times.

Histopathological Findings of Syphilitic Lesions: A Time-Dependent Process

Histopathological examination of the 25 secondary syphilitic lesions revealed a pattern that correlated with the duration of the lesions (Figure 2, A–H), but not with the HIV status of the patients. In lesions existing for fewer than 10 days, the typical histopathological picture was a sparse perivascular lymphocytic infiltrate, with few neutrophils and macrophages admixed, but devoid of plasma cells, in the upper dermis and an overt vacuol-

ization of the basal cell layer of the epidermis mimicking multiforme-like drug eruptions (Figure 2, A and G). Plasma cells were first found 10–20 days after appearance of the rash and gradually increased in numbers with the age of the lesion (Figure 2G). At this early stage, the differential diagnosis with pityriasis lichenoides may be difficult, especially in cases when plasma cells are still few in number. On longer duration of the lesions, the infiltrate became nodular and involved the superficial and deep vascular plexus (Figure 2, B–D and H). In addition to lymphocytes and neutrophils, the plasma cell became a dominant component (Figure 2G). Subsequently, by the admixture of macrophages, the infiltrate exhibited a granulomatous appearance (Figure 2, C, D, and H). In papules of more than 20 days duration a psoriasiform pattern was observed, manifested by epithelial acanthosis, papillomatosis, and parakeratosis as well as dyskeratotic cells within the basal and suprabasal layers (Figure 2, C–F). In early stages the epidermis contained lymphocytes, whereas the psoriasiform hyperplastic epithelium of lesions with more than 20 days duration also contained neutrophils, thereby mimicking psoriasis or epidermomycosis (Figure 2, A–D). A very important histological feature is the involvement of postcapillary venules, which is seen after day ten (Figure 2, A–D). Vessels were occluded by erythrocyte thrombi or even by fibrin thrombi, the endothelial cells were swollen and contained prominent nuclei. Such vessels were always surrounded by extravasated red blood cells. However, this type of vessel involvement can be clearly distinguished from leukocytoclastic vasculitis, where fibrin-occluded vessels are permeated and surrounded by neutrophils and nuclear dust.

IFN- γ as Key Cytokine in Syphilitic Lesions: Differences between HIV⁺ and HIV[−] Patients

Although we could not detect transcripts for the classical Th1 cytokines IL-12p35 or IL-18, quantitative RT-PCR revealed a strong up-regulation of IFN- γ mRNA in syphilitic skin lesions as compared to healthy skin (Figure 3A). Our studies further revealed a strong “Th17” signature, as evidenced by elevated levels of IL-23p19, IL-17, and IL-22 mRNA (Figure 3B). In contrast, stage 2 syphilitic lesions were essentially devoid of transcripts for the Th2 cytokines IL-4 and IL-13 (Figure 3C). When we analyzed the lesional cytokine levels with regard to the patients’ HIV status and time of lesional persistence, we observed significant differences between HIV⁺ and HIV[−] patients (Figure 3, A–C) but not between lesions of different duration (Figure 3, D–F). IFN- γ expression was more than doubled in HIV-infected individuals (Figure 3A), and IL-23p19 as well as IL-17 and IL-22 transcripts were several fold more up-regulated in HIV⁺ patients as compared to HIV[−] individuals. We further observed a tendency for reduced levels of the regulatory cytokines IL-10 and TGF- β in HIV⁺ patients as compared to HIV[−] individuals (Figure 3C). These data suggest that T-cells producing cytokines originally described for Th1 and Th17 cells are operative against *T. pallidum* in secondary syphilitic le-

Table 2. Extent of Skin Involvement in HIV⁺ and HIV[−] Syphilitic Patients

	HIV-positive ($n = 13$)	HIV-negative ($n = 12$)
Localized papules in ≤ 2 sites	5	6
Localized papules in > 2 sites	6	2
Generalized rash	10	7
Localized and generalized manifestations	8	3

Figure 2. Histopathological correlation with the lesional age of secondary syphilitic skin eruptions. Histopathological changes are displayed analogous to clinical alterations in four groups according to lesional persistence. Representative pictures with a magnification $\times 100$ and $\times 400$ are shown for lesional persistence of four different times: (A) 0–10 days ($n = 6$), (B) 11–20 days ($n = 6$), (C) 21–30 days ($n = 7$), and (D) more than 30 days ($n = 6$). Quantifications of epidermal and dermal alterations are shown in E–H.

sions and that such an immune response is amplified in HIV-infected patients.

Composition of the Leukocyte Infiltrate in Syphilitic Lesions of HIV⁺ and HIV[−] Patients

Of 25 syphilitic patients that were included into the study, we obtained snap frozen tissue specimens from 14 patients, seven thereof HIV⁺, that were subjected to immunofluorescence analysis using a broad panel of antibod-

ies listed in Table 1. Normal skin from 10 healthy donors served as control (healthy controls, HC). In stage 2 syphilitic lesions we found the infiltrate to be dominated by CD3⁺ cells with comparatively higher numbers in HIV-infected patients. This increase of CD3⁺ cells in HIV⁺ individuals affected mainly the CD8⁺ T-cell subset (Figure 4, A and B). Regulatory T-cells, visualized by CD25/FoxP3/CD3 triple stainings, were increased in the dermis of HIV[−] patients as compared with normal skin from HC and HIV⁺ individuals (Figure 4, C and D).

Figure 3. Quantitative RT-PCR identifies IFN- γ as master cytokine for secondary syphilitic lesions. Quantitative RT-PCR was performed after trizol fixation of whole biopsies from secondary syphilitic lesions and analyzed for (A and D) so-called Th1 cytokines, (B and E) Th17 cytokines, and (C and F) Th2 and regulatory cytokines. Samples were subdivided according (A–C) the patients' HIV status and (D–F) the time of persistence of the lesions. Data are normalized to β 2-microglobulin of each specimen and represent the mean values of fold changes relative to healthy skin $\pm$ SEM. * $P < 0.05$.

Other constituents of the syphilitic infiltrate included CD138⁺ plasma cells and small numbers of CD3⁺CD56⁺ NK cells, while CD19⁺ B cells and double-negative T-cells as well as inflammatory-type myeloid DCs and plasmacytoid DCs were almost absent (Figures 4A and 5A). Interestingly, CD207⁺CD1a⁺ epidermal Langerhans cells were strongly decreased (Figure 5, A and B) and CD14⁺CD11b⁺ dermal mononuclear phagocytes sharply increased in HIV-infected patients (Figure 5, A and C) as compared with HC or HIV⁻ patients. When analyzing differences of the cellular infiltrate according to the lesional age, we found a rather selective increase of plasma cells in syphilitic lesions persisting for more than 20 days (Figure 4E), while other leukocyte subsets remained unchanged (Figures 4E and 5D). When we subdivided samples from HIV⁺ and HIV⁻ syphilitic patients according to the lesional persistence (>20 days and <20 days), we did not find statistically significant differences (data not shown). However, limited specimen numbers per group (3–4 samples each) reduce the validity of this analysis.

CD8⁺ T-cells as Main Source for IFN- γ and IL-17 in Syphilitic Lesions

To determine the cellular source of those effector cytokines whose expression was increased at the transcriptional level in syphilitic lesions, we performed multicolor immunofluorescence stainings using mAbs against the respective cytokines and defined leukocyte subpopulations. We found T-cells to be the main source of both IFN- γ and IL-17 (Figure 6, A and B). Somewhat to our

surprise, we only rarely observed CD4⁺ T-cells to display anti-IFN- γ and anti-IL-17 reactivity but rather identified CD8⁺ T-cells as the main producers of both cytokines (Figure 6, A and B). Interestingly, anti-IL-17 reactivity was occasionally found to overlap with anti-IFN- γ staining (Figure 6C). Both IFN- γ and IL-17 were stronger expressed in lymphocytes of HIV⁺ patients as compared with HIV⁻ individuals (Figure 6D), which is consistent with the data obtained at the transcriptional level. These CD3⁺CD8⁺IFN- γ ⁺IL-17⁺ cells were localized within the upper dermal infiltrate in close contact to other leukocytes.

Local Host Response of HIV⁺ Patients Fails to Control *T. pallidum* in Late Lesions

Antitreponema immunostainings of paraffin-fixed sections revealed the presence of *T. pallidum* in the epidermis and dermis of syphilitic lesions of all 25 patients. Depending on whether the spirochetes were encountered horizontally or vertically they appeared as spiral or punctual structures (Figure 7A). Quantification of *T. pallidum* by immunohistochemistry revealed a significant increase of spirochetes within the dermis of HIV-infected patients (Figure 7B). This result was substantiated when we subjected tissue specimens of lesions from HIV⁺ and HIV⁻ patients to *T. pallidum*-specific quantitative RT-PCR analysis (Figure 7D). Combined evaluation of the age of the lesions and the HIV status of the patients disclosed a sharp increase of the causative pathogens of syphilis only in late lesions of HIV-infected patients, while the number of spirochetes in early lesions was comparably

Figure 4. Alterations of the lymphocytic infiltrate in syphilitic lesions of HIV⁺ and HIV⁻ patients. **A:** Detailed quantitative *in situ* analysis of lymphocytes within the epidermis and dermis of secondary syphilitic lesions and normal skin from healthy controls (HC). Single and double immunofluorescence staining was performed with the indicated markers from HIV⁺ and HIV⁻ patients. Data are given as absolute numbers of positive cells $\pm$ SEM. **B:** Triple immunofluorescence labeling of syphilitic lesions from HIV⁺ and HIV⁻ patients with anti-CD3 (FITC), anti-CD4 (TRITC), and anti-CD8 (APC) reveals an increase in CD8⁺ T-cells and total T-cells in biopsies from HIV-infected patients. Original magnification, $\times 400$. **C:** Quantification of CD25⁺FoxP3⁺ T-cells by immunofluorescence double labeling. Data are given as absolute numbers of positive cells $\pm$ SEM. **D:** Triple immunofluorescence labeling of syphilitic lesions from HIV-infected and HIV⁻ patients with anti-CD3 (FITC), anti-CD25 (TRITC), and anti-FoxP3 (Cy5) reveals decreased numbers of regulatory T-cells in HIV⁺ patients. **Arrows** denote triple positive cells. Original magnification, $\times 400$. **E:** Quantification of cell subsets was performed by single and double immunofluorescence staining with the indicated markers. Samples were subdivided according their lesional age. Data are given as the absolute numbers of positive cells $\pm$ SEM.

low in both HIV⁺ and HIV⁻ patients (Figure 7C). This observation suggests that an important antitreponemal effector mechanism is impaired in late lesions of HIV-infected patients.

Discussion

The immune response against *T. pallidum* was controversially discussed in the past. After early findings suggest-

Figure 5. Dendritic cells and mononuclear phagocytes within secondary syphilitic lesions. **A:** Detailed quantitative *in situ* analysis of dendritic cells and mononuclear phagocytes within the epidermis and dermis of secondary syphilitic lesions divided according to their HIV status and normal skin from healthy controls (HC). Double immunofluorescence staining was performed with the indicated markers. Data are given as the absolute numbers of positive cells $\pm$ SEM. **B:** Double immunofluorescence labeling of syphilitic lesions from HIV⁺ and HIV⁻ patients with anti-CD1a (FITC) and anti-Langerin (TRITC) elucidates a deficit in epidermal Langerhans cells of HIV⁺ patients. **C:** Triple immunofluorescence labeling of syphilitic lesions from HIV-infected and HIV⁻ patients with anti-CD11b (FITC), anti-CD14 (TRITC), and anti-HLA-DR (APC) reveals an increase of triple-positive mononuclear phagocytes in biopsies from HIV-infected patients. **B** and **C:** Original magnification, $\times 400$. **D:** Quantification of cell subsets with the indicated markers subdivided according to their lesional age. Data are given as the absolute numbers of positive cells $\pm$ SEM.

ing that syphilitic infection results in immunosuppression,¹⁶ studies in experimental animals gave evidence for a protective humoral immunity in early syphilis.^{17,18} It soon became clear by rabbit models that *T. pallidum* induces a strong cellular immune response⁴ dominated by T-cells and classical Th1 cytokines, consistent with a

delayed-type hypersensitivity reaction.⁷ In kinetic studies with rabbits infected with *T. pallidum*, T-cells were found to peak roughly at the same time as *T. pallidum* within the tissue. Lesional macrophages appear a few days later, which is paralleled by a sharp decline of lesional spirochetes.^{4,19,20} Similarly, T-cells and macrophages are

Figure 6. CD8⁺ T-cells as the main source of IFN-γ and IL-17. Triple immunofluorescence labeling of syphilitic lesions from HIV-infected and HIV⁻ patients with anti-CD3 (FITC), anti-CD8 (APC), and (A) anti-IFN-γ (TRITC), or (B) anti-IL-17 (TRITC) reveals that CD8-positive T-cells are IFN-γ- and IL-17-producing cells. C: Double labeling of IFN-γ and IL-17 showed an overlapping staining pattern. A–C: Arrows denote triple-positive cells. Original magnification, ×400. D: Data are given as the absolute numbers of positive cells ± SEM.

found in human primary chancres^{21,22} and secondary lesions.^{11,6} In functional studies phagocytosis of opsonized *T. pallidum* by macrophages,^{9,10} activated by IFN-γ,¹⁰ has been demonstrated, and degraded *T. pallidum*

organisms were found in phagocytic vacuoles of macrophages.^{23,5} These observations combined with the fact that treponemas are extracellular bacteria gave reason to speculate that macrophages, activated by IFN-γ-pro-

Figure 7. Increase of *T. pallidum* in late secondary syphilitic lesions. **A** and **B**: Immunohistochemistry of paraffin sections with an anti-*T. pallidum* antibody shows an increase of treponemas in HIV-infected patients. Original magnification, $\times 100$ of the total pictures and $\times 400$ of the insets. **B**: Quantification of treponemas in the epidermis and dermis of syphilitic patients by immunohistochemistry. **C**: Further splitting of the samples to early and late secondary syphilitic lesions reveals that the increase of treponemas in HIV⁺ patients accounts only for late lesions of HIV-infected individuals. **B** and **C**: Data are given as the absolute numbers of treponemas $\pm$ SEM. * $P < 0.05$. **D**: Quantification of *T. pallidum* transcripts by quantitative RT-PCR. Data represent the mean values of copies per ml eluate $\pm$ SEM. * $P < 0.05$.

ducing T-cells, are the pathogen-clearing cells in syphilis. The exact nature of the IFN- γ -producing T-cell subset has been a matter of debate. While CD4⁺ T-cells were the most likely candidates in rabbit models of syphilis infection,¹⁹ other studies indicated a possible role for CD8⁺ T-cells in providing this macrophage-activating cytokine.⁷

The cellular course of events described in animals and also observed in humans infected with *T. pallidum* is similar to what we observed in our HIV⁻ syphilitic patients. The earliest inflammatory cell was the lymphocyte, followed by neutrophils, macrophages, and plasma cells. Phenotypically, the T-cells belonged at equal parts to the CD4⁺ and CD8⁺ lineage. We now provide evidence that the antisiphilic immune response is not only mediated by IFN- γ -only-producing T-cells but also by IFN- γ /IL-17-double positive T-cells, a phenomenon not described before in syphilis. Attention has recently focused on a new effector cell subset, termed Th17/Tc17 cells, which supposedly play an important role in the protection against extracellular bacteria, in antitumor immunity as well as in the development of autoimmune and (auto)inflammatory diseases.^{24–29} In hepatitis C virus-infected patients a defined subset of CD8⁺ T-cells was observed to co-express IL-17 and high levels of IFN- γ ,³⁰ a phenotype consistent with what we found in syphilitic lesions especially from HIV⁺ individuals. Thus, our data fit well to and are an extension of the concept that *T. pallidum* elicits a delayed-type hypersensitivity reaction of mainly CD8⁺ T-cells producing IL-17 and IFN- γ , similar to what has been described for contact hypersensitivity reactions.^{26,27,29,31} One can speculate that these IFN- γ /IL-17-double positive CD8⁺ T-cells also play a role in the antisiphilic host response by IL-17- and IL-22-mediated antibacterial effector functions and IFN- γ -depen-

dent activation of macrophages that then clear opsonized treponemas. In addition, cytotoxic properties of CD8⁺ T-cells could be responsible for tissue destruction and inflammation that is particularly pronounced in syphilitic lesions of HIV⁺ patients.^{32–34} Activation of the newly discovered CD8⁺ T-cells by *T. pallidum* could take place directly via Toll-like receptor 2³⁵ or indirectly after intracellular uptake of the bacteria by macrophages and presentation via MHC-I.

Somewhat surprising was our finding of only few IL-17⁺ and/or IFN- γ ⁺ CD4⁺ T-cells in syphilitic lesions. One can argue that i) CD4⁺ T-cells are needed as “helper cells” for CD8⁺ T-cells, ii) a substantial proportion of CD4⁺ T-cells has a regulatory phenotype leading to the limitation of the immune reaction against treponemas, iii) IL-17 and IFN- γ are produced by CD4⁺ T-cells below the threshold of our detection system, or iv) CD4⁺ T-cells are “innocent” bystander cells in the anti-treponemal immune response. The last argument is favored by the clinical observation that the severity of clinical manifestations in HIV-infected patients does not strictly correlate with their CD4 cell count, as it is the case in most opportunistic infections.

The situation in HIV-infected syphilitic patients is more difficult to explain. Although studies exist which indicate that the clinical course of syphilis is not significantly distinct between HIV-infected and -uninfected individuals,^{36,37} our findings of increased numbers of *T. pallidum* in lesions of HIV⁺ patients would suggest that the antisiphilic immune response in these individuals is not very effective, and certainly not protective. It therefore appears that IFN- γ /IL-17-double-positive CD8⁺ T-cells and macrophages are perhaps needed but do not suffice to eliminate *T. pallidum*. This implies that an additional effector function is absent or at least down-regulated in

HIV-infected subjects. Different mechanisms come to mind. Exhaustion of CD8⁺ T-cells in HIV-infected individuals^{38,39} could result in an accumulation of these cells as compensation mechanism, incapable of eliminating the bacteria but causing increased tissue destruction, as shown for Th17/Tc17 cells in different chronic inflammatory diseases.^{40–42} On the other hand it is conceivable that in HIV-infected patients reduced numbers of dermal regulatory T-cells⁴³ and/or epidermal Langerhans cells⁴⁴ with regulatory properties⁴⁵ could cause an excessive, but not effective, cellular immune response. In this regard, it is noteworthy that IFN- γ -stimulated plasma cells express B7-H1 (PD-L1) and thereby inhibit T-cell functions.⁴⁶ Thus, increased levels of plasma cells in late syphilitic lesions combined with elevated IFN- γ levels could thereby create an immunosuppressive milieu resulting in diminished pathogen clearance.

The occurrence of IFN- γ /IL-17-producing CD8⁺ T-cells in syphilitic lesions should focus attention to this cell subset and could have implications for other infections with extracellular bacteria. Functional analysis should clarify their role in pathogen clearance and/or tissue destruction.

Acknowledgment

We thank Dr. Sheila Lukehart for helpful discussions and critical arguments.

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