Healthy HIV-1-Infected Individuals on Highly Active Antiretroviral Therapy Harbor HIV-1 in Their Alveolar Macrophages

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Abstract

In a prospective cross-sectional study we quantified HIV viral load within the alveolar macrophage in a cohort of healthy HIV-infected subjects who did not have medical comorbidities or smoke cigarettes to determine if alveolar macrophage proviral DNA was associated with alveolar macrophage phagocytic immune dysfunction. We enrolled 23 subjects who underwent bronchoscopy and bronchoalveolar lavage. Alveolar macrophages were isolated and HIV-1 RNA was quantified in the cells using the Abbott RealTime HIV-1 Assay. Proviral DNA was qualitatively measured using a modified version of the HIV-1 RNA assay. Phagocytosis measured by incubating alveolar macrophages with FITC-labeled Staphylococcus aureus and determining fluorescence with a Zeiss inverted microscope. Phagocytic index was calculated as (% positive cells×mean channel fluorescence)/100. Sixteen subjects had (+) proviral DNA and seven had (-) proviral DNA in their alveolar macrophages. Of all subjects 100% in both groups were on highly active antiretroviral therapy (HAART). The median plasma viral load was 0 in both groups. HIV-1-infected subjects with (+) proviral DNA in their alveolar macrophages had a significantly lower median alveolar macrophage phagocytic index compared to those with (-) proviral DNA in their alveolar macrophages [11.8 (IQR 4.8-39.0) vs. 64.9 (IQR 14.0-166.0), p=0.05]. Alveolar macrophages harbor HIV even in otherwise healthy subjects with undetectable plasma viral loads, representing a potential reservoir for the virus. In addition, HIV viral replication within the macrophage may impair phagocytosis and other immune functions in the lung, leading to an increased risk for lung infection.

Introduction

D ESPITE ADVANCES IN HUMAN immunodeficiency virus-1 (HIV-1) treatment including highly active antiretroviral therapy (HAART) individuals infected with HIV-1 continue to remain susceptible to serious bacterial and viral infections, such as pneumococcus and influenza, which continue to cause significant morbidity and mortality.^{1–3} A recent study in the Veterans Administration (VA) HIV-1 cohort found that bacterial pneumonia was the most common pulmonary disease, with an incidence of 28.0 per 1,000 person-years [95% confidence interval (CI), 27.2–28.8] compared with 5.8 per 1,000 person-years (95% CI, 5.6–6.0) among HIV-1-uninfected individuals (p = 0.001).⁴ Alveolar macrophage immune dysfunction could contribute to the increased risk of pneumonia in HIV-1-infected individuals, but the exact mechanisms are currently unknown. Alveolar macrophages are the primary innate immune defense cell in the lung and also the most plentiful cells in the lung. They are resistant to apoptosis, thus resulting in a half-life that is approximately 30 days.^{5,6} There is abundant evidence that HIV-1 infection impairs alveolar macrophage immune functions.^{7–9} Experimental studies have shown that macrophage phagocytic function by alveolar macrophages was significantly decreased in HIV-1 transgenic animals compared to wild-type animals.⁷

Eradication of HIV-1 has been difficult to achieve. Evidence for ongoing viral replication and the persistence of latently infected cells, including CD4⁺ T cells and cells from the monocyte/macrophage lineage, has been shown in HIV-1-infected individuals.^{10–12} HAART has been ineffective in eliminating these latent reservoirs. Furthermore, studies have shown that despite the lack of detectable replication-competent

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HIV-1 in patients on HAART, plasma HIV-1 reemerges when HAART is discontinued.¹³ Studies have also demonstrated the lack of correlation between viral rebound in plasma upon discontinuation of HAART and the size of the pool of latent HIV-1 reservoir prior to discontinuation of therapy, suggesting that the source of rebound HIV-1 viremia may not have come from a pool that is measurable in the circulating blood.^{14,15}

HIV-1 can infect long-lived alveolar macrophages and HIV-1 entry into these cells is preferentially mediated by the CCR5 receptor.¹⁶ However, reports comparing the burden of HIV-1 in alveolar macrophages relative to peripheral blood monocytes are conflicting and it is unclear what the exact viral burden is in the lung, as percentages of alveolar macrophages expressing HIV-1 antigens vary considerably.¹⁷ Phylogenetic evidence supports compartmentalization of HIV-1 between lungs and peripheral blood, suggesting that alveolar macrophages could be another reservoir for HIV-1.^{18,19}

We hypothesize that the alveolar macrophage is an important reservoir for HIV-1, contributing to cell–cell spread of the virus within the lung and exhibiting abnormal function, which contributes to the increase risk of severe lung infections. Therefore, we sought to quantify HIV-1 viral load within the alveolar macrophage and determine if alveolar macrophage proviral DNA was associated with alveolar macrophage phagocytic immune dysfunction.

Materials and Methods

Study design and protocol

All participants signed an Emory University Institutional Review Board approved consent to participate in the study. The research study purpose, procedures, risks and benefits, and alternative treatment options were described in detail and discussed with each subject. The subject was given an opportunity to read the informed consent form and ask questions. The subject also verbalized understanding of the study and all related visits and procedures, verbalized understanding of the HIPAA authorization, and signed and received a copy of the informed consent form. No study procedures were performed prior to obtaining informed consent. We performed a cross-sectional study of HIV-1-infected subjects without other medical problems within the Grady Health System in Atlanta, GA. Inclusion criteria included all subjects with HIV-1 infection. Exclusion criteria included active liver disease (known cirrhosis and/or total bilirubin > 2.0 mg/dl), heart disease [ejection fraction <50%, history of acute myocardial infarction, New York Heart Association (NYHA) II-IV cardiac symptoms, severe valvular dysfunction], current renal disease (dialysis dependent or creatinine > 2.0 mg/dl), current lung disease [spirometry revealing a forced vital capacity (FVC) or forced expiratory volume in 1 s (FEV₁) < 80% of predicted], diabetes, current pregnancy, malnutrition (body mass index < 17), current tobacco use, and age < 25 years.

The study coordinator entered the date, patient's name, medical record number, and study ID number in the patient enrollment log. Data were initially collected using our established case report forms and then entered into a HIPAAcompliant, secure database. All subjects completed a preenrollment evaluation (visit 1) that included (1) a complete history and physical examination, (2) routine blood chemistries (basic chemistry, liver function tests, complete blood count, coagulation parameters, hemoglobin A_{1C}), (3) urine pregnancy test (qualitative β -HCG) for women of childbearing potential, (4) urine dipstick for cotinine, (5) spirometry (FEV₁, FVC), (6) Short Michigan Alcohol Screening Test (SMAST) and Alcohol Use Disorders Identification Test (AUDIT) alcohol use questionnaires, and (7) body mass index (BMI). Demographic data were collected. Subjects with exclusions identified at the time of the preenrollment evaluation were excluded from further participation.

After completing the preenrollment evaluation to confirm eligibility, subjects presented to the Grady Memorial Hospital Clinical Interaction Site after an overnight fast. Enrolled volunteers had the following interventions on visit 2: (1) CD4⁺ count and viral load for HIV-1-infected subjects only and (2) flexible fiberoptic bronchoscopy with standardized bronchoalveolar lavage (BAL) performed using standard conscious sedation techniques. BAL was performed by installing 30-ml aliquots of 0.9% nonbacteriostatic normal saline solution, followed by withdrawal with low-pressure hand suction until a total of 180 ml had been administered. Subjects were contacted by phone 24 h after completing the study to ensure patient safety. A Data Safety Monitoring Representative was appointed to review each patient's enrollment procedures and bronchoscopy report on a monthly basis.

Alveolar macrophage phagocytosis

BAL fluid was centrifuged at 1,200 rpm for 7 min for recovery of cell pellet. Manual cell counts were performed with a hemocytometer and differentials were obtained from 300 consecutive cells after Diff-Quik staining (Andwin Scientific, Addison, IL). The cell pellet was resuspended in 10 ml of 1:1 Dulbecco's modified Eagle's medium/Ham's F-12 solution containing 2% fetal bovine serum (FBS), L-glutamine, 15 mmol/liter HEPES, penicillin (10,000 U), streptomycin (10,000 mg/ml), amphotericin (25 mg/ml), and gentamicin $(4 \,\mu g/ml)$. Alveolar macrophages (100,000 cells) were added to glass-chamber slides containing $100 \,\mu$ l of medium and $20\,\mu$ l of phosphate-buffered saline (PBS). Alveolar macrophages were incubated at 37°C with 10% CO₂ for 15 h, after which 10×10^{5} particles of pH-sensitive pHrodo fluorescein isothiocyanate-conjugated inactivated Staphylococcus aure*us* were added (10:1 ratio of *S. aureus*/alveolar macrophages) to the cultures and incubated for 2 h. Cells were washed, fixed with 4% paraformaldehyde, and stored at 4°C until analysis. Fluorescence was determined by quantitative computer analysis of images taken with a Zeiss inverted microscope using 10 fields per experimental condition with the same pinhole, detector gain, and amplifier offset. The basis of this assay is that S. aureus is internalized and the pHrodo component fluoresces once it reaches the lower pH (pH=4)present in the phagolysosome. Macrophages with any internalized bacteria were considered positive for phagocytosis. Phagocytosis was quantified by phagocytic index, which was calculated as the percentage of cells positive for phagocytosis multiplied by the mean relative fluorescence units of S. aureus per cell.

HIV-1 RNA and DNA tests

The Abbott RealTime HIV-1 Assay (Abbott Molecular Inc., Des Plaines, IL) is a real-time reverse transcription polymerase chain reaction (RT-PCR) assay for the quantitation of HIV-1

RNA on the automated m2000 System over the range of 40 to 10,000,000 copies/ml. The assay is standardized against a viral standard from the Virology Quality Assurance laboratory of the AIDS Clinical Trial Group and against the WHO 1st International standard for HIV-1 RNA. An internal control is introduced into specimens at the beginning of the sample preparation that is an RNA sequence unrelated to the HIV-1 target (pumpkin plant hydroxypyruvate reductase gene). An input of 1 million alveolar macrophages suspended in 100 μ l PBS was used in the HIV-1 RNA test, the elution volume is 50 μ l, all of which is used for amplification.

The Abbott RealTime HIV-1 Proviral DNA research use only assay is a qualitative modification of the above mentioned HIV-1 RNA viral load assay. The modifications allow amplification of HIV-1 proviral DNA only by removing the reverse transcription stage of the amplification protocol, modifying the nucleic acid extraction procedure from and RNA extraction to a DNA extraction, and adding a DNA internal control. The HIV-1 Proviral DNA test is qualitative and can detect an input of 153 copies of HIV-1 DNA. An input of 600,000 alveolar macrophages suspended in 100 μ l of PBS was used in the HIV-1 DNA test; the elution volume is 50 μ l, all of which is used for amplification.

Statistical analyses

Univariate comparisons between HIV-1 subjects with and without proviral DNA were calculated and evaluated for a significance level of 0.05 using a chi-squared test for categorical variables and a two-sample *t*-test for continuous variables. The data were log-transformed or a Wilcoxon rank-sum test was used when the data were not normally distributed.

Results

Baseline characteristics of HIV-1-infected subjects

A total of 23 HIV-1 subjects were enrolled. Table 1 shows the demographic characteristics of HIV-1-infected subjects enrolled in the study divided into two groups—those who had HIV-1 proviral DNA present in their alveolar macrophages compared to those who did not. We had 16 HIV-1-infected subjects with (+) proviral DNA in their alveolar macrophages and 7 HIV-1-infected subjects with (-) proviral DNA in their alveolar macrophages. There were no significant differences in median age, gender, or race. The majority of subjects in both groups were African American. Of the subjects, 100% in both groups were on HAART. HAART regimens varied and are listed in Table 2. HIV-1-infected subjects with (+) proviral DNA trended toward a lower median CD4 count compared to HIV-1-infected subjects with (-) proviral DNA [349.5 (IQR 248.5–452.3) vs. 560 (390–730), p=0.06]. Approximately 71.4% of subjects with (+) proviral DNA had an undetectable plasma viral load vs. 100% of subjects with (-) proviral DNA in their alveolar macrophages. However, the median plasma viral load was 0 in both groups.

HIV-1 Proviral DNA and RNA are detectable and quantifiable in alveolar macrophages

Sixteen HIV-1-infected subjects had (+) proviral DNA present in their alveolar macrophages (Fig. 1). Eight of these subjects had (+) proviral DNA and (+) HIV-1 RNA. Of these eight subjects, four had quantifiable HIV-1 RNA in their alveolar macrophages. Table 3 details the four subjects who, in addition to detectable proviral DNA, had quantifiable HIV-1 RNA in their alveolar macrophages. Subject 1 had the highest HIV-1 RNA count in the plasma and the second highest count within the alveolar macrophage. However, Subjects 2–4 had quantifiable HIV-1 RNA within their alveolar macrophages, whereas their plasma HIV-1 RNA levels were undetectable.

Proviral DNA within alveolar macrophages is associated with impaired bacterial phagocytosis

Twelve out of 16 HIV-1-infected subjects with (+) proviral DNA and six out of seven HIV-1-infected subjects with (-) proviral DNA had alveolar macrophage phagocytosis data. HIV-1-infected subjects with (+) proviral DNA in their alveolar macrophages had a significantly lower median alveolar macrophage phagocytic index compared to those with (-) proviral DNA in their alveolar macrophages [11.8 (IQR 4.8–39.0 vs. 64.9 (IQR 14.0–166.0), p=0.05] (Fig. 2).

Discussion

The results from this study showed that HIV-1 was both detectable and quantifiable within alveolar macrophages even in otherwise healthy, nonsmoking, HIV-1-infected individuals with plasma viral suppression. HIV-1 DNA was present in approximately 70% of this "healthy" cohort, while

Variables	(+) Proviral DNA	(-) Proviral DNA	р
N	16	7	
Median age (IQR)	49.5 (44–54.3)	50 (47-53)	0.84
Gender (% male)	43.8	57.1	0.55
Race			0.53
White $(n, \%)$	1 (6.3)	1 (14.3)	
Black $(n, \%)$	15 (93.7)	6 (85.7)	
On HAART $(n, \%)$	16 (100)	7 (100)	
Median CD4 (IOR)	349.5 (248.5-452.3)	560 (390-730)	0.06
% Undetectable plasma viral load	71.4%	100%	0.12
Median plasma viral load copies/ml (IQR)	0 (0–27.3)	0 (0–0)	0.18

HAART, highly active antiretroviral therapy.

HIV-1 IN ALVEOLAR MACROPHAGES

Proviral DNA detected in alveolar

Subject macrophage

TABLE 2. HIGHLY	ACTIVE A	Antiretrov	IRAL	THERAPY
REGIMENS OF	THE HIV-	-1-INFECTED	SUB.	JECTS
Enre	OLLED IN	THE STUDY	7	

TABLE 3. SUBJECTS WITH QUANTIFIABLE HIV-1 RNA in Their Alveolar Macrophages

Subjects	Alveolar macrophage HIV RNA (copies/million cells)	Plasma HIV RNA (copies/ml)
1	182	27,542
2	588	Undetectable
3	70	Undetectable
4	48	Undetectable

ID	(+/-)	HAART regimen
1	+	Emtricitabine/tenofovir/efavirenz
2	+	Raltegravir + emtricitabine/tenofovir
3	_	Emtricitabine/tenofovir/efavirenz
4	_	Raltegravir + emtricitabine/
		tenofovir + lopinavir/ritonavir
5	+	Raltegravir + lamivudine/zidovudine
6	+	Emtricitabine/tenofovir/efavirenz
7	_	Emtricitabine/tenofovir/efavirenz
8	+	Efavirenz + abacavir
9	+	Emtricitabine/tenofovir+
		darunavir + ritonavir
10	+	Emtricitabine/tenofovir/efavirenz
11	+	Ritonavir
12	+	Lopinavir/ritonavir + abacavir
13	+	Atazanavir + ritonavir + abacavir/
		lamivudine
14	+	Raltegravir + tenofovir/emtricitabine
15	-	Tenofovir/emtricitabine
16	+	Emtricitabine/tenofovir/efavirenz
17	+	Emtricitabine/tenofovir/efavirenz
18	_	Emtricitabine/tenofovir/efavirenz
19	+	Raltegravir + emtricitabine/tenofovir
20	-	Raltegravir + tenofovir/
		emtricitabine + etravirine
21	_	Emtricitabine/tenofovir/efavirenz
22	+	Emtricitabine/tenofovir/efavirenz
23	+	Atazanavir + ritonavir

both HIV-1 DNA and RNA were detectable in 34.8% and HIV-1 RNA was quantifiable in about 17.4% of all subjects. In addition, the presence of proviral DNA within alveolar macrophages was associated with impaired alveolar macrophage bacterial phagocytic capacity, even in this population with no history of lung infections or lung disease. The alveolar macrophage may, therefore, represent a potential reservoir for the virus. This is the first study demonstrating that (1) HIV-1 DNA and RNA are present within alveolar macrophages in otherwise healthy HIV-1-infected subjects with systemic viral suppression and (2) that the presence of HIV-1 DNA within the alveolar macrophages was associated with impaired lung immune function, as reflected by impairment in alveolar macrophage phagocytic capacity.

This study adds to the accumulating evidence supporting the hypothesis that the lung may be a reservoir for HIV-1. Certain studies have shown a greater HIV-1 burden in unfractionated BAL fluid and undifferentiated BAL cells compared to the periphery.^{20–22} In one of the largest studies, Clarke *et al.* compared HIV-1 DNA copy number within lung cells from BAL specimens to peripheral blood cells from 78 HIV-1-infected individuals not on HAART.²⁰ They determined that the mean HIV-1 DNA copy number was greater in BAL inflammatory cells compared to peripheral blood in



FIG. 1. HIV proviral DNA and RNA are detectable and quantifiable in alveolar macrophages. Alveolar macrophages were isolated from otherwise healthy HIV-infected subjects. The Abbott RealTime HIV-1 Assay was used to measure both HIV DNA and RNA.



FIG. 2. Proviral DNA within alveolar macrophages is associated with decreased bacterial phagocytosis. Alveolar macrophages were isolated from otherwise healthy HIV-infected subjects and incubated with FITC-labeled *S. aureus*. Phagocytosis was determined by measuring fluorescence with a Zeiss inverted microscope. The phagocytic index was calculated as (% positive cells×mean channel fluorescence)/ 100. The Abbott RealTime HIV-1 Assay was used to measure HIV proviral DNA.

approximately 56% of subjects. Furthermore, in both BAL and peripheral blood, a higher level of HIV-1 DNA was detected in the monocyte/macrophage lineage compared to leukocytes. In a subgroup of individuals receiving azidothymidine (AZT) therapy, the authors found a reduced mean HIV-1 proviral DNA copy number in each subpopulation of cells.²⁰ However, in contrast to this current study, all of these individuals were undergoing bronchoscopy for pulmonary symptoms and approximately 38% of the HIV-1-infected individuals proceeded to AIDS with a diagnosis of Pneumocystis carinii (now known as Pneumocystis jiroveci) pneumonia (PCP). Although this evidence suggests that the lung is a significant focus of HIV-1 infection and may be important in the pathogenesis of disease, there is significant heterogeneity in the current literature including the presence of lung infections, other lung diseases, and tobacco use in the different study populations.

We also found that HIV-1 replication in alveolar macrophages was compartmentalized compared to the serum in eight subjects in whom HIV-1 RNA and DNA were detected. In these subjects, three had undetectable plasma viral loads. Furthermore, three out of four subjects with quantifiable HIV-1 RNA had undetectable viral loads in the periphery. These data suggest that HIV-1 replication in the lungs occurs separately from the blood and is not just from simple diffusion from the periphery. Koziel et al. examined the relationship of viral burden in cell-free BAL fluid to that of serum in asymptomatic HIV-1-infected individuals and found that 16% of asymptomatic subjects had evidence of HIV-1 RNA in cell-free BAL fluid.²¹ However, in contrast to this current study, all seven asymptomatic subjects with HIV-1 RNA in BAL fluid had detectable HIV-1 RNA in their serum and six of these seven subjects were not taking antiretroviral medications. Furthermore, unlike this current study, all seven were active tobacco users.

In addition, in this study of otherwise healthy HIV-1infected subjects, we found that the presence of HIV-1 DNA within alveolar macrophages significantly impaired alveolar macrophage phagocytic capacity. Experimental data show us that HIV-1 infection can affect alveolar macrophage immune functions,^{7–9} including the finding that abnormal activation impaired oxidative burst and cytokine secretion.^{23–25} In humans, similar macrophage dysfunctions are seen^{26,27}; however, the data with respect to phagocytic functions are conflicting.^{28–31} Gordon et al. found that alveolar macrophages from healthy, nonsmoking HIV-1-infected adults showed normal internalization of opsonized pneumococci,²⁹ whereas others have shown that HIV-1-infected adults have impaired macrophage phagocytic capacity.^{30,32,33} Most recently, Jambo et al. found that HIV-1 present in small alveolar macrophages from asymptomatic HIV-1-infected HAART-naive individuals was associated with impaired phagocytic capacity and proteolytic activity.³² However, prior to the current study, the presence of intracellular HIV-1 DNA or RNA has not been associated with impaired alveolar macrophage phagocytic function.

The effect of HAART on lung viral load is still being studied. In a prospective, longitudinal study, Twigg *et al.* determined that HAART was associated with a significant decline in plasma viral load, but the decline in lung viral load, both in the acellular and cellular compartments, was more rapid even after just 4 weeks of HAART, which was sustained for 6 months.²² These results suggest that the lung

seemed to respond to HAART just as well as the vascular compartment. These subjects all had CD4 counts \geq 500 cells/ μ l and a plasma HIV-1 RNA load of \geq 5,000 copies/ml. The finding of 34.8% of our asymptomatic HIV-1-infected adults with HIV-1 RNA within their alveolar macrophages, despite HAART, suggests that HIV-1 may be compartmentalized to an area in the lung where HAART is not as effective. At this time, it is still unclear what the pharmacokinetics of HAART in the lung are and whether HAART can restore lung immune function. Currently, CD4 counts and viral loads are used as prognostic markers in this population. However, our study revealed that even otherwise healthy HIV-1-infected individuals have persistent HIV-1 within alveolar macrophages that is associated with macrophage immune dysfunction, despite systemic viral suppression and higher CD4 counts. If this is corroborated in larger clinical trials, then clinicians will need new diagnostic and prognostic biomarkers of susceptibility to lung infection in this vulnerable population.

There are a number of limitations of this study. This is an observational cross-sectional study and so is subject to bias when comparing the two groups. However, confounding was minimized by restricting the population to a select group of individuals who were otherwise healthy, nonsmokers, and without any major medical problems. HAART regimens were self-reported by the study subjects. Although we confirmed regimens from subjects who were followed in the clinic, there were a number of subjects who received their care at other hospitals and we were not able to confirm their medications. In addition, we are currently unable to confirm the duration of HAART. Furthermore, although HIV-1 RNA and DNA were measured in the alveolar macrophages, we did not determine whether the HIV-1 within these alveolar macrophages was replication competent. Although we are unable to assess this on the current samples, we intend to address this important question in the future.

In conclusion, HIV-1 viral infection of alveolar macrophages may impair macrophage phagocytosis and other immune functions, leading to an increased risk for lung infections. At this time, there are a number of unanswered questions that are critical to understanding the effects of HIV-1 infection in the lung. It is unknown what the clinical impact of persistent viral replication in the lung is: Does it, in fact, lead to an increase risk of pulmonary infections? Is impairment of macrophage function implicated in the pathogenesis of other lung diseases, such as obstructive lung disease? Furthermore, if even otherwise healthy HIV-1infected individuals on HAART with systemic viral suppression have HIV-1 present in the lung, and this presence is associated with impaired immune function, then clinicians need better diagnostics to determine which individuals are at risk for lung infections. Finally, supplemental therapies to HAART may be necessary to target the alveolar macrophage reservoir and improve lung immune function.

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Author Disclosure Statement

No competing financial interests exist.

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