Cerebrospinal Fluid HIV-1 Compartmentalization in a Patient With AIDS and Acute Varicella-Zoster Virus Meningomyeloradiculitis

E. Liana Falcone,¹ Ademiposi A. Adegbulugbe,³ Virginia Sheikh,² Hiromi Imamichi,² Robin L. Dewar,⁴ Dima A. Hammoud,⁵ Irini Sereti,² and H. Clifford Lane²

¹ Laboratory of Clinical Infectious Diseases and ² Clinical and Molecular Retrovirology Section, Laboratory of Immunoregulation, National Institute of Allergy and Infectious Diseases, National Institutes of Health, Bethesda, Maryland; ³Section of Infectious Diseases, Department of Medicine, Washington Hospital Center, Washington, District of Columbia; ⁴Clinical Services Program, SAIC-Frederick, NCI-Frederick, and ⁵Center for Infectious Disease Imaging, Radiology and Imaging Sciences, National Institutes of Health, Bethesda, Maryland

We report a case of AIDS presenting as varicella-zoster virus (VZV) meningomyeloradiculitis associated with human immunodeficiency virus (HIV) quasispecies compartmentalization within the cerebrospinal fluid (CSF), and a CSF viral load that was 1 log higher than in peripheral blood. Prolonged antiviral therapy for both VZV and HIV type 1 was associated with partial resolution.

Keywords. human immunodeficiency virus; viral quasispecies; central nervous system compartmentalization; varicellazoster virus; meningomyeloradiculitis.

Viral diversity is an important feature of human immunodeficiency virus (HIV) pathogenesis. Within an individual, HIV exists as a population of related yet distinct viral variants termed viral quasispecies [\[1\]](#page-6-0). These quasispecies evolve as an adaptation to selection pressures such as antiviral therapy and/ or cellular microenvironment, and can lead to changes in key virologic properties including cellular tropism and coreceptor utilization [\[2\]](#page-6-0). Viral quasispecies are not evenly distributed throughout the body, and analysis of their diversity suggests

Received 1 February 2013; accepted 16 May 2013; electronically published 31 May 2013. Correspondence: E. Liana Falcone, MD, MSc, Laboratory of Clinical Infectious Diseases, Na-

tional Institute of Allergy and Infectious Diseases, National Institutes of Health, 10 Center Drive, CRC, Rm B3-4141, MSC 1684, Bethesda, MD, 20892-1684 [\(emilia.falcone@nih.gov](mailto:emilia.falcone@nih.gov)). Clinical Infectious Diseases 2013;57(5):e135–42

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that there may be cell type and/or tissue-specific viral variants that may exist within an infected individual [[3](#page-6-0)]. The term compartmentalization has been attributed to the restriction of viral quasispecies between cells or tissues [[4](#page-6-0)]. One compartment of great interest is the central nervous system (CNS), where there have been reports of selective replication of distinct viral quasispecies leading to resistance confined to the CNS [[5](#page-6-0)], as well as an association between increased cerebrospinal fluid (CSF) viral load and HIV dementia [[6](#page-6-0)].

It is not yet known whether quasispecies compartmentalization in the brain and CSF is associated with other CNS infections in patients with HIV. This patient population does, however, have an increased risk of developing reactivation of varicella-zoster virus (VZV) infection that can lead to serious neurological complications due to subclinical extension of VZV into the CNS. These CNS complications include postherpetic neuralgia, ophthalmic zoster with contralateral hemiparesis, acute or chronic encephalitis, myelitis, polyradiculitis, motor neuropathies, and cranial nerve palsies [\[7,](#page-6-0) [8](#page-6-0)], as well as immune reconstitution inflammatory syndrome (IRIS) [[9\]](#page-6-0).

We report a case of AIDS presenting as VZV meningomyeloradiculitis, which was associated with an HIV type 1 (HIV-1) load in the CSF that was approximately 1 log higher than that in peripheral blood, and with HIV viral quasispecies compartmentalization within the CSF. The patient's neurological syndrome was progressive. A prolonged course of antiviral therapy for both VZV and HIV-1 was associated with stabilization and partial resolution.

CASE REPORT

A 50-year-old African American man presented with a 4-day history of left-sided chest pain and progressive left leg weakness resulting in multiple falls. Two days prior to admission, he noted a painful vesicular rash on the left side of his chest, but denied having any fever, headache, visual disturbances, seizures, neck stiffness, or incontinence. He had a 15-pack-year smoking history, and reported remote intranasal cocaine use, as well as having unprotected intercourse with multiple female partners. His physical exam revealed oropharyngeal thrush, a vesicular rash in the left T4 dermatome, and 0/5 strength in the left lower extremity in association with an increased left patellar reflex and an extensor plantar response on the left. Temperature and pinprick sensation were decreased in the right leg, and torso below T5.

Magnetic resonance imaging (MRI) of the spine showed a focal, centrally located intramedullary hyperintensity on T2 weighted images, at the T3 level that was associated with slight cord expansion and subtle enhancement. There was also marked diffuse meningeal thickening and enhancement involving the cervical and thoracic cord as well as the cauda equina nerve roots (Figure [1\)](#page-2-0). Laboratory testing was significant for a positive HIV 1/2 rapid antibody screen, a CD4⁺ T-cell count of 35 cells/μL, and an HIV-1 load of 14 163 copies/mL. Plasma HIV-1 viral sequence analysis for reverse transcriptase (RT) and protease resistance mutations was negative. Additional blood tests included negative rapid plasma reagin, cytomegalovirus polymerase chain reaction (PCR), Toxoplasma serologies, and serum antigen for Cryptococcus. A blood PCR for Epstein-Barr virus was positive at 350 copies/mL. The CSF contained 132 white blood cells (WBCs)/mm³ (27% neutrophils and 60% lymphocytes), 130 red blood cells/mm³, 486 mg/dL of protein, and 63 mg/dL of glucose. Cytomegalovirus, herpes simplex virus, and John Cunningham virus PCRs were all negative in the CSF, and the Venereal Disease Research Laboratory test was nonreactive. VZV PCR was positive in the CSF. The CSF HIV-1 load was found to be 108 982 copies/mL.

Given the significant discrepancy in HIV-1 load between the plasma and CSF, phylogenetic analysis of the HIV-1 envelope (env) gene using cell-free virion RNA as a template from plasma and CSF samples was performed as previously described [[10\]](#page-6-0), except that the following 2 sets of primers were used to amplify full-length envelope genes: +5954 (sense) 5′- GGCTTAGGCATCTCCTATGGCAGGAAGAA-3′ and –9146 (antisense) 5′-CTGCCAATCAGGGAAGTAGCCTTGTGT-3′ in a first-round reaction; +6202 (sense) 5′-AGAAAGAGCAGA AGACAGTGGCAATGA-3′ and –9068 (antisense) 5′-TAGCC CTTCCAGTCCCCCCTTTTCTTTTA-3′ in a second-round reaction. HIV-1 sequences from the patient clustered into 3 phylogenetically distinct groups: one group consisting of plasma virus sequences, a second group containing plasma and CSF virus sequences, and a third group consisting exclusively of CSF virus sequences. The compartmentalization of HIV-1 between the peripheral blood and spinal cord is represented by separate clusters for plasma, and CSF sequences in the phylogenetic tree (Figure [2\)](#page-3-0). Moreover, the major separations observed in the phylogenetic tree were associated with predicted CXC chemokine receptor type 4 (CXCR4) versus CC chemokine receptor type 5 (CCR5) usage of the virus [[11\]](#page-6-0). Based on the Web PSSM analysis (a bioinformatics tool for predicting HIV-1 coreceptor usage), the viral sequences clustering in the CSF showed CXCR4 use, whereas the viral sequences predominating in the plasma predicted CCR5 use.

Soon after admission, the patient was started on antiretroviral therapy (emtricitabine, tenofovir, efavirenz), intravenous acyclovir, and methylprednisolone. Following 7 days of treatment, he experienced progression of his neurologic impairment with an increase in CSF pleocytosis (132–306 WBCs/ mm³). VZV DNA, however, was no longer detected in his CSF by PCR. Thus, after 14 days of acyclovir, he was switched to famciclovir. Nevertheless, follow-up MRI showed several new T2-hyperintense lesions in the thoracic cord with persistent abnormal cord meningeal and cauda equina nerve root enhancement. The patient was therefore restarted on intravenous acyclovir for an additional 4 weeks. He was then switched to valacyclovir for 8 weeks followed by an additional 8 weeks of valacyclovir at prophylactic doses. His follow-up MRI at the end of his treatment course showed significant improvement with almost complete resolution of the meningeal enhancement, as well as decreased size and conspicuity of the cord lesions. There was, however, persistent evidence of cauda equina nerve root clumping and enhancement suggestive of irreversible arachnoiditis (Figure [1](#page-2-0)). Follow-up plasma HIV loads eventually became <50 copies/mL, and CSF HIV loads declined from 108 982 copies/mL to 211 copies/mL (Table [1](#page-3-0)). The patient is currently much improved neurologically and is ambulating with a walker.

DISCUSSION

To our knowledge, this is the first report of a case of HIV quasispecies compartmentalization in the CNS, in a patient presenting with VZV meningomyeloradiculitis and incomplete Browne-Séquard syndrome. Our patient improved significantly with a prolonged course of acyclovir followed by valacyclovir, and prompt initiation of antiretroviral therapy.

Herpes zoster and its related neurological complications are more common in the HIV-infected population, especially when $CD4^+$ T-cell counts are <100 cells/ μ L [[7,](#page-6-0) [12](#page-6-0)]. Such complications are often diagnosed based on the detection of VZV DNA in the CSF and a temporal association with the appearance of skin lesions. However, zoster reactivation can occur in the absence of any skin manifestations [[7](#page-6-0), [13](#page-6-0)–[15](#page-6-0)]. We reviewed reported cases of VZV myelitis or myeloradiculitis in patients with HIV/AIDS (Table [2](#page-4-0)). Seven of 14 reported patients died, most of whom were from before the era of combined antiretroviral therapy. In at least 9 out of the 14 cases, patients were treated with antivirals (mainly intravenous acyclovir) with or without a short course of steroids. CSF HIV loads were not reported in any of the cases.

HIV infects the CNS during acute primary infection, and compartmentalizes within this site by infecting macrophages and microglia [\[16](#page-6-0)]. The CNS is an important reservoir for long-term viral persistence (and potential neurocognitive impairment), partly due to the relatively poor CNS penetration of many antiretrovirals, as well as the fact that drug resistance can evolve independently within the CNS [[17\]](#page-6-0). In fact, studies have shown that there can be CSF HIV escape with reported CSF

Figure 1. Magnetic resonance imaging findings at presentation and after treatment. Sagittal T2-weighted and sagittal enhanced T1-weighted images of the cervical, thoracic, and lumbosacral spine upon presentation (A, C, A) and E respectively) and after treatment (B, D, A) and F respectively). Thickening and enhancement of the meningeal covers (hollow arrows in A and C) is noted upon presentation. This improves significantly on the follow-up examination (hollow arrows in B and D). Parenchymal cord involvement with abnormal T2-hyperintense lesion at T3 level noted upon presentation also improves significantly on follow-up (solid arrows in A and B). Enhancement of the cauda equina nerve roots (hollow arrows in E) is later replaced by nerve root clumping and enhancement (hollow arrows in F) suggesting the development of chronic arachnoiditis. Note the development of adhesions along the ventral aspect of the cord at T3–T4 level (white solid arrows in D).

Figure 2. Phylogenetic analysis of human immunodeficiency virus type 1 (HIV-1) env sequences derived from cerebrospinal fluid (CSF) and plasma samples. Phylogenetic relationships among the full-length envelope gene sequences were estimated using the neighbor-joining method. Bootstrap percentile values from 1000 replications are shown at nodes defining major grouping of sequences. Statistical support of 50% or greater is shown. HIV-1 U455 was used as an outgroup. Cluster 1 contains plasma sequences, cluster 2 contains plasma and CSF sequences, and cluster 3 contains CSF sequences. The majority of CSF sequences predict CXCR4 usage, while all plasma sequences predict CCR5 usage.

viral loads as high as 12 885 copies/mL while plasma viral loads remain suppressed due to combined antiretroviral therapy [[18,](#page-6-0) [19](#page-7-0)]. In our patient, however, the discrepancy between the CSF and plasma viral loads was present prior to beginning combined antiretroviral therapy, during a severe episode of zoster reactivation. These findings suggest that CNS coinfection with VZV may have played a role in the enhanced HIV replication in the CSF.

The primary determinant of HIV macrophage–mediated neurotropism is genetic variation of the viral env gene. In our patient, phylogenetic analysis of env sequences demonstrated a first viral cluster containing plasma sequences, a second viral cluster containing both plasma and CSF sequences, and a third viral cluster consisting of CSF sequences. Interestingly, the clustering pattern was associated with predicted coreceptor usage: the first and second clusters with CCR5 usage, and the third

Table 2. Reported Cases of Varicella-Zoster Virus Myelitis or Myeloradiculitis in Patients With HIV

Abbreviations: ARV, antiretrovirals; CSF, cerebrospinal fluid; IV, intravenous; MRI, magnetic resonance imaging; NR, not reported.

cluster with CXCR4 usage. However, consistent with previous reports [13], the predicted coreceptor usage did not distinguish viruses in CSF from those in plasma.

Tissue-specific compartmentalization of HIV-1 variants in the brain has been demonstrated in a number of studies [13, [20](#page-7-0)–[22\]](#page-7-0). The CSF is a compartment that bridges the brain and the periphery. Thus, it is plausible that HIV-1 variants in the CSF can be genetically similar to those in the brain and the periphery. In fact, there has been a report demonstrating a close genetic relationship between HIV-1 variants in CSF and plasma [[20](#page-7-0)]. This intermingling of CSF and plasma HIV-1 sequences was also observed in our patient. These findings suggest that the CSF compartment was infiltrated by HIV-1 from the periphery (represented by the second cluster) and subsequently underwent local amplification of the HIV-1 variants (resulting in the formation of the third cluster), possibly driven by the selective pressure that was unique to the CSF compartment. From a clinical standpoint, viral load discrepancies between the plasma and CSF should, at the very least, prompt sequencing of the RT, protease, and integrase portions of the genome to assist in clinical decision making, and the selection of the appropriate antiretroviral regimen.

Our patient presented with a severe neurological complication of zoster reactivation in the setting of HIV load discrepancy between the plasma and CSF with quasispecies compartmentalization in the CNS. He required a prolonged course of intravenous acyclovir and was started on antiretrovirals early in his course in order to achieve a significant resolution of his neurological symptoms. It is possible that the initial worsening of the patient's neurological status may have been due to IRIS, especially in the context of initiation of ART 1 week prior. However, the patient was already on IRIS treatment doses of steroids at the time that ART was initiated, and they did not appear to be of benefit in this setting. Moreover, the CSF VZV PCR had become negative, and the CSF HIV load was increased relative to the plasma viral load.

It is unclear whether localized HIV replication in the CNS played a role in the severity of the zoster reactivation in this patient or whether concomitant VZV infection facilitated CNS HIV replication. Future phylogenetic analyses of HIV sequences in the CSF during zoster reactivation or other acute CNS infections may be of interest in better understanding the potential synergy between these 2 viruses.

Notes

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