

Plasma Cytokine Levels are Related to Brain Volumes in HIV-infected Individuals

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Abstract HIV-infected individuals frequently exhibit brain dysfunction despite antiretroviral treatment. The neuropathological mechanisms underlying these abnormalities remain unclear, pointing to the importance of identifying biomarkers sensitive to brain dysfunction. We examined 74 medically stable HIV-infected individuals using T1-weighted MRI. Volumes of the cortical grey matter (GM), white matter (WM), caudate, putamen, globus pallidus, thalamus, hippocampus, amygdala, and ventricles were derived using automated parcellation. A panel of plasma cytokines was measured using multiplexed bead array immunoassay. A model selection algorithm was used to select the combination of clinical and cytokine markers that best predicted each brain volumetric measure in a series of linear regression models. Higher CD4 nadir, shorter HIV infection duration, and antiretroviral treatment were significantly related to higher volumes of the putamen, thalamus, hippocampus, and WM. Older age

was related to lower volumes in most brain regions and higher ventricular volume. Higher IFN- γ , MCP-1, and TNF- α were related to higher volumes of the putamen, pallidum, amygdala, GM, and WM. Higher IL-1 β , IL-6, IL-16, IL-18, IP-10, MIP-1 β , and SDF-1 α were related to lower volumes of the putamen, pallidum, thalamus, hippocampus, amygdala, GM, and WM; and higher ventricular volume. The current findings provide evidence linking smaller brain volumes to HIV disease history, antiretroviral treatment, and advanced age. Cytokine markers, especially IL-6 and IL-16, showed robust association with brain volumes even after accounting for other clinical variables, demonstrating their utility in examining the mechanisms of HIV-associated brain abnormalities.

Keywords HIV infection · Hepatitis C infection · Antiretroviral treatment · MRI

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Introduction

People living with HIV continue to exhibit brain dysfunction despite effective antiretroviral treatment. Although the prevalence of advanced dementia has dramatically declined since the advent of combination antiretroviral therapy (CART), the overall rates of neurocognitive dysfunction in people living with HIV remain strikingly similar to those observed pre-CART (Tozzi et al. 2007; Heaton et al. 2010). Evidence from neuroimaging studies has corroborated this impression, demonstrating prevalent structural, metabolic, and cerebrovascular abnormalities even in medically stable HIV-infected individuals (Cardenas et al. 2009; Harezlak et al. 2011; Jernigan et al. 2011; Gongvatana et al. 2009; Roc et al. 2007; Ances et al. 2009; Becker et al. 2011; Lepore et al. 2008).

The etiology of this persistence of HIV-associated brain dysfunction in the CART era remains unclear. Recent findings have linked neurocognitive function and structural and metabolic MRI markers to nadir CD4 levels (Jernigan et al. 2011; Cohen et al. 2010; Ellis et al. 2011), suggesting the importance of distant history of immune suppression on current brain integrity. Hepatitis C (HCV) coinfection has recently emerged as an important contributor to both neurocognitive and structural brain abnormalities observed in HIV-infected individuals (Jernigan et al. 2011; Gongvatana et al. 2011; Cherner et al. 2005; Clifford et al. 2005; Hinkin et al. 2008; Devlin et al. 2012). In addition, HIV-associated neurocognitive impairment has been linked to chronic systemic immune activation and inflammation (Cohen et al. 2011; Burdo et al. 2013; Ryan et al. 2001; Sevigny et al. 2004), pointing to the potential utility of peripheral cytokine markers both for probing the underlying neuropathological mechanisms and as biomarkers sensitive to brain dysfunction.

The current study examines the independent contributions and relative importance of these diverse clinical markers on brain volume alterations. Volumes of the cerebral cortex, cerebral white matter, subcortical structures, and ventricles were measured from automated parcellation of high-resolution T1-weighted MRI images. Volumetric measures were examined in the context of markers of HIV disease, antiretroviral treatment, HCV coinfection, and a panel of thirteen plasma cytokine markers measured using a multiplexed bead array immunoassay. We utilized multivariate statistical models to examine the independent contributions of clinical and cytokine markers on brain volumetric measures, and a model selection algorithm was used to identify the subset of markers that best predict each brain volume. We hypothesized that advanced HIV disease, lack of antiretroviral treatment, HCV coinfection, older age, and abnormal cytokine markers would

be related to lower volumes of brain structures and greater ventricular volume.

Methods

Participants

Participants include 74 HIV-infected (HIV+) individuals recruited as part of an NIH-sponsored study of HIV-associated brain dysfunction at The Miriam Hospital / Brown University. The study was approved by the institutional review board, and informed consent was obtained from each participant prior to enrollment. Exclusion criteria included 1) history of head injury with loss of consciousness > 10 min; 2) neurological conditions including dementia unrelated to HIV, seizure disorder, stroke, and opportunistic infection of the brain; 3) severe psychiatric illness that may impact brain function, e.g., schizophrenia; and 4) diagnosis of alcohol or substance abuse or dependence based on DSM-IV criteria within 6 months prior to neuroimaging. HIV serostatus was documented by ELISA and confirmed by Western blot test.

Participants were between 24 and 65 years of age (mean = 45, SD = 9.66). Average duration of HIV infection was about 13 years. Most participants (82 %) received combination antiretroviral therapy (CART), and generally had well-controlled HIV viral load and intact immune function: 72 % had undetectable plasma HIV RNA (<75 copies/ml), average CD4 count was 523. More than half (54 %) of the participants reported nadir CD4 < 200, indicating history of significant immune suppression. A significant number (34 %) of participants had active HCV infection documented by detectable serum HCV RNA by PCR. Table 1 shows relevant demographic and clinical information of participants.

Cytokine Measurements

Blood samples acquired for each participant were separated, and plasma samples were immediately frozen and stored at -80°C . Aliquots were used to measure chemokine and cytokine levels using an xMAP multiplexed bead array immunoassay. This approach permits simultaneous quantification of multiple cytokines in solution by capturing them onto antibody coated spectrally distinct fluorescent microspheres, and measuring fluorescence intensity using the Luminex-100 system (Luminex Corp., Austin, TX). Levels of IFN- γ , IL-1 β , IL-6, IL-8, IL-10, IL-16, IL-18, IP-10, MCP-1, MIP-1 β , SDF-1 α , TNF- α , and TRAIL were measured. These specific markers were selected for their sensitivity to

Table 1 Relevant demographic and clinical information of study participants

N	74
Age (years)	45.28±9.66
Gender (% male)	66.22 %
Ethnicity (% Caucasian)	50.00 %
Current CD4 (cells/ml)	523.32±270.92
Nadir CD4 (cells/ml)	189.27±159.10
Plasma HIV RNA (% undetectable)	71.62 %
HIV infection duration (years)	12.66±7.01
Receiving CART (%)	82.43 %
Current hepatitis C infection (%)	33.78 %
Lifetime Alcohol Dependence (%)	51.35 %
Lifetime Substance Dependence (%)	54.05 %

Note: CART = combination antiretroviral treatment. Continuous variables are reported as mean±SD. Proportions are reported in %

different aspects of systemic immune activation and inflammatory processes relevant to neuroAIDS based on findings by our group and others (Cohen et al. 2011; Ragin et al. 2010; Letendre et al. 2011). All cytokines reached the prerequisite concentration levels, and measurements were verified to be reliable across samples. Brief descriptions of measured cytokines are provided in Table 2.

MRI Data Acquisition and Analysis

All neuroimaging was performed on the same Siemens Tim Trio 3-tesla MRI scanner located at Brown University MRI Research Facility. High-resolution structural MRI of the whole brain was acquired in the sagittal plane using a T1-weighted MPRAGE pulse sequence with TE/TR = 3.06/2250 ms, flip angle = 9°, FOV = 220 mm, matrix=256×256, slice thickness = 0.86 mm. Head restraint was used during image acquisition and quality control of T1 images was performed immediately following acquisition to identify excessive movement and other artifacts. Repeated acquisitions were prescribed as necessary, and images with inadequate quality were excluded from analysis. Parcellation of brain regions was performed on each T1-weighted MRI image using automated algorithms implemented in Freesurfer, where anatomical labels are assigned to each voxel based on probabilistic estimates after a nonlinear registration to an atlas, and volumetric measures were derived for each brain (Fischl et al. 2004). Combined bilateral volumes of cortical grey matter (GM), cerebral white matter (WM), caudate, putamen, globus pallidus, thalamus, hippocampus, amygdala, and overall ventricles

were thus derived (Fig. 1). Intracranial volumes were also measured and included in all statistical models to control for variability in head size.

Statistical Analysis

All statistical analyses were performed in R version 2.13.0 (<http://www.r-project.org>). A series of linear multiple regression models were used to examine the independent contributions of clinical variables and cytokine markers on each brain volumetric measure. Each full model consisted of clinical variables (age, current and nadir CD4 levels, detectable plasma HIV RNA, years of HIV infection, current CART regimen, and detectable serum HCV RNA), thirteen cytokine measures as noted above, and intracranial volumes to control for head size. Age, CD4 levels, and infection duration were included as continuous covariates, while the remaining clinical variables were included as dummy coded binary covariates. Cytokine and nadir CD4 levels were log-transformed to handle the skewed distributions of these variables. Due to the large number of variables in the full models under consideration, Akaike information criterion (AIC) was used to select the subset of markers that best predicted each brain volume (Akaike 1974). Final linear regression models were selected by minimizing AIC, which takes into account both the goodness of fit and the model complexity.

Results

Table 3 shows the unstandardized regression coefficients and associated p-values for clinical and cytokine markers that were included in the final regression model following the model selection algorithm for each brain volumetric measure. Results are also presented graphically in Fig. 2. As expected, intracranial volume was a highly significant covariate and was included in all final models to control for head size.

Global Volumetric Measures

Cortical Grey Matter Lower GM volume was significantly associated with older age, in addition to lower level of IFN- γ and higher levels of IL16 and IL18.

Cerebral White Matter Lower WM volume was significantly associated with longer duration of HIV infection, lack of current CART, in addition to lower IFN- γ level and higher IL6 level.

Table 2 Brief descriptions of measured cytokines

IFN- γ	Interferon-gamma	Produced by innate NK cells, acquired antigen-specific cytotoxic CD4+ and effector CD8+ T cells. Activates macrophages and critical for innate and adaptive immune responses to intracellular pathogens, tumor control, and inhibition of viral replication.
IL-1 β	Interleukin-1 beta	Produced by activated macrophages; mediates inflammatory responses, cell proliferation, apoptosis. Induces Cox-2 in CNS, causing inflammatory pain
IL-6	Interleukin-6	Secreted by T cells and macrophages; triggers inflammation, acute phase response, fever. Anti-inflammatory effects include inhibiting TNF- α and IL-1, and activating IL-1ra and IL-10.
IL-8	Interleukin-8	Made by macrophages and some epithelial and endothelial cells; Role in innate immune response. Major role in chemotaxis of neutrophils. Also mediates inflammatory response and angiogenesis.
IL-10	Interleukin-10	Produced by monocytes. Pleiotropic cytokine. As an anti-inflammatory cytokine, it inhibits macrophage and dendritic cell function, suppresses TNF- α . Acquires pro-inflammatory activity during immune response with IFN- α stimulation.
IL-16	Interleukin-16	Secreted by lymphocytes. Pleiotropic cytokine. Functions as a chemoattractant (CD4+ cells), modulates T cell activation, and inhibits HIV replication.
IL-18	Interleukin-18	Produced by macrophages and monocytes. Pro-inflammatory cytokine interacts with IL-12 to induce cell-mediated immune response with microbial infection and LPS, inducing severe inflammatory reactions. Stimulates NK and T cell release of IFN- γ , which activates macrophages. Inhibits IL4-dependent IgE, enhances B cell production.
IP-10	Interferon-inducible protein 10	Produced by various cell types including monocytes, endothelial cells, fibroblasts, keratinocytes. Induced by IFN- γ and TNF- α . Chemoattractant for activated T cells.
MCP-1	Monocyte chemoattractant protein-1	Expressed in monocytes, vascular endothelial cells, smooth muscle cells. CCL2 chemokine, induces monocyte attraction, and degranulation of basophils with histamine release. Induced by IL-1, TNF- α , PDGF, TGF- β , and LIF
MIP-1 β	Macrophage inflammatory protein-1 beta	Produced by macrophages. CCL4 chemokine that generates local inflammatory responses, induces superoxide production by neutrophils. Chemotactic activity for lymphocytes, macrophages, NK cells, and monocytes with inflammation; down-regulates CCR5, inhibiting HIV-1 blocking.
SDF-1 α	Stromal cell-derived factor-1 alpha	Expressed ubiquitously, except in blood cells. Small cytokine member of CXCL12 family of chemokines. Activates leukocytes due to strong chemotactic effects. Induced by pro-inflammatory stimuli, e.g. TNF- α and IL-1 β .
TNF- α	Tumor necrosis factor-alpha	Secreted by macrophages, monocytes, neutrophils, T cells, NK cells after stimulation with LPS. CD4+ cells secrete TNF- α . Also made by astrocytes, microglial cells, smooth muscle cells, and fibroblasts. Mediates systemic inflammation, inhibits viral replication, and inhibits tumorigenesis.
TRAIL	TNF-related apoptosis-inducing ligand	Expressed broadly in tissues. Cytokine induces proapoptotic caspase activity by up-regulating pro-apoptotic Bcl proteins. Causes apoptosis in hepatocytes, neural cells, and thymocytes

Fig. 1 Segmented T1-weighted axial slices showing volumes of interest, including cortical grey matter (red), cerebral white matter (white), caudate (light blue), putamen (pink), globus pallidus (dark blue), thalamus (green), hippocampus (yellow), amygdala (teal), and overall ventricles (shown unmasked)

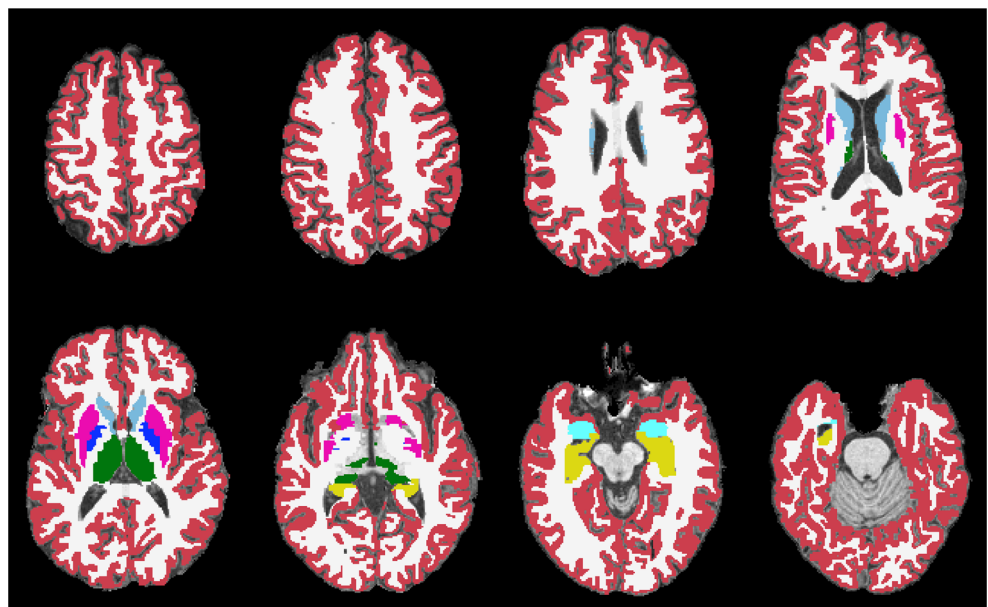


Table 3 Unstandardized regression coefficients and associated p-values (in parentheses) for clinical and cytokine markers included in the final model selected by minimizing AIC for each brain volumetric measure. Rows indicate the predictors under consideration, and each final

regression model is represented by one column of the table. Statistically significant covariates are highlighted in grey. AIC and adjusted R² values of the final models are included in the final rows. Note: ICV = intracranial volume

	Cortex	White Matter	Caudate	Putamen	Globus Pallidus	Thalamus	Hippocampus	Amygdala	Ventricles
Age	-1.95e+3 ($<.0001$)	-7.30e+2 (0.0635)	-1.62e+1 (0.0764)	-4.30e+1 (0.0001)	-1.18e+1 (0.0019)	-5.81e+1 ($<.0001$)	-2.10e+1 (0.0093)	-8.44e+0 (0.0428)	4.41e+2 ($<.0001$)
CD4 current				-5.60e-1 (0.1468)				2.78e-1 (0.0626)	7.59e+0 (0.0353)
CD4 nadir				1.75e+2 (0.0204)	4.73e+1 (0.0795)	1.97e+2 (0.0311)	1.21e+2 (0.0445)		
HIV RNA									
HIV duration		-1.21e+3 (0.0355)				-3.18e+1 (0.0672)			
CART		2.58e+4 (0.0166)				6.86e+2 (0.0342)	6.35e+2 (0.0053)		
HCV			3.48e+2 (0.0964)	4.96e+2 (0.0361)	3.48e+2 (0.0009)				
IFN- γ	2.23e+4 (0.0114)	2.41e+4 (0.0306)				4.79e+2 (0.1370)	4.24e+2 (0.0687)		
IL1- β	-1.22e+4 (0.1490)				-2.32e+2 (0.0069)				
IL6		-2.19e+4 (0.0095)		-6.21e+2 (0.0030)		-8.36e+2 (0.0021)	-4.82e+2 (0.0081)		6.56e+3 (0.0003)
IL8									
IL10								-2.53e+2 (0.0526)	-9.03e+3 (0.0190)
IL16	-2.48e+4 (0.0001)		-2.55e+2 (0.1253)	-6.63e+2 (0.0014)	-2.18e+2 (0.0038)		-2.37e+2 (0.1263)	-2.04e+2 (0.0091)	
IL18	-1.21e+4 (0.0023)					2.37e+2 (0.1184)	1.52e+2 (0.1436)		
IP10	3.82e+3 (0.1153)	4.99e+3 (0.1240)			-5.15e+1 (0.1450)				
MCP1	7.44e+3 (0.0924)			4.01e+2 (0.0127)	1.38e+2 (0.0150)	3.03e+2 (0.0691)		1.29e+2 (0.0218)	
MIP-1 β			1.39e+2 (0.1325)						2.47e+3 (0.0184)
SDF1- α									
TNF- α									
TRAIL				4.46e+2 (0.0663)					
ICV	2.31e-1 ($<.0001$)	2.70e-1 ($<.0001$)	4.87e-3 ($<.0001$)	4.51e-3 ($<.0001$)	1.64e-3 ($<.0001$)	7.57e-3 ($<.0001$)	3.78e-3 ($<.0001$)	1.63e-3 ($<.0001$)	2.17e-2 (0.0001)
Final AIC	1494.88	1528.72	977.03	999.36	842.38	1016.12	959.31	865.53	1336.36
Adjusted R ²	0.8377	0.7692	0.6283	0.6764	0.6289	0.7893	0.6477	0.5357	0.4142

Ventricles Higher ventricular volume was significantly associated with age and higher current CD4 level, in addition to higher levels of IL6 and MIP-1 β , and lower IL10 level.

Subcortical and Medial Temporal Structures

Caudate Caudate nucleus volume was not significantly associated with any clinical or cytokine markers.

Putamen Lower putamen volume was significantly associated with older age, lower CD4 nadir level, undetectable HCV RNA, in addition to higher levels of IL6 and IL16, and lower MCP1 level.

Globus Pallidus Lower pallidum volume was significantly associated with older age, undetectable HCV RNA, in addition to higher levels of IL1- β and IL16, and lower MCP1 level.

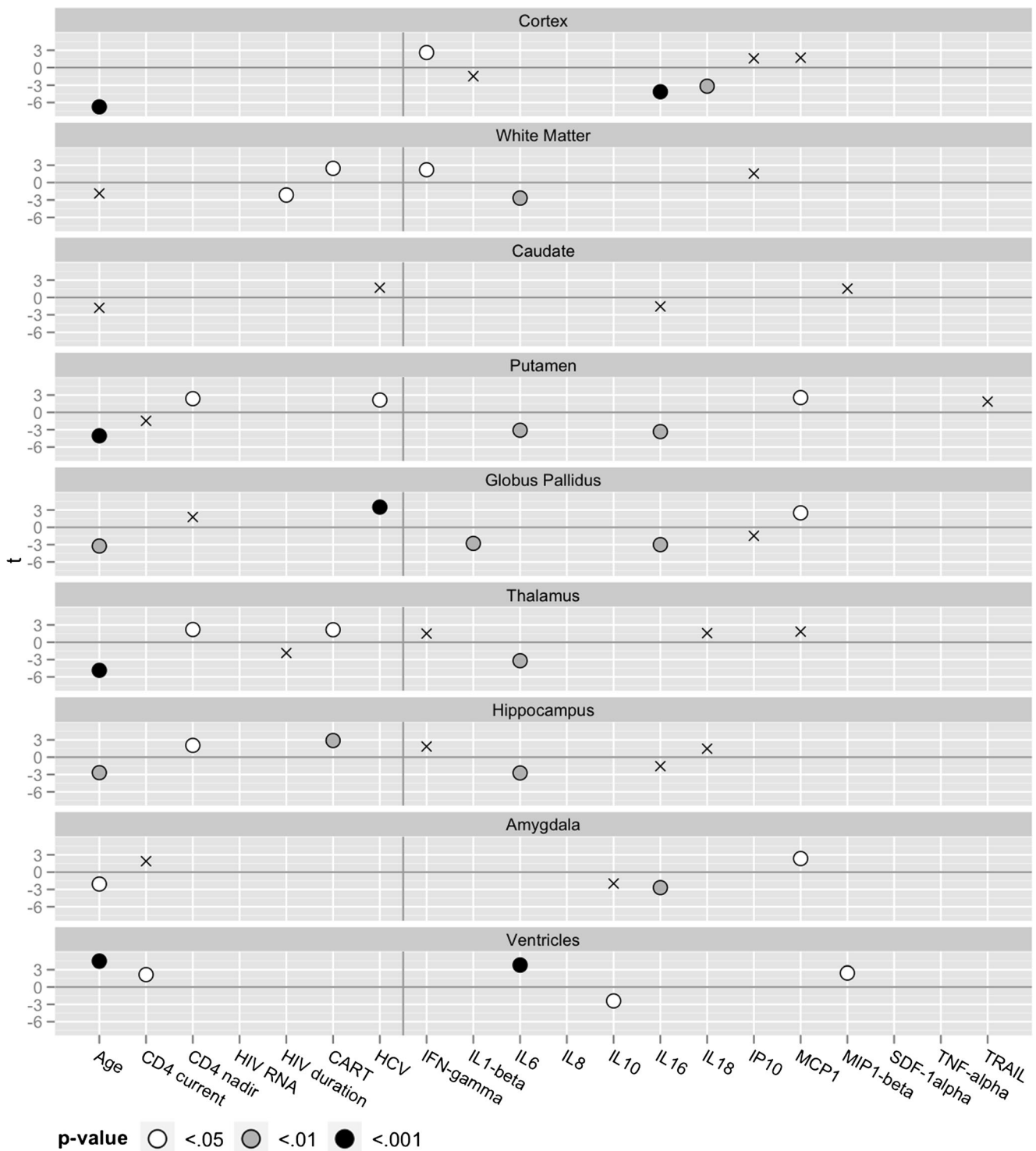


Fig. 2 Final linear regression models selected by minimizing AIC. Each model includes the subset of clinical and cytokine markers that best predict each brain volumetric measure. Significant predictors are shaded to indicate p-value levels (see legend). Non-significant predictors retained

in the models are marked with crosses (X). Regression coefficients were transformed to *t*-statistics to scale the plots for visualization. Refer to Table 3 for individual regression coefficient estimates and associated p-values of all covariates

Thalamus Lower thalamus volume was significantly associated with older age, lower CD4 nadir level, and lack of current CART, in addition to higher IL6 level.

Hippocampus Lower hippocampal volume was significantly associated with older age, lower CD4 nadir level, lack of current CART, in addition to higher IL6 level.

Amygdala Lower amygdala volume was significantly associated with older age ($p=0.0428$), in addition to higher IL16 level and lower MCP1 level.

Impact of Lifetime Alcohol and Substance Use Disorders

To further examine the possible influence of more distant alcohol/substance use history, we repeated the regression analyses predicting each volumetric measure with an additional binary covariate indicating lifetime alcohol or substance dependence history based on DSM-IV criteria. Following the model selection algorithm, this variable was retained in models predicting volumes of the amygdala ($\beta=135.97$, $p=0.1420$), putamen ($\beta=-430.05$, $p=0.0797$), and thalamus ($\beta=797.85$, $p=0.0018$). The addition of this variable did not alter the patterns of the volumetric relationships with other predictors in these three models, while all other models remain identical to those reported above.

Discussion

Initial optimism that CART would lead to an eradication of HIV-associated brain dysfunction has been tempered by persistent neurocognitive impairment observed in people living with HIV despite virologic suppression. Chronic systemic immune activation and inflammation have been among the mechanisms proposed for these disturbances, though the underlying pathophysiology is still not fully understood. Using a multivariate statistical approach, we examined the independent contributions of age, clinical status, and an extensive panel of plasma cytokine markers on brain volumetric alterations in this cohort of 74 medically stable individuals with chronic HIV infection. Our findings corroborate existing evidence demonstrating pervasive brain atrophy in the context of HIV disease and older age (Jernigan et al. 2011; Lockhart and DeCarli 2014). These structural alterations appear to be partially mitigated by antiretroviral treatment. Cytokine levels, most notably IL-6, IL-16, and MCP-1, were significantly associated with brain volumes even after accounting for the effects of the above clinical variables. The current findings demonstrated the potential utility of these markers both in elucidating the pathophysiological mechanisms underlying brain damage associated with HIV infection, and as systemic markers of HIV-associated brain dysfunction in the CART era.

A major strength of this study is the use of multiplexed assay to simultaneously investigate an extensive list of markers potentially relevant to brain structural abnormalities. Our findings extend those from previous studies in a number of ways. This is among the first studies to demonstrate associations between plasma cytokine and structural brain

alterations, in addition to previously reported associations to neurocognitive dysfunction (Cohen et al. 2011; Ryan et al. 2001). To our knowledge, one published study has examined this link with brain volumetric alteration in a relatively small cohort of 10 HIV-infected individuals (Ragin et al. 2010). In addition to expanding the sample size, we also examined volumes of specific subcortical nuclei and medial temporal structures, along with the global volumetric measure of the grey/white matter and the ventricles as were reported in the previous study. The use of a multivariate model selection approach permitted the independent contributions and relative importance of each marker to be examined, while ensuring parsimony of the statistical models.

IL-6 and IL-16 emerged as the most robust cytokine predictors of smaller brain volumes in these models, together accounting for alterations in virtually all brain regions examined. These findings are consistent with expectation given the principal roles of these molecules in systemic immune responses and inflammatory processes. IL-16 in particular is an important chemoattractant cytokine for CD4+ T cells and contributes to the regulatory process of recruitment and activation of T cells at inflammation sites (Cruikshank et al. 2000). Corroborating the current findings, a previous study from our group showed IL-16 elevation to be among the strongest predictors of impairment in attention and executive and psychomotor functions (Cohen et al. 2011). IL-6 functions as both a pro- and anti-inflammatory cytokine and has been associated with HIV RNA level and implicated as a mediator of activation-induced CD4+ T cell losses (Lederman et al. 2000; Scheller et al. 2011). Although CSF IL-6 level has been linked to neurocognitive impairment in treatment-naïve HIV-infected individuals (Airoldi et al. 2012), to our knowledge, no previous studies have shown associations between systemic IL-6 level and structural brain measures in the context of HIV. Less widespread associations were found between higher IL-1 β , IL18, and MIP-1 β levels and smaller volumes of the globus pallidus and cerebral cortex, and larger ventricular volume. These latter findings are likewise novel and should be considered in the context of future replicating studies.

Higher MCP-1 levels were associated with greater basal ganglia and amygdala volumes. This is contrary to expectation given previous findings linking neurocognitive impairment and cerebral white matter and metabolite abnormalities to increased MCP-1 levels both in plasma (Cohen et al. 2011; Ragin et al. 2010; Woods et al. 2006) and CSF (Letendre et al. 2011; Yuan et al. 2013). Plasma level of MCP-1 has also been positively related to HIV RNA level (Weiss et al. 1997). MCP-1 is an important chemotactic factor for monocytes and CD4+ T cells, and serves to recruit immune cells to the site of inflammation (Ajuebor et al. 1998). Given its direct involvement with these immune cells, in addition to the fact that monocytes/macrophages are major producers of these

molecules, MCP-1 is believed to have important roles in HIV pathogenesis (Deshmane et al. 2009). While the basis for the positive relationships between MCP-1 and brain volumes cannot be ascertained from this study, it is possible that these findings reflect hypertrophy resulting from fluid accumulation secondary to inflammation of these brain structures. This possibility is further supported by findings suggesting that MCP-1 in plasma may largely be derived originally from within the CNS (Monteiro de Almeida et al. 2005).

Among the HIV disease markers, nadir CD4 suppression was most consistently associated with smaller brain volume. This is consistent with emerging literature indicating the effects of distant history of immune suppression on current brain integrity (Jernigan et al. 2011; Cohen et al. 2010; Ellis et al. 2011). These findings relate to the possibility that viral replication and the resulting immune suppression occurring at an early stage of HIV disease have long term effects on the brain that may persist even after effective viral suppression and immune reconstitution. It is notable that the current findings contrast with recently published results from our group indicating a lack of association between nadir CD4 and diffusion tensor MRI measures of white matter integrity (Gongvatana et al. 2011). This dissociation may reflect differential effects of distant immunologic history on macroscopic (i.e., volumetric alteration) versus microstructural (i.e., cellular water diffusion) measures of brain integrity. Although it is possible that chronic measures such as nadir CD4 may also be a proxy for secondary factors such as poor access to health care, comorbid medical conditions, and socioeconomic status, these findings suggest the potentially beneficial role of early antiretroviral treatment to prevent significant immune suppression.

The finding related to duration of HIV infection is consistent with existing evidence linking general brain atrophy to infection chronicity (Becker et al. 2011; Elovaara et al. 1990). Specifically, our finding in this medically stable, primarily CART-treated cohort suggests particular susceptibility of cerebral white matter to chronic infection. The positive association between current CD4 and ventricular volume was contrary to expectation, and needs to be further examined in future studies. As expected, lack of antiretroviral treatment was associated with smaller brain volumes, including those of the white matter, thalamus, and hippocampus. These findings suggest a beneficial role of CART in mitigating brain atrophy in the context of HIV infection. It should be noted that the observed relationships with CART were not in the direction of antiretroviral toxicity as has been reported in recent studies (Ciccarelli et al. 2011).

To further examine the impact of CART history in this chronically infected cohort, we classified participants based on estimated year of infection. Using 1995 as the cut point for CART availability, 41 % of our sample was diagnosed with HIV pre-CART. As expected, individuals diagnosed prior to 1995 were significantly more likely to have history of immune

suppression: 77 % of pre-CART individuals vs. 41 % of those diagnosed in the CART era presented with CD4 nadir < 200, $\chi^2=7.8404$, $p=0.0051$. It should be noted that while this difference in immune suppression history between the groups is likely related to CART availability at the time of diagnosis, it is likely also impacted by the longer duration of illness in the pre-CART group ($t=16.08$, $p<.0001$). We repeated the regression analyses reported above with an additional binary covariate indicating an HIV diagnosis before or after 1995. Interestingly, this additional predictor did not show significant relationships with any of the volumetric variables. While it is possible that these statistical outcomes were impacted by multicollinearity with HIV disease factors discussed above, it should be noted that our binary classifier based on diagnosis year likely failed to capture the complex and diverse antiretroviral treatment time course among these individuals, and future studies with more comprehensive assessment of ART history are needed to adequately examine this important question.

HCV coinfection was found to be associated with larger volumes of basal ganglia structures. Basal ganglia hypertrophy has been previously reported in the context of substance abuse (Jernigan et al. 2005; Jacobsen et al. 2001), and it is possible that the current findings reflect analogous neuroinflammatory processes specifically affecting these structures. HCV infection has been linked to microglial activation and neuroinflammation (Grover et al. 2012). The extent to which such events are contributed by direct viral infiltration of the CNS or indirectly through HCV-associated hepatic disturbances is unclear. However, multiple lines of evidence indicating direct HCV neuroinvasion have recently emerged (Laskus et al. 2005), pointing to the importance of virologic control in preventing brain dysfunction. Regardless of the underlying mechanism, existing findings clearly indicate the presence of neuroinflammatory processes in HCV-infected individuals (Bokemeyer et al. 2011; Forton et al. 2001, 2008). The pathophysiological basis of brain dysfunction associated with HCV infection, particularly in the context of HIV coinfection, remains to be fully delineated. Previous findings from our group and others have linked HIV/HCV coinfection to increased neurocognitive impairment (Cherner et al. 2005; Clifford et al. 2005; Hinkin et al. 2008; Devlin et al. 2012) and cerebral white matter abnormalities (Jernigan et al. 2011; Gongvatana et al. 2011). Additional studies are needed specifically examining neuroinflammation in the context of HIV/HCV coinfection.

While the current analyses excluded individuals with alcohol or substance use disorder within 6 months prior to the study, the possible influence of more distant alcohol/substance use history cannot be ruled out. To further examine this, we repeated the regression analyses predicting each volumetric measure with an additional binary covariate indicating lifetime alcohol or substance dependence history. Following the

model selection algorithm, this variable was a significant predictor only of thalamic volume. While hypertrophy of the basal ganglia in the context of substance abuse has been reported (Jernigan et al. 2005; Jacobsen et al. 2001), the significant positive relationship between thalamic volume and lifetime alcohol/substance dependence in this cohort is a noteworthy and novel finding. Future studies that include more detailed assessment of distant alcohol and substance use are needed to further address this important question.

Not surprisingly, older age was among the most robust predictors of smaller volumes in virtually all measured brain structures. Although these findings are consistent with the substantial literature on age-related brain atrophy (Zimmerman et al. 2006; Brickman et al. 2006; Raz et al. 2010), it is possible that the robust associations observed here may reflect the augmented effects of aging in the context of HIV such as has been suggested (Valcour et al. 2004; Kirk and Goetz 2009). Future studies are needed that are adequately powered to examine the possibility of an HIV-associated accelerated aging phenomenon relevant to structural and functional brain changes.

In summary, the current findings provide novel evidence that cortical, subcortical and white matter volumes vary as a function of plasma cytokine levels among people with chronic HIV infection. These results from a cohort with an average infection duration of about 13 years corroborate an emerging body of evidence linking HIV-associated chronic systemic immune activation and inflammatory processes to compromises in brain integrity (Williams et al. 2005; Lentz et al. 2009; Pulliam et al. 2011). The fact that these cerebral correlates were observed for plasma cytokine levels is noteworthy, and indicates potential utility of these peripheral markers in predicting and monitoring brain alterations associated with chronic HIV infection. The relative importance of specific cytokines to brain structural integrity demonstrated here also contributes to the current understanding of mechanisms underlying HIV neuropathogenesis, which may aid in the development of beneficial therapies for HIV-associated neurocognitive dysfunction.

Inferences from this study are limited by its cross-sectional and observation nature. Future longitudinal studies are necessary to examine the causal relationships between the immunologic and inflammatory markers with structural and functional brain measures, especially in the context of antiretroviral treatment, aging, and comorbid conditions including HCV infection. Simultaneous examinations of the CNS and systemic trajectories of cytokine levels, in addition to those of other biomarkers, will also be important to further elucidate the underlying mechanisms of abnormal brain alterations in people living with HIV. Finally, novel neuroimaging markers are rapidly being developed that enable concurrent examination of multiple aspects of brain integrity, including those relevant to structural and functional

connectivities and vascular integrity, which together with other biomarkers will help to extend the current understanding of the underlying pathophysiological mechanisms of HIV-associated brain dysfunction.

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