

HHS Public Access

Author manuscript *AIDS*. Author manuscript; available in PMC 2021 March 01.

Published in final edited form as: *AIDS*. 2020 March 01; 34(3): 415–426. doi:10.1097/QAD.0000000002436.

Regional Brain Volumetric Changes Despite Two Years of Treatment Initiated During Acute HIV Infection

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Abstract

Objective: To assess changes in regional brain volumes after 24 months among individuals who initiated combination antiretroviral therapy (cART) within weeks of HIV exposure.

Design: Prospective cohort study of Thai participants in the earliest stages of HIV-1 infection.

Methods: Thirty-four acutely HIV-infected individuals (AHI; Fiebig I-V) underwent brain magnetic resonance (MR) imaging and MR spectroscopy at 1.5T and immediately initiated cART. Imaging was repeated at 24 months. Regional brain volumes were quantified using FreeSurfer's longitudinal pipeline. Voxel-wise analyses using tensor-based morphometry (TBM) were conducted to verify regional assessments. Baseline brain metabolite levels, blood and cerebrospinal fluid biomarkers assessed by ELISA, and peripheral blood monocyte phenotypes measured by flow cytometry were examined as predictors of significant volumetric change.

Results: Participants were 31 ± 8 years old. The estimated mean duration of infection at cART initiation was 15 days. Longitudinal analyses revealed reductions in volumes of putamen (p<0.001) and caudate (p=0.006). TBM confirmed significant atrophy in the putamen and caudate, as well as in thalamic and hippocampal regions. In exploratory post-hoc analyses, higher baseline frequency of P-selectin glycoprotein ligand-1 (PSGL-1)-expressing total monocytes correlated with greater caudate volumetric decrease (ρ =0.67, p=0.017), while the baseline density of PSGL-1-expressing inflammatory (CD14⁺CD16⁺) monocytes correlated with putamen atrophy (ρ =0.65, p=0.022).

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Conclusion: Suppressive cART initiated during AHI may not prevent brain atrophy. Volumetric decrease appears greater than expected age-related decline, although examination of longitudinal change in demographically similar HIV-uninfected Thai individuals is needed. Mechanisms underlying progressive HIV-related atrophy may include early activation and enhanced adhesive and migratory capacity of circulating monocyte populations.

Keywords

Acute HIV infection; brain atrophy; putamen; caudate; monocytes; biomarkers; metabolites

INTRODUCTION

The neuroimaging signature of chronic HIV infection includes gray and white matter atrophy [1-3]_ENREF_1, disrupted frontostriatal functional connectivity [4, 5], and abnormal white matter (WM) structural connectivity patterns [6]. Many neuroHIV studies have included participants with advanced disease or severe immune compromise prior to treatment [7-10]. However, reductions in total and cortical gray matter (GM) volumes have been identified even during primary HIV infection (PHI, < 1 year post-transmission) [11], and decreased brain parenchymal volume was observed within the first 100 days [12]. Initial brain alterations may be determined by the severity of immunosuppression [13].

Early initiation of combination antiretroviral therapy (cART) is associated with greater probability of restoring normal CD4 counts [14, 15]_ENREF_9 and appears protective. Neurofilament light protein, a marker of ongoing HIV-related neurodegeneration [16], is elevated in cerebrospinal fluid (CSF) during untreated PHI [17] but is normal after 6 months of cART initiated during acute HIV infection (AHI) [18]. Treatment that is implemented within 6 months of infection greatly decreases CD4+ T-cell activation and HIV DNA and RNA reservoir size [19, 20]. The HIV DNA reservoir in peripheral blood mononuclear cells (PBMC), established early in infection [21], predicts HIV disease progression and the time to viremic rebound following cessation of cART [22, 23]. We and others have shown that peripheral monocytes harboring HIV in the setting of chronic HIV infection, and monocyte phenotypic changes associated with migration and activation, are linked to a range of brain abnormalities that persist despite therapy [24]. Plasma elevations of soluble CD163 (sCD163) in PHI correlate with the proportion of CD14⁺CD16⁺ monocytes and decrease with effective cART [25]. Similar changes have been observed in AHI [26].

HIV invasion of the central nervous system (CNS) occurs early. The virus has been isolated in the brain 15 days after iatrogenic HIV infection, only one day after it was first found in blood _ENREF_28[27]. _ENREF_28Computational modeling of clinical data has identified the first three weeks post-infection as a window of opportunity in which cART initiation may optimally control HIV replication and prevent immune activation and the establishment of viral reservoirs [28]. Whether HIV-induced brain injury can be arrested or slowed by immediate treatment remains unknown. SEARCH010/RV254, a longitudinal cohort study of AHI, enrolls and evaluates participants 2–3 weeks following history of exposure. Thereafter cART is commenced at once. Matched samples at baseline and two-year follow-up are a strength of this study. The current work evaluated SEARCH010/RV254 participants for changes in volumes of nine *a priori* brain regions of interest after 24 months of cART. Although HIV-infected individuals over age 60 have exhibited progressive brain atrophy despite > 3 years of viral suppression [29], previous longitudinal studies suggest that early cART may prevent such injury. In a PHI cohort (infected for ~3.7 months and followed for up to 6 years), duration of untreated infection correlated with brain volume loss prior to cART initiation; however, after cART no further atrophy was observed [30]. Two-year changes in subcortical volumes of aviremic chronically HIV-infected individuals on cART were comparable to those of HIV-negative controls, although the HIV+ group had smaller volumes [31, 32]. We therefore hypothesized the absence of HIV-related brain atrophy; namely, volumetric loss beyond that attributable to normal aging, which in the SEARCH010/RV254 cohort would be small. The link between brain atrophy in chronic, stable HIV disease and a history of immunosuppression supports our conjecture that very early antiretroviral intervention may prevent brain damage [33].

Brain volume deficits have been associated with disrupted levels of cerebral metabolites [33] and plasma cytokines [34] in chronic HIV infection. Plasma neopterin, a marker of immune activation, correlates with peripheral blood HIV DNA burden [35] which in turn relates to HIV-associated brain injury [36] and cognitive disorders [35]. Soluble CD14 (sCD14) is a marker of monocyte activation [37]; high plasma sCD14 levels predict morbidity and mortality in HIV [38]. CSF concentrations of the neuronal injury marker S100B may be elevated in HIV-infected individuals [39] and, in Alzheimer's disease, correlate with brain atrophy [40]. Brain inflammation at baseline in the SEARCH010/RV254 cohort has been identified by magnetic resonance spectroscopy (MRS) and by elevated CSF levels of inflammatory markers such as neopterin [41, 42]. The present work investigated baseline brain metabolites as predictors of volumetric changes found to be significant. Exploratory analyses examined plasma and CSF biomarkers, and monocyte phenotypes potentially implicated in HIV neuropathogenesis, for their predictive value.

METHODS

Participants

Our study sample was drawn from the first 38 AHI enrollees with magnetic resonance imaging (MRI) in the SEARCH010/RV254 study in Bangkok, Thailand (ClinicalTrials.gov #) [41, 43]. Participants were >18 years old, had confirmed acute HIV-1 infection, were cART-naïve, and consented to initiating protocol-defined cART. Testing was performed to categorize them into Fiebig stages I to V by a hierarchical algorithm from pooled nucleic acid, HIV RNA, sequential immunoassay, p24 antigen, and Western Blot testing [44]: Fiebig I (RNA+, p24 antigen–), Fiebig II (p24 antigen+, IgM–), Fiebig III (IgM+, Western Blot–), Fiebig IV (Western Blot indeterminate), and Fiebig V (Western Blot+ without p31) [45]. Neurologic interviews and examinations were performed by physicians at study entry and longitudinally, using tools developed by the AIDS Clinical Trials Group [46]. Treatment was initiated immediately after baseline evaluations. Exclusion criteria for this sub-study were excessive motion on MRI and poor cART adherence resulting in plasma HIV RNA (viral load, VL) >200 copies/mL at 24 months. One individual who chose not to start treatment

soon after entry was dropped from analysis. Two were excluded because of poor image quality and one because of high 24-month VL, yielding a study sample of 34 with baseline and two-year follow-up data (baseline MRI dates: 5/15/2009–11/2/2012). Antiretroviral combinations consisted of efavirenz, tenofovir, and emtricitabine (standard cART), with 20 of the 34 participants randomly assigned to receive additional maraviroc and raltegravir (cART intensification, or "mega-cART") during the first 24 weeks. Viral load was also assessed at intermediate timepoints (weeks 4, 12, 24, 36, 48, 72 post-baseline).

All participants underwent hepatitis C antibody testing, blood Treponema pallidum hemagglutination assay (TPHA), and either Venereal Disease Research Laboratory (VDRL) or Rapid Plasma Reagin (RPR) testing (hereafter denoted as VDRL). Presence of chlamydia or gonorrhea was determined by clinical evaluation. Syphilis and treatment history were collected through interviews conducted by research physicians at enrollment. Syphilis (at the time of AHI) was defined by positive TPHA and VDRL in the absence of prior syphilis treatment.

Information on substance use was obtained for most participants by physician interviews conducted informally at study entry. Participants were asked whether they had used any illicit substance during the 4 months prior to enrollment. Those who responded affirmatively were questioned about their use, within the same period, of substances including cannabis, poppers (inhaled nitrites), amphetamines (with street names such as speed, whizz and base), and methamphetamine (crystal meth, ice, glass, tina).

This study was approved by institutional review boards of the participating sites. All participants provided signed written consent.

Neuroimaging data acquisition and processing

Anatomical MRI data at both timepoints were acquired by an axial 3D T1-weighted spoiled gradient echo image (3D SPGR, TE= 7 ms, TR= 11.2 ms, flip angle=25°, with 1 mm voxel resolution) on a GE Signa HDx 1.5T clinical scanner (GE Healthcare, Milwaukee, WI, software version 12-M4) using an 8-channel head coil for data reception and a standard body coil for transmission. Single-voxel MRS was performed using a double spin echo data acquisition (PROBE-P) with TE= 35 ms, TR= 1.5s at four locations: left frontal white matter (FWM, 8cc), midline frontal gray matter (FGM, 8cc), occipital gray matter (OGM, 8cc) and basal ganglia (BG, 12cc) [42]. At the end of each scan, single-voxel MRS was conducted on a standard spectroscopy phantom (GE Healthcare) to verify scanner stability [42, 47]. MRI/MRS data acquisition and quality assurance were performed by an experienced technician.

To obtain regional brain volumes, T1-weighted MRI data at baseline (month 0, or M0) and month 24 (M24) were processed using the longitudinal pipeline [48] of the FreeSurfer image analysis suite (http://surfer.nmr.mgh.harvard.edu/, v.5.3)[49]. Quality assurance of the FreeSurfer-derived volumes was performed by visual inspection of cortical and subcortical segmentations. Manual corrections were applied when necessary and were followed by reprocessing, a method that improves gray/white segmentation [50]. Total volumes of regions of interest (caudate, putamen, pallidum, thalamus, hippocampus, amygdala, nucleus

accumbens, cortical GM, and cerebral WM) were computed by summing over left and right brain hemispheres. Baseline MRS data were processed and levels of N-acetylaspartate (NAA), choline (Cho), creatine (Cr), myoinositol (MI) and glutamate-glutamine (Glx) quantified as previously reported [42].

Viral load measurements

HIV-1 RNA (VL) was measured in M0 and M24 paired plasma and CSF samples using the Roche Amplicor HIV-1 Monitor, Test v1.5 or the Roche COBAS AmpliPrep/COBAS TaqMan HIV-1 Test, v2.0 (Roche Diagnostics, Branchburg, NJ, USA); the lower limits of plasma VL detection were 50 and 20 copies/mL, respectively. The lower limit of detection of CSF HIV RNA was 80 copies/mL.

Clinical and biological markers and monocyte populations

Risk factors for HIV-associated cognitive impairment and/or decreased brain volumes include high plasma and CSF HIV RNA [7]; low CD4 count (especially nadir) [7, 51]; and elevated plasma sCD163 [52] and cytokines TNF-a, IL-1 β , IL-6, MCP-1, and IP-10 [34]. To identify factors that may help to preserve or monitor brain integrity, we assessed biomarkers measured in plasma and CSF (neopterin, sCD14, sCD163, IL-6, IL-1 β , TNF-a, MCP-1, IP-10); and CSF S100B protein, a marker of astrocytic activation seldom studied in HIV. Also examined were monocyte subpopulations classified by CD14 and CD16 expression: classical (CD14+CD16–), intermediate/inflammatory (CD14+CD16+), and non-classical/patrolling (CD14lowCD16+). These subsets were further analyzed for expression of CCR2, CCR5, CX3CR1, P-selectin glycoprotein ligand-1 (PSGL-1), and CD163.

Levels of soluble biomarkers were assessed in plasma at months 0 and 24 for all participants, and for CSF where available as not all participants elected to undergo the lumbar puncture procedure. Multiplex Luminex® array was used to measure TNF-α, IL-1β, IL-6, MCP-1, S100B (CSF only) (Millipore, Burlington MA) and sCD163 (BioRad, Hercules CA). Standard ELISA assays measured neopterin (GenWay Biotech, San Diego CA) and sCD14 (R&D Systems, Minneapolis MN). Monocyte phenotyping based on CD14, CD16, CCR2, CCR5, CX3CR1, PSGL-1 and CD163 expression was assessed by flow cytometry and is described elsewhere [26, 53]_ENREF_24. All data were analyzed by Prism 6.0 (GraphPad, La Jolla, CA) using a 4-parameter fit standard curve.

Statistical analysis

Wilcoxon signed-rank tests assessed regional volumetric changes from baseline (M0) to 24 months (M24). A Holm-Bonferroni correction for multiple comparisons was applied [54]. For each participant we computed percentage volume change from baseline: i.e., (M0-M24)/M0 \times 100. Wilcoxon signed-rank tests also evaluated changes in neuronal and inflammatory markers, and the glial marker S100B, from baseline to 24 months for individuals who had data at both timepoints. Univariate relationships between variables were assessed by Pearson or Spearman correlation, as appropriate.

For each brain region showing significant volumetric change, Spearman correlation was used to examine associations of baseline brain NAA, Cho, MI and Glx concentration ratios

over creatine with percentage volume change. Statistical significance was defined by p<0.05. Additionally, volumetric changes were investigated in relation to baseline plasma and intrathecal inflammatory markers and to baseline monocyte subsets; these analyses were exploratory as the predictor data were available only for participant subsets, limiting statistical power. SPSS V.23.0 (SPSS Inc., Chicago, Illinois, USA) was used.

Tensor-based morphometry

Longitudinal tensor-based morphometry (TBM) was performed as in Nir et al, 2019 [1], using ANTS registration software [55].

RESULTS

Participant characteristics

After quality control, we evaluated 34 acute HIV participants (29 male) with median (range) age=29.0 (18–48) years; estimated time since exposure=15 (4–32) days; CD4 count=398 (132–1127) cells/mm3; plasma VL=5.52 (2.78–7.49) log10 copies/mL; and who were at Fiebig stages I (N=12), II (N=2), III (N=17), IV (N=2), and V (N=1) at baseline. Demographic and clinical characteristics of the study sample at baseline and 24 months are summarized in Table 1. Optional CSF sampling was performed at study entry on 25 of the 34 participants.

At M0, the sample included one case of untreated systemic syphilis without neurosyphilis, one case of gonorrhea, and no cases of chlamydia. One individual was co-infected with the hepatitis C virus (HCV), which was not treated. The participant who had syphilis showed an elevated CSF white blood cell count at baseline; following systematic treatment, CSF white blood count levels returned to normal at week 24. Neurosyphilis was deemed unlikely as the individual was neuroasymptomatic and remained so. The participant with gonorrhea was treated immediately.

Data on illicit substance use were unavailable for 3 individuals, but the majority of our cohort (N=21, or 68%) denied use within the 4 months prior to baseline. Ten participants reported using illicit substances during that time: poppers (N=4); methamphetamine (N=3); poppers and methamphetamine 5(N=2); and no specific substance information (N=1; only "illicit drug use" was noted).

Clinical variables and inflammatory markers at baseline and follow-up

Plasma and CSF VL decreased from M0 to M24, while CD4 count increased (Table 1). As well, decreases were seen in plasma and CSF neopterin, with median reductions from baseline of 33.1% and 59.1% respectively. IP-10 levels dropped by 84.6% (in plasma) and 59.2% (CSF). Plasma sCD14, plasma IL-1β, and CSF TNF-α decreased also. Among the 10 participants who had CSF S100B data at both timepoints, CSF S100B showed a non-significant increase (median 970.3 to 988.6 pg/mL; p=0.11).

Review of data at weeks 4, 12, 24, 36, 48, 72 showed that plasma VL first dropped to <50 copies/mL at week 4 in 8 participants, week 12 in 15, week 24 in 9, and week 36 in 2. We note that 94% of cases had suppressed viremia by the 24-week follow-up. All participants

were fully suppressed by week 36 and all remained aviremic after achieving suppression. The median (IQR) time to suppression of plasma VL to below 50 copies/mL was 12 (6–12) weeks in participants on mega-cART vs. 18 (10–24) weeks in those on standard cART (p=0.17).

Regional brain volumetric changes after 24 months of cART

Volumetric reductions were observed in the putamen (p<0.001) and caudate (p=0.006) over the first 24 months of cART (Table 1). As computed by the median percentage change, caudate and putamen volumes decreased by approximately 3%. The changes were significant after Holm-Bonferroni correction and were subsequently examined for relationships with baseline brain metabolites, plasma and CSF biomarkers, and monocyte subsets.

TBM results can be visualized in Figure 3. Briefly, TBM showed that the most significant change from baseline occurred bilaterally in the putamen (up to 5.2% atrophy in 24 months, $p<5\times10^{-6}$, corrected), confirming FreeSurfer findings. Atrophy of the caudate was significant, although less striking at 2.5% ($p<1.4\times10^{-4}$, corrected). In the thalamus, atrophy of ~4% was detected over 24 months ($p<2\times10^{-4}$, corrected). Thalamic atrophy appears to be in regions representing the ventral posterior nucleus, pulvinar, lateral geniculate nucleus, and/or medial geniculate nucleus (individual nuclei cannot be distinguished without higher field strength and spatial resolution, so their identities were inferred from the observed locations of atrophy). The head of the hippocampus exhibited nearly 3% atrophy in 24 months ($p<2\times10^{-4}$, corrected). Significant volumetric loss was also noted outside subcortical regions; e.g., in the orbitofrontal cortex and the cingulate gyrus. Large portions of temporal white matter were observed to undergo atrophy of up to 3% ($p<2\times10^{-6}$, corrected), along with the isthmus of the corpus callosum (2.5% atrophy; $p<1\times10^{-4}$, corrected). In addition, CSF expansion was detected in sulcal regions.

Baseline clinical variables and regional brain volumetric change

Changes in caudate and putamen volumes did not correlate with age, years of education, baseline CD4 count, plasma VL, CSF VL, or mega-cART status. Suppression of plasma VL below 50 copies/mL by week 4 (N=8) was associated with a smaller decrease from M0 to M24 in caudate volume (median 0.66% vs. 3.1%, p=0.009), and a trend toward lower putamen volume loss (1.9% vs. 3.5%, p=0.07).

Baseline brain metabolites as predictors of regional brain volumetric change

Greater change (decrease) in putamen volume correlated with lower baseline Cho/Cr (ρ = -0.43, p=0.015) and lower Glx/Cr (ρ = -0.39, p=0.027) in frontal GM (Figure 1). The results do not survive Holm-Bonferroni correction for comparisons involving four metabolite ratios, four MRS voxel locations and two regional volumes. Brain metabolite ratios at baseline did not relate to reduced caudate volume.

Substance use, co-infections and regional brain volumetric change

To examine the possible confounding effects of substance use on our results, we compared percentage volume changes of caudate and putamen between participants who had used

illicit substances during the 4 months preceding enrollment (N=10) and those who had not (N=21). The groups did not differ in either putamen (Mann-Whitney p=0.72) or caudate (p=0.35) volume loss. Additionally, we assessed caudate and putamen atrophy from M0 to M24 separately in the users and non-users. For the putamen, p-values of 0.013 and p=0.012 for users and non-users, respectively, revealed volumetric changes in both groups (less significant than in the combined sample because of reduced power). Similarly, caudate volume loss was comparable between users (p=0.059) and non-users (p=0.063).

Volumetric loss in the putamen and caudate remained essentially unaltered when the three cases of co-infection with syphilis, gonorrhea or Hepatitis C at baseline were excluded from analyses.

Plasma and CSF biomarkers and monocyte subsets

Exploratory analyses examined baseline values of plasma and CSF biomarkers as predictors of percentage volume changes of the caudate and putamen. No significant correlations were found. Associations were also assessed between the volumetric changes and baseline blood monocyte subpopulations, including those expressing CCR2, CCR5, CX3CR1, PSGL-1, and CD163. Greater decrease in caudate volume correlated with higher total frequency and density (mean fluorescence intensity [MFI]) of PSGL-1-expressing monocytes at baseline (ρ =0.67, p=0.017 and ρ =0.60, p=0.038, respectively). Putamen volume decrease correlated positively with the baseline frequency (ρ =0.61, p=0.037) and density (ρ =0.65, p=0.022) of PSGL-1-expressing inflammatory monocytes. Volumetric associations with PSGL-1-expressing monocytes are plotted in Figure 2.

DISCUSSION

Contrary to expectations, AHI participants demonstrated decreased volumes of caudate and putamen after two years of cART. The finding that brain injury may occur despite very early implementation of therapy is somewhat surprising given the anticipated neuroprotective effects of acutely administered cART_ENREF_31. However, HIV invasion of the CNS within 8 days of exposure [41] and CNS inflammation during AHI [41] constitute indirect evidence that the impact of HIV on the brain is almost immediate and may compete with mitigating effects of concomitant cART initiation. Our results are bolstered by data from separate cohorts indicating early brain damage and preferential susceptibility of the putamen during PHI [11, 56]. Impaired cognition seen in a subset of SEARCH010/RV254 participants after 6 months of treatment [57] suggests that brain injury may be incompletely ameliorated. Moreover, consistent with our findings, the cognitive deficits were manifested in domains in which basal ganglia play crucial roles (psychomotor speed, fine motor skills, and executive functioning [57-59]). Reduced putamen volume in PHI has been correlated with psychomotor slowing [56], and basal ganglia reductions in chronic HIV disease are associated with motor [60] and executive [61] dysfunction.

The decreases of about 3% in putamen and caudate volumes over two years, or annual atrophy rates of 1.5% which we note are greater than the 0.63%/year rate reported even in untreated PHI [30], should be compared with normal age-related loss. Published rates of brain atrophy in healthy individuals are inconsistent [62]. Cross-sectional studies may not

accurately characterize normal change [63, 64], and many longitudinal studies, in addressing brain development or aging, overlook younger adults. Research on older adults typically focuses on global and medial temporal lobe atrophy in the context of aging-related dementia, with consequent neglect of other regions [63]. A 5-year study of healthy individuals (baseline ages 20-77 years) measured respective annual atrophy rates of 0.83% and 0.73% for the caudate and putamen independently of age [65]. In a different, similarly aged cohort, caudate and putamen shrinkage over 2 years was 0.44% and 0.47% (i.e., negligible) [66]. The decreases we observed in putamen and caudate volumes most likely reflect atrophy due to HIV (or to neurotoxic effects of cART), although this remains speculative in the absence of an HIV-uninfected Thai comparison group. Caudate and putamen atrophy was lowest in participants who achieved plasma HIV RNA <50 copies/mL by week 4. The trend association observed between mega-cART and rapidity of viral suppression is supported by previous observations of the SEARCH 010/RV254 cohort [67] and by neurocognitive improvement linked to maraviroc-intensified cART [68]. Taken together, these findings indicate that cART intensification by raltegravir and maraviroc should be further examined for its ability to preserve brain integrity in HIV-infected individuals.

A potential limitation of morphometric MRI studies is the reliability of regional brain volume estimates. Longitudinal investigations of atrophy are facilitated by automated segmentation software but confounded by multiple factors including variability due to data processing techniques. Gunter et al (2003) conducted a careful examination of sources of error in a study of semi-automated methods for computing brain volume loss [69]. Volunteers aged 23–45 years were scanned every 6 weeks over a 5-month period to gauge errors inherent in the measurement process. With atrophy assumed to be imperceptible, image processing accounted for only $\frac{1}{5}$ to $\frac{1}{2}$ of test-retest variability [69]. Other variations reflected MRI acquisition differences (head position, motion, etc.) or real changes (e.g., from hydration effects [70]) [69] which, together with variability due to quantification methods, make "true" rates of atrophy elusive [69]. Recent versions of FreeSurfer are competitive with gold standard manual measurements in cognitively normal and impaired individuals [71], and FreeSurfer was validated for subcortical volumetry in HIV [72]. Importantly, for healthy young volunteers (age 26–31) scanned repeatedly over 31 days (20 sessions; 2 scans per session), FreeSurfer's inter-session volumetric variability did not exceed intra-session variability for the caudate, putamen or any structure except lateral ventricles [73]. (Intra-session variability is presumably unaffected by biological change.) The reliability and consistency of FreeSurfer volume estimates in health and disease [72, 74-79] imparts confidence in our findings, which were confirmed by TBM analyses.

Decreased astrocytic uptake and glutamate metabolism is a major contributing factor in excitotoxic neuronal death linked to cognitive impairment in HIV [80, 81]. We observed relationships between putamen atrophy and lower baseline Cho/Cr and Glx/Cr in frontal GM, which did not, however, overcome correction for multiple comparisons. Our ability to identify such associations may have been constrained by the modest sample size. The analyses of monocyte inflammatory markers also were exploratory and not corrected for multiple comparisons. However, the link between baseline monocyte expression of P-selectin glycoprotein ligand-1, (PSGL-1) and basal ganglia shrinkage suggests that increased trafficking of HIV-infected monocytes into the brain in AHI may play a causative role in

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subsequent atrophy. PSGL-1 promotes leukocyte trafficking across the blood-brain barrier (BBB) during inflammation by tethering the leukocytes to endothelial surfaces of blood vessels [82, 83]. The intermediate/inflammatory CD14+CD16+ monocyte subset, in particular, appears to be involved in putamen atrophy. CD16-expressing monocytes are highly susceptible to HIV infection [84], selectively transmigrate across the BBB [84], and preferentially harbor the virus in cART-treated individuals [85]. New findings indicate that PSGL-1 promotes T-cell immune exhaustion, and it has also become evident that migration-independent functions of PSGL-1 in myeloid cells lead to immune activation and increased MCP-1 secretion through a mammalian target of rapamycin (mTOR) pathway [86]. Collectively these findings provide clues to mechanisms associated with brain atrophy that warrant further investigation.

As CSF S100B and monocyte data were available only for participant subsets, our analyses of brain volumetric change in relation to baseline S100B and monocyte subsets had limited power. The sample size did not allow statistical adjustments for several known modifiers of brain aging, either adverse (e.g., vascular risk factors and stress) or beneficial (educational level, aerobic fitness) [62], which could affect brain volumes in this cohort. Comparison with normal brain alterations was hampered by the lack of an HIV-uninfected Thai comparison group. Nevertheless, likely confounding factors such as substance use and sexually transmitted diseases (STDs) did not appear to influence our results. STD prevalence was <6% and had no effect. The relatively low rate of illicit substance use in the SEARCH010/RV254 cohort [57] distinguishes it from other HIV studies that may have been confounded by drug or alcohol consumption (reviewed by Zahr et al, 2018 [87]). Recreational substance use data were self-reported, available only at M0, and during the period of our study, not systematically collected, which limited their usefulness. However, individuals recruited into the cohort reported no history of substance use judged severe enough to compromise patient care (as evidenced by viral suppression by week 36 in 100% of the sample). Sensitivity analyses confirmed that illicit substance use in the 4 months prior to baseline did not affect atrophy of the caudate or putamen.

A unique strength of the SEARCH010/RV254 study is its well-characterized cohort, a significant proportion of which initiates cART during Fiebig I/II, permitting assessment of the earliest phase of the disease and the impact of immediate treatment. This eliminates what usually is a major source of uncertainty, the duration of infection, and minimizes the occurrence of pre-cART damage, a critical obstacle in determining whether cART is protective.

CONCLUSIONS

Participants enrolled during AHI exhibited progressive brain atrophy two years after infection despite very early initiation of cART. Longitudinal assessment of HIV-uninfected Thai individuals is required to determine whether the volumetric decline is steeper than in healthy controls. Intensification of cART by maraviroc and raltegravir may inhibit brain atrophy by promoting faster viral suppression. Caudate and putamen volume loss is supported by a plausible link to early expansion of PSGL-1-expressing monocytes, particularly the intermediate/inflammatory subpopulation. Therapeutic strategies targeting

viral production in PSGL-1 monocytes may reduce HIV-CNS invasion and eventual brain injury.

ACKNOWLEDGMENTS

We thank our study participants and staff from the Thai Red Cross AIDS Research Centre and the Silom Community Clinic in Bangkok for their contributions, and Ms. Shayanne Martin for her helpful suggestions and assistance with data collection. K.J.K. supervised FreeSurfer MRI data processing, performed statistical analyses and data interpretation, prepared tables and figures, and drafted the manuscript. N.J. conducted TBM data analyses, prepared a figure, and helped to edit the manuscript. N.S. supervised neuroimaging data acquisition and processed the MRS data. K.B. assisted with clinical data collection. P.C. was the on-site neurologist who conducted neurological assessment and clinical follow-up. M.P. conducted the MRI/MRS scanning. N.D. was the radiologist in charge of protocol for MRI/MRS. E.L. conducted cytokine measurements. L.C.N. supervised cytokine measurement and helped to edit the manuscript. K.M.C. performed FreeSurfer processing of MRI data. R.P. contributed to data interpretation and manuscript editing. L.J. performed HIV serology and measured CSF HIV RNA. S.K. measured CSF S100B levels. J.A and S.S. participated in study conception and design, data interpretation, and writing and editing of the manuscript. All authors read and approved the final manuscript.

SEARCH is a research collaboration between the Thai Red Cross AIDS Research Centre (TRCARC), the University of Hawaii and the Department of Retrovirology, U.S. Army Medical Component, Armed Forces Research Institute of Medical Sciences (AFRIMS). This work was supported by NIH grants R01-MH-113560, R01-MH-10414, R21-MH0–86341, K24-MH-098759 and R01-MH-095613; and by the US Military HIV Research Program through its cooperative agreement with the Henry M. Jackson Foundation for the Advancement of Military Medicine, Inc. (W81XWH-07–2-0067, W81XWH-11–2-0174). The views, opinions and/or findings contained in this report are those of the authors and should not be construed as an official Department of the Army position, policy or decision unless so designated by other documentation. In addition, the content of this manuscript is solely the responsibility of the authors and does not necessarily represent the official views of the U.S. Department of Defense, National Institutes of Health, Department of Health and Human Services, or the United States government and the Thai Red Cross AIDS Research Centre.

Funding: NIH grants R01-MH-113560, R01-MH-10414, R21-MH0–86341, K24-MH-098759 and R01-MH-095613; and the cooperative agreement of the US Military HIV Research Program with the Henry M. Jackson Foundation (W81XWH-07–2-0067, W81XWH-11–2-0174).

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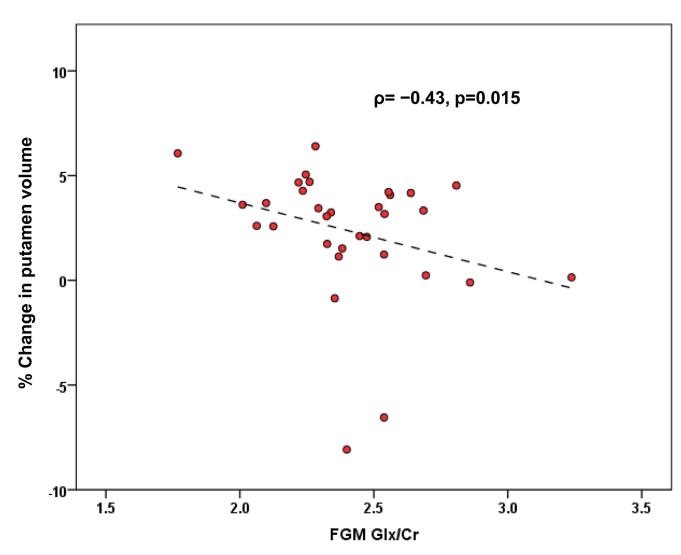


Figure 1.

Putamen volume change at 24 months post-baseline plotted against baseline Glx/Cr in frontal gray matter (FGM). Glx=glutamate (Glu)+glutamine (Gln); percent volume change = (M0-M24)x100/M0, where M0=volume at month 0 and M24=volume at month 24; and ρ is the Spearman correlation coefficient. Putamen volumes were obtained by processing T1-weighted magnetic resonance imaging data with the FreeSurfer longitudinal pipeline.

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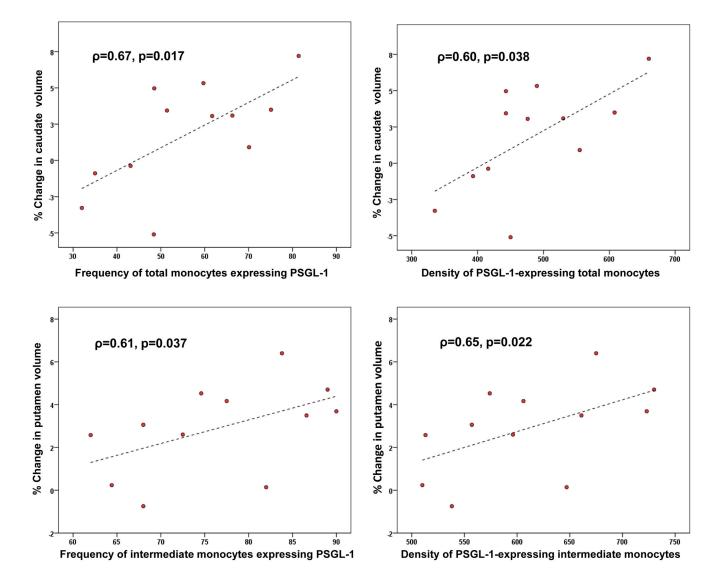
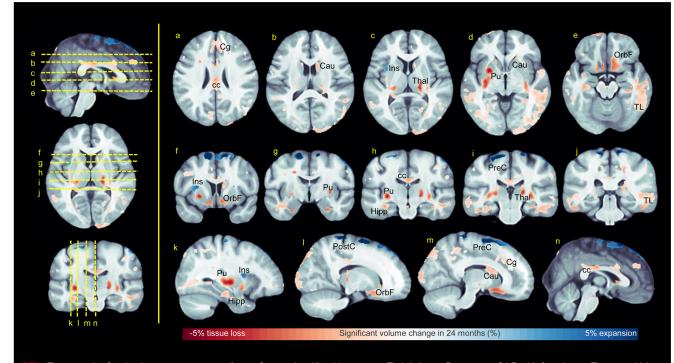


Figure 2.

Caudate and putamen volume changes at 24 months post-baseline plotted against frequencies and densities of circulating intermediate/inflammatory (CD14⁺CD16⁺) and total monocytes expressing P-selectin glycoprotein ligand-1 (PSGL-1). Percent volume change = (M0-M24)x100/M0, where M0=volume at month 0 and M24=volume at month 24; ρ is the Spearman correlation coefficient. T1-weighted magnetic resonance imaging data were processed with the FreeSurfer longitudinal pipeline to obtain regional brain volumes.



RED - Tissue atrophy: Cg: cingulate gyrus; cc: corpus callosum; Cau: caudate; Hipp: hippocampus; Thal: thalamus; Pu: putamen; OrbF: orbitofrontal gyrus; TL: temporal lobe. BLUE - Expansion of cerebral spinal fluid (CSF): PreC/PostC: precentral or postcentral sulcal regions; Ins: insular sulcus.

Figure 3.

Significant changes in regional brain volumes at 24 months post-baseline, identified by using longitudinal tensor-based morphometry (TBM) to process T1-weighted magnetic resonance imaging data.

Table 1.

Characteristics (mean \pm SD, median [minimum-maximum], or N) of study population.

| Demographic/clinical variables | Month 0 (baseline) | Month 24 | p-value |
|---|--------------------------------------|--|--------------------|
| N | 34 | 34 | _ |
| Sex, # male (%) | 29 (85%) | - | - |
| Age (years) | 31.2 ± 8.3 | - | - |
| Education (years) | 16.4 ± 4.0 | - | - |
| Duration of HIV infection (days) | 15 (4–32) | - | - |
| CD4 count (cells/mm ³) | 397.5 (132.0–1127.0) | 644.5 (378.0–1200.0) | < 0.001 |
| Plasma HIV RNA (copies/mL) | 333,971 (604–30,811,000) | 50 (20–58) | < 0.001 |
| CSF HIV RNA (copies/mL) | 1102 (0–201,803) ^A | 0 (0–0) ^B | 0.001 ^B |
| Used illicit substance(s) in past 4 months, # (%) | 10 (32%) [†] | | |
| Co-infections, # (%) | | | |
| Hepatitis C | 1 (3%) | - | - |
| Syphilis | 1 (3%) | - | - |
| Gonorrhea | 1 (3%) | - | - |
| Chlamydia | 0 (0%) | - | - |
| Plasma markers | | | |
| sCD14 (pg/mL) | 1,552,748 (724,106–2,324,882) | 1,106,143 (765,350–1,955,069) ^C | <0.001 C |
| sCD163 (pg/mL) | 99,402 (19,895–377,504) ^C | 85,525 (13,564–380705) ^D | 0.21 C |
| Neopterin (pg/mL) | 2510.8 (227.1-8895.4) | 1399.4 (380.3–6889.9) ^E | 0.001 ^E |
| IL-6 (pg/mL) | 0.43 (0.10–3.14) | 0.25 (0.11–3.42) | 0.68 |
| IL-1β (pg/mL) | 2.03 (0.48–12.59) ^F | 1.50 (0.48–5.83) G | $_{0.014} G$ |
| TNF-a (pg/mL) | 1.46 (0.17–9.10) | 1.05 (0.17–75.82) | 0.17 |
| MCP-1 (pg/mL) | 273.2 (62.2–1781.9) | 352.8 (108.9–1145.1) C | 0.86 C |
| IP-10 (pg/mL) | 2821.8 (33.4 - 13,924.0) | 446.7 (160.9 – 1616.9) C | <0.001 C |
| Cerebrospinal fluid (CSF) markers | | | |
| S100B (pg/mL) | 983.3 (622.3–1795.0) ^H | 1027.2 (807.6 –1590.2) ^I | 0.11 ^J |
| sCD14 (pg/mL) | 45,601 (2057–178,139) G | 58,690 (25,000–1,516,912) ^K | 0.83 ^I |
| CD163 (pg/mL) | 7157.7(3955.1–46,956.1) L | 6281.8 (3253.8–19,266.4) K | 0.055 K |
| Neopterin (pg/mL) | 1817.6 (425.3–5715.5) ^E | 634.6 (469.0–2267.5) ^K | <0.001 L |
| IL-6 (pg/mL) | 2.10 (0.70–20.70) ^H | 1.40 (0.90–8.40) K | 0.07 ^I |
| IL-1β (pg/mL) | $0.55\ (0.500.60)\ ^M$ | $0.50\ (0.500.60)\ ^{N}$ | 0.66 ^N |
| TNF-a (pg/mL) | 0.50 (0.20–7.20) ^H | 0.40 (0.20–3.60) K | 0.011 L |
| MCP-1 (pg/mL) | 903.7 (151.7–4056.8) ^E | 731.2 (386.8–1066.1) ^K | 0.29 ^K |
| IP-10 (pg/mL) | 935.1 (17.7–7006.0) ^E | 552.3 (111.3–2164.0) ^K | 0.004 K |

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P-values were computed by Wilcoxon signed-rank test. S100B = S100 calcium-binding protein B; sCD163 = soluble CD163; IL = interleukin; TNF = tumor necrosis factor; MCP = monocyte chemoattractant protein; IP = interferon- γ -inducible protein 10. Percentages shown are valid percent and exclude missing data.

 † Data were available for 31 participants.

| Data were av |
|--------------------------|
| A _{N=28} |
| B _{N=16} |
| C _{N=30} |
| D _{N=31} |
| E _{N=27} |
| F _{N=24} |
| G _{N=22} |
| H _{N=21} |
| I _{N=14} |
| J _{N=10} |
| <i>K</i> _{N=17} |
| |

L_{N=20}

M_{N=13}

N_{N=11}

AIDS. Author manuscript; available in PMC 2021 March 01.

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Table 2.

Regional brain volumes (mm³, mean \pm SD) at baseline and 24 months.

| Brain region | Month 0 (baseline) | Month 24 | Р |
|-------------------|----------------------------|--------------------------------|---------|
| Caudate nucleus* | 7147.2 ± 867.7 | 7024.5 ± 890.3 | 0.006 |
| Putamen* | 11,364.3 ± 980.4 | $11,102.1 \pm 1031.0$ | < 0.001 |
| Cerebral WM | $434,\!475.2\pm58,\!084.5$ | $432,\!401.3\pm57,\!357.4$ | 0.07 |
| Hippocampus | 9185.4 ± 930.3 | 9219.2 ± 945.0 | 0.08 |
| Cortical GM | $459,\!051.0\pm48,\!940.2$ | $460{,}639{.}4\pm46{,}801{.}2$ | 0.13 |
| Thalamus | $15{,}206.8 \pm 1496.8$ | $15{,}169.2 \pm 1425.9$ | 0.40 |
| Pallidum | 3200.9 ± 430.4 | 3187.8 ± 405.2 | 0.38 |
| Nucleus accumbens | 1149.6 ± 163.0 | 1160.1 ± 275.3 | 0.51 |
| Amygdala | 2985.2 ± 337.5 | 2940.1 ± 443.1 | 0.48 |

P-values were computed by Wilcoxon signed-rank test for related samples.

WM = white matter; GM = gray matter.

* Significant after Holm-Bonferroni correction for multiple comparisons.