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## Oxidative mitochondrial DNA damage in peripheral blood mononuclear cells is associated with reduced volumes of hippocampus and subcortical gray matter in chronically HIV-infected patients

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### Abstract

Cross-sectional relationships were examined between regional brain volumes and mitochondrial DNA (mtDNA) 8-hydroxy-2-deoxyguanosine (8-oxo-dG) in peripheral blood mononuclear cells (PBMCs) of 47 HIV patients [mean age 51 years; 81% with HIV RNA > 50 copies/mL] on combination antiretroviral therapy. The gene-specific DNA damage and repair assay measured mtDNA 8-oxo-dG break frequency. Magnetic resonance imaging was performed at 3 T. Higher mtDNA 8-oxo-dG was associated with lateral ventricular enlargement and with decreased volumes of hippocampus, pallidum, and total subcortical gray matter, suggesting the involvement of systemic mitochondrial-specific oxidative stress in chronic HIV-related structural brain changes and cognitive difficulties. Clarification of the mechanism may provide potential therapeutic targets.

### Keywords

Atrophy; Brain; Neuroinflammation; Oxidative stress; Pallidum; ROS

### 1. Introduction

Entry of the human immunodeficiency virus (HIV) into the brain is believed to occur during acute infection via infected monocytes that migrate across the blood–brain barrier (Gartner,

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2000; Persidsky et al., 2000). HIV antiretroviral drugs suppress plasma viral load and increase CD4 + T-cell count (Collier et al., 1996) but have variable central nervous system (CNS) penetration and effectiveness against infected monocytes (Enting et al., 1998; Letendre et al., 2008). Protected HIV reservoirs are established in brain perivascular macrophages and parenchymal microglia (Kaul et al., 2005) and other CNS cell types (Bagasra et al., 1996). While viral proteins can injure neurons directly (Meucci et al., 1998; Liu et al., 2000), the predominant pathway to neuronal damage and death in HIV/AIDS is probably indirect and mediated by proinflammatory cytokines secreted by HIV-infected or activated macrophages and microglia (Ade-Biassette et al., 1999; Kaul et al., 2001). Immune activation and neuroinflammation often persist even after combination antiretroviral therapy (cART) has reduced plasma HIV RNA to undetectable levels (Anthony et al., 2005; Eden et al., 2007). Many cART-treated patients exhibit clinical features of HIV-associated neurocognitive disorders (HAND) such as psychomotor slowing and impaired verbal and visual memory (Sacktor et al., 2002). Systemic inflammation may also contribute to persistent HAND in this population (Eden et al., 2007).

Studies of Alzheimer's disease (AD) (Griffin et al., 1998; Nunomura et al., 2001), Parkinson's disease (PD) (Sawada et al., 2006; Jenner & Olanow, 1996) and amyotrophic lateral sclerosis (Almer et al., 2002; Ferrante et al., 1997) point to a connection between inflammation and elevated oxidative stress. Oxidative stress is linked to a range of neurodegenerative disorders (reviewed by Andersen (2004)). A mediator of apoptosis and necrosis (Higuchi, 2003), oxidative stress arises from excessive amounts of reactive oxygen species (ROS) which overwhelm cellular antioxidant defenses (Andersen, 2004). In normal and pathological metabolism, most ROS are formed as toxic by-products of oxidative phosphorylation within mitochondria (Turrens, 2003; Lesnefsky et al., 2001; Wallace, 2000) during synthesis of adenosine triphosphate. Mitochondrial DNA (mtDNA) has a high mutation rate compared to that of nuclear DNA (Nachman et al., 1996), possibly due to the proximity of mtDNA to mitochondrial ROS production (Wallace et al., 2010). Emerging data suggest that mitochondrial dysfunction plays a critical role in inflammation. Through ROS production, alterations in mitochondria may amplify cellular responses to cytokine-induced inflammation in osteoarthritis (Vaamonde-Garcia et al., 2012). Mitochondrial dysfunction also increases allergic airway inflammation (Aguilera-Aguirre et al., 2009), which is modulated by mitochondrial ROS production that activates the inflammasome and leads to production of the cytokine IL-1 $\beta$  (Kim et al., 2014). The generation of inflammatory mediators may induce additional mitochondrial dysfunction and oxidative damage, perpetuating a vicious cycle of inflammation.

The majority of metabolic enzymes are downregulated in HIV-infected cells (van't Wout et al., 2003). Neurotoxic HIV proteins that increase ROS production have been implicated in HAND; for example, HIV envelope glycoprotein (gp120) impairs glucose metabolism and neuronal function (Kimes et al., 1991; Vignoli et al., 2000). ROS production is induced by mitochondrial dysfunction in HIV disease (Macho et al., 1995; Gil et al., 2003). Mitochondrial toxicity (Brinkman et al., 1998) and oxidative stress (Lagathu et al., 2007) can be also caused by antiretroviral drugs, especially nucleoside reverse transcriptase inhibitors (NRTIs) which are components of current cART regimens (Hulgan et al., 2003). In particular, use of zidovudine (ZDV) causes mtDNA depletion (Semino-Mora et al., 1994),

shown to impair mitochondrial protein synthesis in a rat model (Masini et al., 1999), and didanosine (ddI) and stavudine (d4T) have been related to mitochondrial damage (Blanco et al., 2003). Oxidative stress can lead to declines in mitochondrial function, which in turn hamper energy production and further elevate ROS levels.

Although neuroimaging of patients on cART reveals HIV-related changes in subcortical and cortical brain structure (Aylward et al., 1993; Thompson et al., 2005; Becker et al., 2011; Kallianpur et al., 2012, 2013), little in vivo research has addressed HIV-related brain structural change in the context of oxidative damage. Neuronal injury detected in the brains of HIV patients (Harezlak et al., 2011; Gray et al., 2000; Ellis et al., 2007) is believed to involve apoptosis and ROS-mediated injury to cellular lipids and proteins (Gray et al., 2000). A widely used marker of mitochondrial oxidative DNA damage is the ROS-induced lesion 8-hydroxy-2-deoxyguanosine (8-oxo-dG) (Valavanidis et al., 2009). Our study investigated the associations of peripheral oxidative stress with regional brain volumes and cognition in chronic HIV infection. Specifically, we examined relationships of mtDNA 8-oxo-dG in peripheral blood mononuclear cells (PBMCs) to regional brain volumes and to neuropsychological (NP) measures.

## 2. Materials and methods

### 2.1. Participants

Neuropsychological and regional brain volume data, obtained in a subset of cART-treated HIV patients from the Hawaii Aging with HIV Cohort-Cardiovascular Disease (HAHC-CVD) study (Shikuma et al., 2012), were examined for cross-sectional associations with mitochondrial data from the parent investigation. The substudy population comprised 47 HIV-seropositive (HIV+) individuals aged 40 years and older. Participants were recruited through referrals from our clinic at the Hawaii Center for AIDS, community physicians, community advisory board members, and AIDS service organizations. Individuals were eligible for the study if they were 40 years old, had a documented history of HIV infection, had been on stable cART 3 months, and if their primary language was English and they could understand and provide informed consent. Exclusion criteria were: (i) uncontrolled major affective disorder; (ii) active psychosis; (iii) loss of consciousness > 5 min; (iv) pregnancy or breast-feeding; (v) factors precluding MRI (e.g., claustrophobia); and (vi) any past or present condition (e.g., central nervous system infection, traumatic brain injury, stroke, substance abuse) deemed by the evaluating physician to introduce confounding variables. All study participants underwent a general and focused HIV/neurological history and physical exam, neuroimaging, and neuropsychological testing. Nadir CD4 count, years since HIV diagnosis, and date of cART initiation were determined by subject self-report. Proximity to HIV diagnosis (i.e., duration of HIV seropositivity) served as a proxy for duration of infection. Past or current use of ZDV, d4T or ddI was noted. Blood specimens were obtained. Plasma HIV RNA and CD4 cell counts were performed by a local commercial Clinical Laboratory Improvement Amendments (CLIA)-certified laboratory. Levels of mtDNA-specific 8-oxo-dG were measured in PBMCs. Each participant provided written informed consent. The University of Hawaii Committee on Human Studies approved the study.

## 2.2. Neuroimaging

MRI was performed on a 3.0-Tesla Philips Medical Systems Achieva scanner equipped with an 8-channel head coil (InVision Imaging, Honolulu). For each subject, a high-resolution anatomical volume was acquired with a sagittal T1-weighted 3D turbo field echo (T1W 3D TFE) sequence (echo time TE/repetition time TR = 3.2 ms/6.9 ms; flip angle 8°; slice thickness 1.2 mm with no gap; in-plane resolution 1.0 mm<sup>2</sup>; field of view 256 × 256 mm<sup>2</sup>). To obtain regional brain volumes, structural MRI data were processed with FreeSurfer (version 5.0, <https://surfer.nmr.mgh.harvard.edu/>) (Dale et al., 1999; Fischl et al., 2002; Fischl et al., 1999, 2004) in a procedure that includes skull-stripping (Segonne et al., 2004), intensity normalization (Sled et al., 1998), Talairach transformation, subcortical white matter and deep gray matter segmentation (Fischl et al., 2002, 2004), and cortical gray/white matter boundary and pial surface reconstruction (Dale et al., 1999). Quality assurance of processed data was performed visually, and cortical surfaces and subcortical segmentations were checked prior to statistical analysis. FreeSurfer's estimate of intracranial volume (ICV) is a reliable measure for regional brain volume normalization (Buckner et al., 2004).

## 2.3. Neuropsychological testing

Subjects completed a comprehensive neuropsychological test battery that assessed cognitive domains affected by HIV. It included the following tests: Choice and Sequential Reaction Time from the California Computerized Assessment Package (CalCAP); the Rey Auditory Verbal Learning Test (RAVLT); Rey-Osterrieth Complex Figure (RCF) Copy and Recall; Trail Making Test, Parts A and B; Wechsler Adult Intelligence Scale–Revised (WAIS-R) Digit Symbol; WAIS-III Letter-Number Sequencing; Grooved Pegboard (dominant and nondominant hands); Verbal Fluency Test (FAS); Animal Naming; Boston Naming Test (BNT), the WAIS-R Digit Span (Forward and Backward); Delis-Kaplan Executive Function System (D-KEFS) Color Word Interference Test, condition 3; and Timed Gait. Raw NP test scores were z-transformed by use of standard demographically adjusted normative data. Composite, domain-specific z-scores were derived by averaging z-scores of the appropriate individual sub-tests (NPZpm [psychomotor speed]: WAIS-R Digit Symbol Subtest; Trail-Making Test, Part B; Grooved Pegboard, non-dominant hand; NPZlrn\_mem [learning and memory]: RAVLT Total; RAVLT Delayed Recall; Rey-Osterrieth Complex Figure Test–Delayed Recall; NPZef [executive functioning]: Verbal Fluency Test (FAS); Stroop Color Word Interference Test; Trail-Making Test, Part B; and NPZwm [working memory]: RAVLT\_B; WAIS-III Letter Number Sequencing). NPZglobal, a global z-score reflecting overall neurocognitive performance, was the arithmetic mean of z-scores on 14 NP tests (WAIS-R Digit Symbol Subtest; Trail-Making Test, Parts A and B; Grooved Pegboard, dominant hand and non-dominant hand; Timed Gait; CalCAP Choice; CalCAP Sequence; RAVLT Total; RAVLT Delayed Recall; Rey-Osterrieth Complex Figure Test–Delayed Recall; Verbal Fluency Test (FAS); Animal Naming; D-KEFS Color Word Interference Test).

## 2.4. Mitochondrial analyses

DNA was isolated using a Qiagen DNA isolation kit (Qiagen, Inc., Valencia, CA). Eight µg of total DNA was digested with Pvu II for one hour at 37 °C. MtDNA 8-oxo-dG was

measured as follows: 4 µg of total DNA was treated with 10 units of human 8-oxo-guanine DNA glycosylase (hOGG1) in 1X REC™ Buffer 6 (20 mM Tris-Cl (pH 8.0), 1 mM DTT, 1 mM EDTA, 0.1 mg/mL BSA) for 1 h at 37 °C in a reaction volume of 20 µL (Trevigen, Inc., Gaithersburg, MD). For analysis, 5 µL of REC 5X Loading Buffer (10 mM HEPES-KOH (pH 7.4)), 10 mM EDTA, 6 M urea, 50% glycerol, and 0.2% bromophenol blue were added. The tubes were heated at 95 °C for 5 min and then fast-cooled to 4 °C. Cleaved and non-cleaved products were resolved on 0.75% alkaline agarose gels. DNA was transferred to nylon membranes using standard Southern Blot methodology. Human mitochondrial probes to cytochrome *b* were labeled with digoxigenin-deoxyuridine triphosphate (Roche Applied Science, Indianapolis, IN) during a linear PCR amplification. Blots were probed and processed for chemiluminescent detection using Roche protocols. Membranes were developed on chemilumiimager (Roche). Quantitation of the break frequency of mtDNA 8-oxo-dG was based on comparing the undigested and digested mtDNA break frequencies calculated using the Poisson distribution, with units reported in break frequency (BF) (Gerschenson et al., 2009).

## 2.5. Statistical analysis

Volumes of 13 brain regions (thalamus, caudate, putamen, globus pallidus, hippocampus, amygdala, nucleus accumbens, cerebellar white matter, cerebellar gray matter, cortical gray matter, cerebral white matter, total subcortical gray matter, and lateral ventricles) were obtained by summing over left and right brain hemispheres. Descriptive statistics were provided for patient demographics, clinical variables, mtDNA 8-oxo-dG, NP z-scores, and regional brain volumes. For ZDV, d4T and ddI (separately), a “history of use” variable was defined to include either prior or current use. Another variable called “history of ZDV, d4T and/or ddI use” reflected past or present use of any of these three NRTIs. MtDNA 8-oxo-dg was treated as a dichotomous categorical variable (zero vs. positive) only for comparisons between subject groups with and without PBMC mtDNA breaks (mtDNA 8-oxo-dG > 0 and mtDNA 8-oxo-dG = 0, respectively). In analyses involving correlation or multivariable linear regression, mtDNA 8-oxo-dG was viewed as a continuous variable. Mann–Whitney or chi-square tests compared differences in demographic and clinical parameters, NP z-scores, and regional brain volumes between groups characterized by the presence or absence of PBMC mtDNA breaks. Relationships of mtDNA 8-oxo-dG levels to continuous demographic and clinical parameters, regional volumes and NP z-scores were also assessed by Pearson correlation (and, when restricted to subjects with mtDNA 8-oxo-dG > 0, by Spearman correlation).

Multiple linear regression examined associations between regional brain volumes (dependent variables) and mtDNA 8-oxo-dG, controlling for age, nadir CD4 cell count and ICV (head size). Low nadir CD4 count has been related to brain atrophy (Cohen et al., 2010) and increased risk of cognitive impairment (Ellis et al., 2011) in HIV-infected individuals on cART. Gender, years of education and years since HIV diagnosis were tested as covariates in the regression but had no significant effects and were not retained. Squares of the semipartial correlations were computed to determine the proportion of variance of each regional volume explained by mtDNA 8-oxo-dG after taking into account the other independent variables. MtDNA 8-oxo-dG data were inspected for outliers and non-normal distribution, the effects

of which were examined in post-hoc analyses. For all regression models, histograms of the standardized residuals were visually checked for normality, and plots of standardized residuals examined to ensure that the residuals were normally distributed around the regression line. Homogeneity of variances was assessed using plots of the standardized residuals against predicted (fitted) values.

SPSS and StatView 5.0 (SAS Institute Inc., Cary, NC) were utilized for statistical analyses. P-values  $<0.05$  were considered statistically significant and  $0.05 < p < 0.1$  indicative of trends. For multivariable regression analyses, all p-values are uncorrected for multiple comparisons, but associations meeting the Bonferroni criterion ( $p < 0.05/13 = 0.004$ ) are noted.

### 3. Results

#### 3.1. Subject characteristics

We studied 47 HIV-infected individuals [mean age:  $50.9 \pm 7.3$  years; predominantly male (87%); mean CD4 cell count =  $462.6 \pm 201.4$  cells/mm<sup>3</sup>; median (interquartile range [IQR]) nadir CD4 count = 140.0 (34.0–251.5) cells/mm<sup>3</sup>; mean duration of HIV infection =  $13.5 \pm 6.8$  years; mean duration of cART =  $11.5 \pm 6.0$  years]. Plasma HIV RNA was undetectable ( $< 50$  copies/mL) in 38 (81%) of the study participants, and the remaining 9 subjects had a median plasma HIV RNA count of 180 copies/mL (range: 53–15,700). Twenty-eight individuals had nonzero (positive) mtDNA 8-oxo-dG levels in PBMCs. Median (IQR) mtDNA 8-oxo-dG values for the entire study sample and for subjects with positive mtDNA 8-oxo-dG were 0.04 (0–0.13) and 0.10 (0.06–0.21) BF, respectively. Distributions of the mtDNA 8-oxo-dG data are shown in Fig. 1. Most demographic and clinical parameters were well balanced between groups defined by the presence or absence of PBMC mtDNA breaks (Table 1), although subjects with mtDNA 8-oxo-dG  $>$  showed trends toward higher current and nadir CD4 counts.

Complete neuropsychological data were obtained for 38 participants, with NP test scores incomplete for 5 subjects and lacking for 4. NP test performance was slightly worse in working memory and learning and memory than in other neurocognitive domains: NPZl<sub>rn</sub>\_mem =  $-0.20 \pm 1.04$  (N = 41); NPZ<sub>wm</sub> =  $-0.18 \pm 0.78$  (N = 43); NPZ<sub>pm</sub> =  $0.28 \pm 0.63$  (N = 42); NPZ<sub>ef</sub> =  $0.16 \pm 0.99$  (N = 43); and NPZ<sub>global</sub> =  $-0.05 \pm 0.50$  (N = 38). Structural MRI yielded volumes of the following brain regions: thalamus ( $13,023.6 \pm 1186.2$  mm<sup>3</sup>), caudate ( $7292.3 \pm 768.3$  mm<sup>3</sup>), putamen ( $11,438.0 \pm 1406.7$  mm<sup>3</sup>), pallidum ( $3315.7 \pm 404.8$  mm<sup>3</sup>), hippocampus ( $7934.8 \pm 804.8$  mm<sup>3</sup>), amygdala ( $3803.2 \pm 495.9$  mm<sup>3</sup>), nucleus accumbens ( $1325.8 \pm 209.4$  mm<sup>3</sup>), cerebellar white matter ( $30,936.5 \pm 3805.4$  mm<sup>3</sup>), cerebellar gray matter ( $87,417.6 \pm 9232.1$  mm<sup>3</sup>), cortical gray matter ( $430,782.3 \pm 35,072.00$  mm<sup>3</sup>), cerebral white matter ( $490,764.5 \pm 55,875.4$  mm<sup>3</sup>), subcortical gray matter ( $164,534.5 \pm 14,469.7$  mm<sup>3</sup>), and lateral ventricles ( $22,798.9 \pm 12,097.5$  mm<sup>3</sup>). Intracranial volume, a covariate in the statistical analyses, was  $1451.2 \pm 204.8$  cm<sup>3</sup>.

### 3.2. Comparison of subject groups with mtDNA 8-oxo-dG = 0 and mtDNA 8-oxo-dG > 0

The groups with and without mtDNA breaks did not differ significantly in demographic or clinical characteristics (Table 1). Compared to subjects with mtDNA 8-oxo-dG = 0, those with mtDNA 8-oxo-dG > 0 had higher current and nadir CD4 count but the differences did not reach significance. NP z-scores did not differ between groups. There were no group differences in history of ZDV use; history of d4T use; history of ddI use; or history of ZDV, d4T and/or ddI use. Current use and past use were also examined separately for each drug, and the group differences (not shown in Table 1) were non-significant. Similarly, Mann–Whitney tests found no significant difference in any regional brain volumes between subjects with mtDNA 8-oxo-dG = 0 and mtDNA 8-oxo-dG > 0. The group with mtDNA breaks showed a trend toward smaller volumes of hippocampus (positive mtDNA 8-oxo-dG:  $7765.5 \pm 816.2 \text{ mm}^3$  vs. zero mtDNA 8-oxo-dG:  $8184.3 \pm 738.6 \text{ mm}^3$ ;  $p = 0.07$ ) and pallidum (positive mtDNA 8-oxo-dG:  $3224.9 \pm 378.2 \text{ mm}^3$  vs. zero mtDNA 8-oxo-dG:  $3449.5 \pm 415.3 \text{ mm}^3$ ;  $p = 0.09$ ). (Both p-values were 0.03 when group differences were assessed by analysis of covariance adjusting for ICV.) Fig. 2 (top) shows the distributions of hippocampal volume for both the zero and positive mtDNA 8-oxo-dG groups.

### 3.3. Correlation of mtDNA 8-oxo-dG with other parameters

Pearson correlation showed no relationship between mtDNA 8-oxo-dG and years of education, years since HIV diagnosis, or years on cART. MtDNA 8-oxo-dG correlated with age ( $R = 0.31$ ,  $p = 0.04$ ), and at trend-level with NPZ lrm\_mem ( $R = 0.30$ ,  $p = 0.06$ ) nadir CD4 ( $R = 0.29$ ,  $p = 0.05$ ) and current CD4 count ( $R = 0.27$ ,  $p = 0.06$ ). Higher mtDNA 8-oxo-dG levels correlated with reduced volumes of hippocampus ( $R = -0.56$ ,  $p = 0.000046$ ), amygdala ( $R = -0.48$ ,  $p = 0.001$ ), pallidum ( $R = -0.32$ ,  $p = 0.027$ ) and total subcortical gray matter ( $R = -0.36$ ,  $p = 0.013$ ). Hippocampal volumes are plotted against mtDNA 8-oxo-dG in Fig. 2 (bottom). When analyses were restricted to subjects with mtDNA 8-oxo-dG > 0, mtDNA 8-oxo-dG still correlated with hippocampal (Spearman  $\rho = -0.55$ ,  $p = 0.003$ ) and amygdalar ( $\rho = -0.52$ ,  $p = 0.004$ ) volumes, and there was a trend correlation between mtDNA 8-oxo-dG and volume of the thalamus ( $\rho = -0.33$ ,  $p = 0.09$ ).

### 3.4. Adjusted associations between mtDNA 8-oxo-dG and regional brain volumes

Higher mtDNA 8-oxo-dG levels correlated with decreased volumes of several brain regions after controlling for age, CD4 nadir count and ICV (Table 2). P-values in the table are uncorrected for multiple comparisons, but volumetric associations of mtDNA 8-oxo-dG with hippocampus, pallidum, total subcortical gray matter and lateral ventricles survive application of a stringent Bonferroni correction ( $p < 0.0039 = .05/13$ ). (The same relationships satisfy the sequentially rejective Holm–Bonferroni criterion (Holm, 1979), often considered more powerful than the classical Bonferroni test.) MtDNA 8-oxo-dG was directly associated with lateral ventricular volume ( $p = 0.0023$ ) and inversely associated with volumes of hippocampus ( $p = 0.0003$ ), pallidum ( $p = 0.0032$ ) and subcortical gray matter ( $p = 0.0013$ ). Relationships between higher mtDNA 8-oxo-dG and decreased volumes of amygdala ( $p = 0.013$ ), putamen ( $p = 0.025$ ) and cerebellar gray matter ( $p = 0.015$ ) did not satisfy multiple comparison correction criteria.

Associations between regional volumes and mtDNA 8-oxo-dG, adjusted for age, ICV and nadir CD4, were examined for dependence on the extreme mtDNA 8-oxo-dG outlier (0.74). The same data point corresponds to the outlier (5349 mm<sup>3</sup>) in the hippocampal volume distribution (Fig. 2, top). With the outlier removed from analyses, brain regions whose volumetric associations with mtDNA 8-oxo-dG had been significant after correction for multiple comparisons remained significantly related to mtDNA 8-oxo-dG: hippocampus ( $\beta = -0.40$ ,  $p = 0.0071$ ), pallidum ( $\beta = -0.35$ ,  $p = 0.015$ ), subcortical gray matter ( $\beta = -0.31$ ,  $p = 0.019$ ), and lateral ventricles ( $\beta = 0.26$ ,  $p = 0.030$ ). Discarding large mtDNA 8-oxo-dG values is somewhat undesirable since the impact of high break frequency is of interest, but the regression coefficients and p-values obtained when the outlier is excluded indicate the robustness of the main results.

Given the skewed distribution of mtDNA 8-oxo-dG (Fig. 1), we tested the effects of transforming these data although predictor variables in regression models need not be normally distributed. (The dependent variables, regional volumes, were approximately Gaussian in distribution.) For the full study population, skewness due to zero values of mtDNA 8-oxo-dG could not be eliminated but was reduced by square root and cube root transformations. The associations between mtDNA 8-oxo-dG and volumes of hippocampus, pallidum, subcortical gray matter, and lateral ventricles (all  $p < 0.01$ ) showed little change in  $\beta$  coefficients. For the subject group with nonzero (positive) values of mtDNA 8-oxo-dG, log-transformation of the data achieved a normal mtDNA 8-oxo-dG distribution. MtDNA 8-oxo-dG was negatively associated with hippocampal volume ( $\beta = -0.38$ ,  $p = 0.021$ ), controlling for age and ICV (nadir CD4 was dropped as it had a non-significant effect and because of low sample size). Square root and cube root transformations of mtDNA 8-oxo-dG produced results roughly similar to those found for the log-transformed data. (Regression coefficients cannot be directly compared when using different transformations of the data.)

#### 4. Discussion

ROS-mediated mtDNA injury, mitochondrial dysfunction and oxidative stress play important roles in neurodegenerative processes. MtDNA is 10 to 100 times more sensitive than nuclear DNA to oxidative stress (Yakes & Van Houten, 1997) and thus highly susceptible to oxidative damage. In a cycle central to the free radical theory of aging, elevated mitochondrial ROS levels lead to mtDNA damage which then causes further mitochondrial and respiratory chain deficiency and accumulation of ROS (Harman, 1956, 1972). HIV-associated mitochondrial dysfunction results in increases in ROS production and indices of overall oxidative stress (Macho et al., 1995; Gil et al., 2003). Peripheral macrophages are related to activated microglia (Davis et al., 1994) which are largely responsible for generation of free radicals in the CNS (Klegeris & McGeer, 2000). Neurotoxins released by the activated microglia trigger inflammatory processes linked to neurodegeneration. Oxidative stress-mediated neuroinflammation is a probable contributor to neuronal loss in HIV (reviewed by Gray et al. (2000)); pronounced neuroinflammation occurs even in cART-treated, cognitively normal HIV patients (Anthony et al., 2005). Our study identified a link between brain structural change and elevated peripheral oxidative stress in well-controlled HIV disease. Volumes of multiple brain regions correlated negatively with mitochondrial DNA 8-oxo-dG in PBMCs of HIV-infected individuals after



adjustment for age, intracranial volume and nadir CD4 count. These findings were substantiated by an association between increased mtDNA 8-oxo-dG and lateral ventricular dilatation indicating central brain atrophy. After correction for multiple comparisons, higher mtDNA 8-oxo-dG levels were associated with enlargement of the lateral ventricles and with smaller volumes of hippocampus, pallidum, and subcortical gray matter in regression models with moderate R (Persidsky et al., 2000). The amygdala, putamen, and cerebellar gray matter were also volumetrically associated with PBMC mtDNA 8-oxo-dG though the p-values did not pass the strict Bonferroni criterion.

Our most statistically significant result was the relationship between increased PBMC mtDNA 8-oxo-dG break frequency and reduced hippocampal volume. HIV proteins damage the hippocampus via oxidative stress: for example, hippocampal neurons are susceptible to transregulatory protein (Tat)-induced injury and apoptosis (Maragos et al., 2003) which occurs through mitochondrial dysfunction (Kruman et al., 1998). The hippocampus is a site of ongoing neuroinflammation even in HIV patients on suppressive cART. A postmortem study by Anthony et al. (Anthony et al., 2005) examined microglial and macrophage activation in the basal ganglia and hippocampus of HIV-infected, cognitively normal individuals whose plasma viral load had been suppressed by cART to below 50 copies/mL. Hippocampal neuroinflammation in these subjects exceeded that seen in HIV-encephalitis patients during the pre-cART era. It was concluded that the site of microglial/macrophage activation in patients with chronic suppressed HIV infection may be shifting to the hippocampus, perhaps because of therapy itself (Anthony et al., 2005). The basal ganglia, too, showed a pre- and post-cART difference in neuroinflammation (Anthony et al., 2005) that is consistent with the associations we observed between mtDNA break frequency and volumes of pallidum, putamen, and total subcortical gray matter.

The hippocampus is vulnerable to oxidative stress in general. In a rat model of neuroinflammation in AD, increased microglial activation and working memory deficits (Hausse-Wegrzyniak et al., 1998, 1999a, 1999b) as well as striking hippocampal atrophy (Hausse-Wegrzyniak et al., 2000) were produced by infusion of lipopolysaccharide into the ventricular system. Severity of neuronal damage in rat hippocampal slices correlates with ROS levels (Schurr & Gozal, 2011). Prominent in the human hippocampus are fast neuronal network oscillations in the gamma frequency range (~ 30–90 Hz), which involve high oxygen consumption requiring maximal mitochondrial oxidative capacity (Kann et al., 2011). These gamma oscillations provide a structure for information processing and contribute to cognitive memory formation and sensory processing (reviewed by Bartos et al. (2007)). The role played by hippocampal gamma oscillations in encoding and retrieving working memories (Montgomery & Buzsaki, 2007; van Vugt et al., 2010), and the well-established involvement of the hippocampus in learning and memory (O'Reilly & Rudy, 2001), support our study findings since the participants had lower test scores both in working memory and in learning and memory. Because the amygdala was also volumetrically related to mtDNA 8-oxo-dG, the limbic system may be particularly susceptible to ROS-induced oxidative stress. This conjecture is strengthened by the fact that the relationships between mtDNA oxidative damage and volumes of hippocampus and amygdala appeared to be driven by the subject group with positive mtDNA 8-oxo-dG values. Moreover, the associations we found between mtDNA 8-oxo-dG and volumes of

hippocampus and cerebellar gray matter are interesting in light of published work on the effect of oxidative stress on brain neurons. Hippocampal CA1 neurons are selectively vulnerable to oxidative stress (Wang et al., 2005; Wilde et al., 1997). Neurons in cerebellar gray matter (but not cerebral cortex) show a differential response to conditions involving oxidative stress: in particular, the cerebellar granule layer is sensitive to oxidative stress and can undergo extensive neuronal death (Wang et al., 2009).

The trend associations of mtDNA 8-oxo-dG with higher current and nadir CD4 counts may be attributable to earlier initiation of therapy in subjects with positive mtDNA 8-oxo-dG, although duration of cART did not correlate with mtDNA 8-oxo-dG or differ significantly between groups with and without PBMC mtDNA oxidative damage. A greater proportion of subjects with mtDNA 8-oxo-dG > 0 had a history of ZDV, d4T and/or ddI use. Antiretroviral drugs may cause accumulation of ROS and induce neuronal injury and death in vitro (Akay et al., 2014), and were proven centrally neurotoxic in vivo in two animal models (Akay et al., 2014). HIV patients treated with NRTIs have reduced mtDNA content in PBMCs (Chiappini et al., 2004). However, PBMCs of HIV-infected individuals naïve to antiretroviral therapy also exhibit mitochondrial dysfunction involving decreased oxidative phosphorylation activity (Miro et al., 2004) and mtDNA levels (Chiappini et al., 2004; Miro et al., 2004; Casula et al., 2005; Miura et al., 2003), suggesting that HIV infection itself contributes to mitochondrial toxicity (Maagaard & Kvale, 2009).

Our study limitations included the cross-sectional design, modest sample size and lack of HIV-negative subjects. Additionally, the participants were aged 40 years and older. A larger, longitudinal study that includes younger individuals is warranted to confirm and clarify the impact of HIV-induced oxidative stress on the brain.

## 5. Conclusion

Peripheral mitochondrial DNA oxidative damage was associated with reduced hippocampal and subcortical gray matter volumes in chronic stable HIV disease. To our knowledge this is the first reported connection between oxidative stress in the peripheral compartment and brain structure in HIV infection. Our data corroborate the hypothesized involvement of oxidative stress in HIV-related brain neuronal loss in the era of effective cART, and suggest that PBMC mtDNA 8-oxo-dG should be investigated as a predictor of brain atrophy in individuals with mild symptoms of HAND. Regional brain volumetric loss may be moderated in part by mitochondrial dysfunction or by a common HIV-related process in the midst of viral suppression that affects both mtDNA and the brain. Clarification of the mechanism may eventually lead to treatments that inhibit neurodegeneration by decreasing oxidative stress and restoring mitochondrial function in this vulnerable population.

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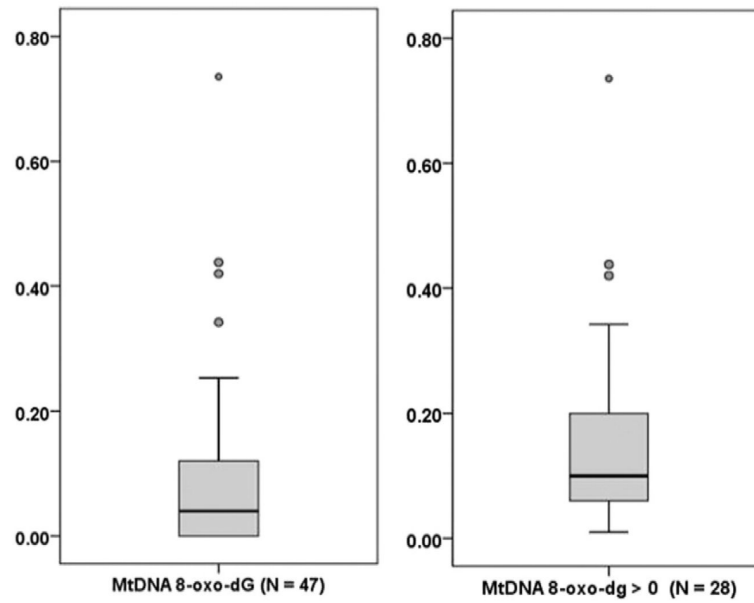
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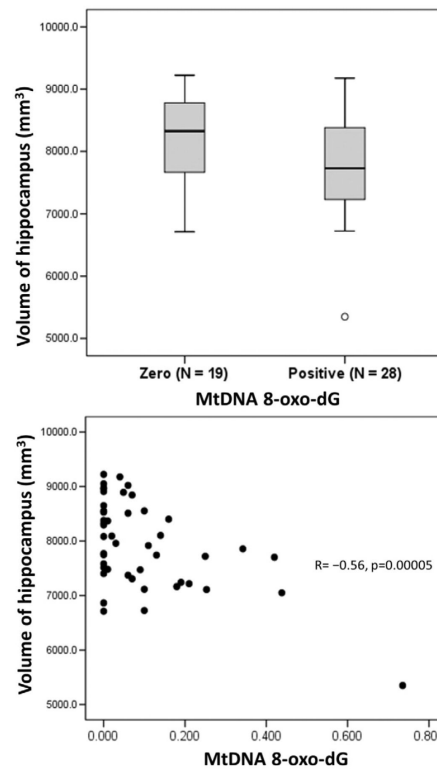
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**Fig. 1.**

Data distributions of peripheral blood mononuclear cell mitochondrial DNA (mtDNA) 8-hydroxy-2-deoxyguanosine (8-oxo-dG) (break frequency). Left: full distribution; right: distribution of positive values. For each boxplot, the middle dark line shows the median; the shaded box indicates the 1st (Q1) and 3rd (Q3) quartiles. Whiskers extend the box range to the extreme data points or, if there are outliers, by  $1.5 \times (Q3 - Q1)$  in either direction; circles are outliers ( $< Q1 - 1.5 \times [Q3 - Q1]$  or  $> Q3 + 1.5 \times [Q3 - Q1]$ ).





**Fig. 2.** Top: boxplots showing total volumes of hippocampus for subject groups with zero and positive values of mitochondrial DNA (mtDNA) 8-hydroxy-2-deoxyguanosine (8-oxo-dG) in peripheral blood mononuclear cells;  $p = 0.07$  by Mann–Whitney test. Bottom: total volumes of hippocampus plotted against mtDNA 8-oxo-dG for entire study population.

**Table 1**

Study population: demographics, clinical variables, and mitochondrial DNA 8-oxo-dG in peripheral blood mononuclear cells. Data are given as mean  $\pm$  standard deviation,  $n$  (%) or median (interquartile range [IQR]). P-values for group differences were obtained by chi-squared or Mann–Whitney test.

Variable	All subjects	MtDNA 8-oxo-dG = 0	MtDNA 8-oxo-dG > 0	p
N	47	19	2	8–
Age (years)	50.9 $\pm$ 7.3	49.4 $\pm$ 6.7	51.9 $\pm$ 7.6	0.24
Male gender	41 (87.2%)	17 (90%)	24 (86%)	0.99
Education (years) <sup>a</sup>	14.4 $\pm$ 2.1	14.2 $\pm$ 2.1	14.6 $\pm$ 2.1	0.58
Race/ethnicity (Caucasian)	24 (51%)	10 (53%)	14 (50%)	0.99
Duration of HIV infection (years)	13.5 $\pm$ 6.8	11.7 $\pm$ 6.9	14.8 $\pm$ 6.5	0.12
Duration of cART (years)	11.5 $\pm$ 6.0	10.0 $\pm$ 6.1	12.4 $\pm$ 5.9	0.18
CD4 count (cells/mm <sup>3</sup> )	462.6 $\pm$ 201.4	393.3 $\pm$ 205.6	509.6 $\pm$ 187.7	0.05
Nadir CD4 count (cells/mm <sup>3</sup> ) <sup>b</sup>	140.0 (34.0–251.5)	67.0 (10.0–230.0)	183.5 (76.3–269.8)	0.07
Undetectable plasma HIV RNA (<50 copies/mL)	38 (81%)	14 (74%)	24 (86%)	0.45
MtDNA 8-oxo-dG (BF)	0.04 (0–0.13)	0 (0–0)	0.10 (0.06–0.21)	–
History of ZDV use	25 (53%)	8 (42%)	17 (61%)	0.21
History of d4T use	18 (38%)	6 (32%)	12 (43%)	0.44
History of ddI use	10 (21%)	3 (16%)	7 (25%)	0.45
History of ZDV, d4T and/or ddI use	31 (66%)	10 (53%)	21 (75%)	0.11

<sup>a</sup>N = 44.

<sup>b</sup>N = 45.

**Table 2**

Associations between regional brain volumes and mitochondrial 8-oxo-dG by multiple regression.

Brain region (volume)	Predictor variables	$\beta$	p	Fraction of variance due to mtDNA 8-oxo-dG
Hippocampus	Age	-0.14	0.273	0.21
	CD4 nadir	0.10	0.412	
	ICV	0.36	0.0040	
	MtDNA 8-oxo-dG	-0.52	<b>0.0003*</b>	
Amygdala	Age	-0.28	0.045	0.10
	CD4 nadir	0.07	0.591	
	ICV	0.34	0.0086	
	MtDNA 8-oxo-dG	-0.36	0.013	
<b>Pallidum</b>	Age	-0.16	0.256	0.15
	CD4 nadir	0.34	0.013	
	ICV	0.35	0.0077	
	MtDNA 8-oxo-dG	-0.44	<b>0.0032*</b>	
Thalamus	Age	-0.23	0.092	0.04
	CD4 nadir	-0.02	0.853	
	ICV	0.50	0.0002	
	MtDNA 8-oxo-dG	-0.23	0.096	
Caudate	Age	-0.26	0.082	0.02
	CD4 nadir	0.11	0.415	
	ICV	0.46	0.0014	
	MtDNA 8-oxo-dG	-0.17	0.248	
Putamen	Age	-0.28	0.040	0.08
	CD4 nadir	0.29	0.027	
	ICV	0.42	0.0014	
	MtDNA 8-oxo-dG	-0.32	0.025	
Nucleus accumbens	Age	-0.25	0.105	0.05
	CD4 nadir	0.12	0.397	
	ICV	0.31	0.032	
	MtDNA 8-oxo-dG	-0.25	0.112	
Cerebellar WM	Age	-0.18	0.230	
	CD4 nadir	-0.08	0.585	
	ICV	0.32	0.024	

Brain region (volume)	Predictor variables	$\beta$	p	Fraction of variance due to mtDNA 8-oxo-dG
	MtDNA 8-oxo-dG	-0.27	0.089	
				0.06
Cerebellar GM	Age	-0.06	0.688	
	CD4 nadir	0.25	0.079	
	ICV	0.43	0.0023	
	MtDNA 8-oxo-dG	-0.38	0.015	
				0.11
Cortical GM	Age	-0.14	0.199	
	CD4 nadir	-0.01	0.934	
	ICV	0.72	<0.0001	
	MtDNA 8-oxo-dG	-0.16	0.183	
				0.02
Cerebral WM	Age	-0.14	0.189	
	CD4 nadir	0.01	0.950	
	ICV	0.72	<0.0001	
	MtDNA 8-oxo-dG	-0.21	0.064	
				0.03
Subcortical GM	Age	-0.13	0.296	
	CD4 nadir	0.25	0.039	
	ICV	0.55	<0.0001	
	MtDNA 8-oxo-dG	-0.43	<b>0.0013*</b>	
				0.14
Lateral ventricles	Age	0.33	0.006	
	CD4 nadir	-0.11	0.307	
	ICV	0.49	<0.0001	
	MtDNA 8-oxo-dG	0.38	<b>0.0023*</b>	
				0.11

\* Significant when Bonferroni-corrected for multiple comparisons ( $p < 0.004$ ).