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# Plasma Biomarker Factors Associated with Neurodevelopmental Outcomes in Children with Perinatal HIV Infection and Controlled Viremia

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# Abstract

**Objective:** We examined relationships between plasma biomarkers and neurodevelopment in children from Sub-Saharan Africa with perinatal HIV (PHIV) with controlled viremia on antiretroviral therapy (ART).

**Design:** Longitudinal retrospective cohort study of children with controlled blood HIV replication.

**Methods:** Children (N=213; 57% female) started ART at <3 years old, had neurodevelopmental assessments (cognition, attention/impulsivity, motor proficiency, global executive functions) at 5–11 years, and achieved controlled viremia (HIV-1 RNA <400 copies/mL for 9 months before initial assessment). Twenty-three plasma biomarkers were measured at onset of controlled viremia, week 0 (first neurodevelopmental assessment), and week 48 (second neurodevelopmental assessment). Factor analysis was conducted at each time point. Multivariable linear regressions assessed associations between factors and neurodevelopmental scores.

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**Results:** Median age at week 0 was 7.0 years. Nineteen biomarkers loaded on 6 factors: A (L-10, IFN $\gamma$ , IFN $\alpha$ 2, IL-1 $\beta$ , IL-6, IP-10, TNF $\alpha$ ); B (sCD163, sICAM-1, sVCAM-1, CRP); C (sE-selectin, sP-selectin); D (MIP-1 $\beta$ , VEGF-A); E (sCD14, CRP); and F (CX3CL1, MCP-1). Higher Factor B scores were consistently associated with worse cognition and attention/impulsivity, and higher Factor D scores with better attention/impulsivity.

**Conclusions:** These results suggest a detrimental effect of increased endothelial cell activation (sICAM-1, sVCAM-1) and monocyte/macrophage scavenger function (sCD163) and a beneficial effect of increased CCR5 ligand and HIV entry blocker MIP-1 $\beta$  and angiogenesis stimulant VEGF concentrations on the neurodevelopment of children with PHIV. The model that emerges is of vascular inflammation leading to neurodevelopmental deficits. The role of persistent HIV replication in the central nervous system also needs to be further explored.

# INTRODUCTION

Children with perinatal HIV infection (PHIV) have poorer neurodevelopmental (ND) outcomes compared to their uninfected peers, despite initiation of antiretroviral therapy (ART) in early childhood and sustained virologic control.[1] Soluble biomarkers of monocyte/microglial and endothelial activation may have clinical utility in predicting ND outcomes in children with PHIV.[2–5] Expression of CD163, a marker of monocyte/ macrophage scavenger activity, has been associated with developmental delays in a Kenyan cohort of children with PHIV and suppressed viremia who had initiated ART in infancy,[2] but not in a cohort of late-diagnosed and treated school-aged children in Thailand and Cambodia[3], where elevated subsets of activated monocytes following ART initiation were associated with higher neurocognitive scores.[3] Biomarkers of pro-coagulant state (fibrinogen and P-selectin) have been individually associated with adverse ND outcomes in youth with PHIV.[4]

While these studies suggests associations between biomarkers and ND outcomes in children, they are limited due to selective and restrictive panels of biomarkers, analyses at a single time point, research participants in different stages of HIV disease, and variations in treatment status and level of virologic control. Furthermore, most published studies fail to capture the known redundancy of inflammatory and immune responses, such that different individuals may respond to illness or injury with different cytokines or chemokines within the same family.[2–4, 6–7]

One way to account for this complexity is to examine a wide range of candidate biomarkers utilizing a statistical approach that captures potential aggregate effects of subsets of biomarkers. Using this approach, we have previously identified a latent factor based on fibrinogen, CRP, and IL-6 that was negatively associated with a measure of processing speed in a cross-sectional US-based study of youth with virologically suppressed PHIV.[5] The generalizability of this finding was limited, however, due to the small number of biomarkers (n=9), lack of longitudinal data, and assessment of ND outcomes that used only one test (WISC-IV).

In the present hypothesis-generating study, we explore possible pathophysiological pathways and their cross-sectional and longitudinal association with four ND outcomes. The study

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included assays of 23 biomarkers relevant to monocyte/microglial and endothelial activation and was restricted to children with virologically controlled HIV infection to minimize potential confounding effects of uncontrolled viremia. This manuscript summarizes the identified groupings of biomarkers and their longitudinal association with ND outcomes.

# **METHODS**

#### Study Sample, Eligibility Criteria and Time Points of Interest

This analysis used data from two studies: P1060[8] (NCT00307151) and P1104s[9] (NCT02140255). P1060 was a randomized controlled trial comparing nevirapine versus lopinavir/ritonavir-based ART initiated in children with PHIV at age 2 months to 3 years at six study sites in South Africa, Malawi, Uganda and Zimbabwe.[8] P1104s, a sub-study of P1060, assessed the feasibility, reliability, and validity of administering a ND assessment battery to children aged 5–11 years. All children with PHIV in P1104s participated in P1060.[9–10]

The analysis focused on P1104s children with PHIV who had controlled viremia, defined as HIV-1 plasma RNA (viral load; VL) <400 copies/mL, for 9 months prior to P1104s entry. VLs were assessed every 6 months. Eligible children had at least one VL <400 copies/mL between 6–12 months prior to P1104s entry and no VL 1000 copies/mL, and no more than one VL of 400–999, for the 9 months prior to P1104s entry. Children were excluded if ND assessment results were not available at both P1104s entry (+/–6 weeks) and week 48 visits (+/–12 weeks). A VL cutoff of 400 copies/mL was chosen as this was the lower limit of detection for the assay in 2005, when P1060 opened for enrollment.

There were three time points of interest: time of onset of controlled viremia prior to P1104s entry, P1104s entry (week 0), and 48 weeks after P1104s entry (week 48). Start of controlled viremia was defined as the date of the earliest VL <400 copies/mL with no subsequent consecutive measurements 400 copies/mL and no 12-month periods with more than one VL 1000 copies/mL prior to P1104s entry. Participants with subsequent uncontrolled viremia (consecutive VL 400 copies/mL or more than one VL 1000 copies/mL) between P1104s entry and their week 48 visit were included in week 0 analyses but excluded from week 48 analyses.

#### **Biomarkers**

Plasma samples were collected and stored during P1060. The cryopreserved samples closest to and within six months of each of the time-points of interest were identified, shipped and tested in a central laboratory. Plasma levels of 23 biomarkers were assayed, including acute phase reactants (CRP), adhesion factors (ICAM-1, ICAM-5, VCAM-1), anti-inflammatory cytokines (IL-10), chemoattractants (CX3CL1, MCP-1, MIP-1 $\beta$ ), inflammatory cytokines (IFN $\gamma$ , IFN $\alpha$ 2, IL-1 $\beta$ , IL-6, IP-10, TNF $\alpha$ ), and markers of endothelial activation (sEselectin, endothelin-1, VEGF-A), matrix digestion (MMP9), monocyte activation (CD14, CD163), neuronal toxicity (neurofilament-light [NFL]), and pro-coagulant state (fibrinogen, P-selectin). Analytes were measured as per manufacturers' instructions using the following commercial kits: MMP-9, IFN $\alpha$ 2, IFN $\gamma$ , IL-10, IL-1 $\beta$ , IL-6, IP-10, MCP-1, MIP-1 $\beta$ , TNF $\alpha$ , VEGF-A, CRP (lower limit of detection = 1.33 pg/ml), sICAM-1, sVCAM-1, sE-selectin, sP-selectin (Mesoscale Discovery, chemiluminescence microarrays); fibrinogen, CX3CL1 (Millipore; ELISA); sCD14, sCD163, endothelin-1, sICAM-5, NFL (R&D, ELISA).

#### **Neurodevelopmental Outcomes**

The ND measures were previously adapted for pediatric HIV research in Africa, validated for school-age children in Sub-Saharan Africa, and described elsewhere.[1,9–10] Briefly, the Kaufman Assessment Battery for Children, 2nd edition (KABC-II) Mental Processing Index is a standardized composite (mean=100, SD=15) of four cognitive domains (sequential and simultaneous processing, learning ability, and planning ability); higher values indicate better performance. The Tests of Variables of Attention (TOVA) D-prime is an unstandardized signal detection measure assessing attention and impulsivity based on correct responses to signal and withholding of responses to non-signal in proportion to incorrect responses; TOVA D-prime can range from 0 to 10 with higher values indicating better performance. [11-12] The Bruininks-Oseretsky Test of Motor Proficiency, 2nd edition (BOT-2) is a standardized measure (mean=50, SD=10) assessing fine-motor precision, fine-motor integration, manual dexterity, bilateral coordination, balance, running speed and agility, upper-limb coordination and strength; higher values indicate better performance. The Behavior Rating Inventory of Executive Function (BRIEF), Parent Form, Global Executive Composite (GEC) is a standardized measure of parental perceptions of their child's executive functioning (mean=50, SD=10); higher values indicate poorer performance. For this analysis, the BRIEF GEC was subtracted from 100 (reverse scored) so that higher values indicated better performance. Consistent with previous research, [10] the performance on the KABC-II, BOT-2, and BRIEF was age-standardized using American norms, while TOVA Dprime outcomes were adjusted for age and sex.[9–10]

#### **Statistical Methods**

ND measures and biomarker levels at each time point and changes between time points were summarized. Changes greater than zero indicated that the measurement at the later time point was larger than the measurement at the earlier time point. Paired t-tests were used to assess changes in ND measurements between week 0 and week 48.

Biomarker results were  $log_{10}$ -transformed and standardized (mean=0, SD=1) to put all measures on the same scale prior to conducting factor analyses. Factor loadings were calculated using varimax rotations. The number of factors was selected after evaluating scree plots and cumulative variance plots.

After reviewing the resulting seven factors at each time point, we observed six groupings of biomarkers that remained essentially unchanged across timepoints. Therefore, instead of analyzing seven different factors at each of the three timepoints, we simplified the approach post hoc by focusing our exploratory analysis on these six groupings of biomarkers, which we refer to as "consensus factors", defined as groupings of biomarkers with loadings >0.4 on the same factor at 2 time points. Typically, factor loadings above 0.40 are considered meaningful loadings.[13] The consensus factors were labeled "A" through "F".

For each consensus factor, we checked for consistency in directionality of the biomarkers. If a factor loading was negative, the corresponding biomarker value was multiplied by -1 to convert it to the same numerical standard. Consensus factor scores for a given participant at each time point were calculated by summing the standardized values of that participant's biomarker measurements in each grouping and dividing by the number of biomarkers in the grouping.

Multivariable linear regression models were fit to assess associations between the consensus factor scores and ND outcomes. For each consensus factor, separate models were fit using each of the four ND outcomes at week 0, week 48 and the change in ND outcomes between weeks 0 and 48. Each model was adjusted for study site, sex, age at P1104s entry, age at ART initiation, ART regimen at time of biomarker specimen collection (LPV/r-based, NVPbased, or other), and whether the participant switched regimens between P1060 entry and biomarker specimen collection (a proxy for virologic failure). For models involving the change between weeks 0 and 48, we also adjusted for consensus factor scores at week 0. We report regression coefficient estimates and 95% confidence intervals, highlighting estimates whose 95% confidence interval did not contain zero. We used a complete case analysis approach to handle data. Participants that were missing any biomarker measurement at a given time point were excluded from the factor analyses and regression models corresponding to that time point. Given the exploratory nature of this study, no adjustments were made for multiple comparisons.[14–16] Statistical analyses were performed using SAS version 9.4 (SAS Institute, Cary, NC, USA, 2013) and graphics generated in R version 3.2.2 (http://www.R-project.org).

# RESULTS

## **Participant Characteristics**

Of 246 children with PHIV who were enrolled in both P1060 and P1104s, 213 were included in this analysis (Supplemental Figure 1). Excluded children either had uncontrolled viremia in the 9 months prior to P1104s entry (n=25) or did not have biomarker results at any of the key time points (n=8). The remaining 213 children had at least one ND result at both weeks 0 and 48; two children were removed from week 48 analyses due to loss of virologic control following P1104s entry. Table 1 shows demographic and clinical characteristics of the cohort. Supplemental Table 1 shows ART treatment history across multiple time points.

#### ND Results over Time

Mean ND scores at week 0 were: 73.1 for KABC-II (95%CI: 71.7, 74.5), 48.4 for BOT-2 (95%CI: 47.2, 49.6), 2.4 for TOVA D-prime (95%CI: 2.3, 2.6), and 47.2 for BRIEF (95%CI: 45.4, 49.0). There were statistically significant changes between weeks 0 and 48 in three of four ND measures: BOT-2 (p = 0.016), TOVA D-prime (p < 0.001), and BRIEF (p < 0.001). Between weeks 0 and 48, mean [95% CI] scores increased 1.0 for the BOT-2 (0.2, 1.9), 0.6 for TOVA D-prime (0.5, 0.8), and 5.2 for the BRIEF (3.5, 6.8) and decreased 0.3 (-1.4, 0.8) for the KABC-II (Figure 1).

#### **Biomarkers over Time**

Boxplots for the 23 biomarkers at the three time points are provided in Figure 2. Ninety-two percent of sICAM-5 values were below the lower limit of detection (LLOD); all other biomarkers had less than 8% of values above or below the corresponding upper or lower detection limits.

#### **Biomarker Reduction using Factor Analysis**

Two biomarkers (sICAM-5 and endothelin-1) were not included in the factor analyses because of high rates of values below LLOD (92%) or missing observations (30%), respectively, leaving 21 biomarkers for the factor analysis. Additionally, missing data excluded 15 of 184 (8%) children from factor analyses at the start of controlled viremia, 6 of 185 (3%) at week 0, and 8 of 91 (8%) at week 48.

Eigenvalues and scree plots from each factor analysis suggested six or seven factors at each time point. The total variance explained by seven factors was between 68% and 74%. Factor loadings for each biomarker at each time point are shown in Table 2.

The six identified consensus factors were Factor A (IL-10, IFN $\gamma$ , IFN $\alpha$ 2, IL-1 $\beta$ , IL-6, IP-10, and TNF $\alpha$ ), Factor B (sCD163, sICAM-1, sVCAM-1, and CRP), Factor C (sE-selectin and sP-selectin), Factor D (MIP-1 $\beta$  and VEGF-A); Factor E (CRP and sCD14), and Factor F (CX3CL1 and MCP-1). Two biomarkers (MMP9 and NFL) did not meet the post-hoc criteria and were not assigned to any of the six consensus factors. Supplemental Table 2 presents the six consensus factors and the 19 corresponding biomarkers. Supplemental Table 3 shows the factor loadings at each time point, grouped according to consensus factors.

#### Association between Biomarker Factors and ND Outcomes

Table 3 presents the estimated association between the consensus factor scores at Week 0 and the four ND outcomes at week 0 and week 48 and the change in ND outcomes between weeks 0 and 48. At week 0, higher Factor B scores were associated with lower (poorer) KABC-II scores at weeks 0 and 48 and poorer TOVA D-prime scores at weeks 0 and 48. Higher Factor D scores at week 0 were associated with higher (better) TOVA D-prime scores at week 48 and a greater increase in TOVA D-prime scores between weeks 0 and 48. Participants with higher Factor E scores at week 0 had a smaller increase, on average, in KABC-II scores between weeks 0 and 48. Factors A, C, and F at Week 0 were not associated with any of the ND outcomes. None of the six consensus factors at week 0 was associated with the BOT-2 or BRIEF at either Week 0 or Week 48.

Associations between consensus factor scores at the start of controlled viremia and at week 48 are provided in Supplemental Tables 4 and 5, respectively. Higher Factor D scores assessed at the start of controlled viremia were associated with higher TOVA D-prime values at Week 0; consensus factor scores at the start of controlled viremia were not associated with BOT-2, BRIEF, or KABC-II outcomes. At week 48, higher Factor B scores were associated with poorer KABC-II scores, and higher Factor C scores were associated with poorer BOT-2 scores.

# DISCUSSION

We investigated cross-sectional and longitudinal associations between biomarkers reflective of inflammation and immune activation and ND outcomes in a cohort of children with PHIV with controlled viremia. Notably, almost all of the noteworthy associations between biomarker consensus factors and ND outcomes involved consensus factors subsequent to a period of sustained virologic control (i.e., at weeks 0 and 48), rather than at the onset of virologic control. It is possible that the impact of specific inflammatory cascades is not easily discernable proximate to a period of uncontrolled viral replication and may only become apparent after a period of sustained control of viral replication; alternatively, an insult to the CNS that occurred during uncontrolled viremia may only manifest later, despite virological control. One exception, the association between Factor D at the onset of controlled viremia and TOVA D-prime at Week 0, suggests the relatively early prognostic value of this factor. The hypothesis that emerges is that, in addition to suppressing viral replication, modulating inflammation and immune activation may improve ND outcomes.

The consensus factor most consistently associated with ND outcomes was Factor B (sICAM-1, sVCAM-1, sCD163, and CRP). Factor B at week 0 was negatively associated with measures of working memory, learning, visual-spatial analysis, and planning/reasoning (KABC-II) and with measures of attention and impulsivity (TOVA D-prime), at both weeks 0 and 48. This factor was also cross-sectionally negatively associated with the KABC-II at week 48. ICAM-1 and VCAM-1 are adhesion molecules expressed by endothelial and immune system cells. Their expression is induced by inflammatory cytokines and results in leukocyte transmigration across the endothelium into tissues.[17–18] Their expression can be associated with endothelial dysfunction.[17-18] CD163 is a marker of monocyte/ macrophage activation and scavenger activity, which can be triggered by endothelial damage and can further contribute to vascular damage.[7] Previous studies have also shown that sCD163 may be a measure of HIV replication.[19] In children with PHIV initiating ART in infancy, CD163 expression has been associated with developmental delays [2] Conversely, in late-diagnosed and treated school-aged children, elevated subsets of activated monocytes following ART initiation were associated with better neurocognitive scores[3], while in adults with acute HIV infection normalization of CD163 expression after ART initiation was associated with lack of neurological sequelae.[20] Combined, these results suggest that timing of infection and ART initiation with respect to individual immune maturation may affect the balance between damaging and protective roles of CD163. CRP, a marker of increased inflammation, was identified in a latent factor (fibrinogen, CRP, and IL-6) negatively associated with a measure of processing speed in youth with virologically suppressed PHIV.[5] Elevated CRP has also been associated with cardiovascular disorder. Adults with HIV are at high risk of cardiovascular disease that may have neurologic complications. The association of elevated CRP, sCD163 and endothelial activation with ND deficits in children with PHIV suggests a potential role of cardiovascular disorder in the genesis of ND deficits that warrants further investigation.

Factor D, consisting of VEGF-A (which promotes angiogenesis in the developing CNS[21–24]) and MIP-1 $\beta$  (a competitive CCR5 receptor ligand[25]) showed favorable longitudinal associations with a measure of attention and impulsivity (TOVA D-prime). CCR5 is a

coreceptor of HIV entry into T cells and monocytes. CCR5 and CCR2 (another HIV coreceptor) antagonists in adults with controlled HIV replication in the blood compartment improved their neurocognitive performance.[26] Collectively, these results suggest a combined neuroprotective effect of angiogenesis and CCR5 antagonism in the developing brains of children with PHIV.

Associations between Factors C and E and ND outcomes were less consistent across timepoints and, therefore, should be interpreted with caution. Factor C, which included the adhesion molecules E-selectin (expressed solely on endothelial cells) and P-selectin (expressed on both endothelial cells and platelets),[6,27] was cross-sectionally associated with poorer motor proficiency (BOT-2) at week 48, suggesting a detrimental effect of endothelial dysfunction on motor outcomes. Factor E (CRP and CD14) at week 0 was associated with a decline in KABC-II scores between weeks 0 and 48. Similar to sCD163, sCD14 is a marker of monocyte/macrophage activation. In the HIV literature, sCD14 has been used as a marker of microbial translocation.[7] Low-grade microbial translocation and elevated sCD14 persists in individuals with HIV despite virologic suppression, remaining at levels higher than in non-infected individuals.[28,29] However, the association of sCD14 with microbial translocation is based on data generated in white males with HIV and has not been confirmed in black people with HIV,[30] such as our study population. Further studies are needed to determine if persistent residual microbial translocation plays a role in the development of ND deficits in children with PHIV.

Factor A, comprised of inflammatory biomarkers (IL-10, IFN $\gamma$ , IFN $\alpha$ , IL-1 $\beta$ , IL-6, IP-10, and TNF $\alpha$ ) involved in multiple cascades of systemic inflammation,[6,7] and Factor F (chemoattractants MCP-1 and CX3CL-1) were not associated with any of the ND outcomes. ND vulnerability in virologically controlled children with PHIV might be better understood by examining specific inflammatory or neuroimmune pathways (such as the ones reflected by Factors B, C, D, and E) rather than the generic pro-inflammatory milieu reflected by Factors A and F.

None of the consensus factors were associated with the measure of executive function (BRIEF). This measure may be relatively less sensitive to plasma biomarker levels because it is based on parent report and could be influenced by factors such as parental depression[31] or a change in caregiver during the interval between measurements.[10]

This study has limitations. It was intended to be hypothesis-generating rather than hypothesis-testing. We report signals of potentially detrimental or protective effects without adjusting for multiple comparisons; thus, hypotheses generated by this analysis will require confirmation in future studies.[14–16] Although we used reproducible, quantitative criteria to identify consensus factors (biomarkers with loadings >0.4), the consensus-based approach is subjective by definition. Nevertheless, our approach was based on the observation of multiple biomarker groupings that varied little over time, guiding our choice to simplify the factors by creating consensus factors across time points. The generalizability of the findings is constrained by the standards and practices in place in 2005, when P1060 enrollment began. We used a cutoff of 400 HIV RNA copies/mL to identify participants with controlled viremia because that was the lower limit of detection for the assay used for most participants

throughout P1060. This cutoff is higher than contemporary assays with limits of detection of <20 to <40 copies/mL. It is possible that viral loads in the range between 20 and 400 copies could have affected levels of inflammatory biomarkers and/or ND outcomes in some participants. Although our multivariable models controlled for potential effects of ART regimen at time of specimen collection, some newer ART agents developed since the completion of P1060 might have different effects on inflammatory markers or ND outcomes compared to the agents used in P1060.

In summary, we identified several biomarker groupings associated with ND outcomes in children with PHIV with controlled viremia. The factor analysis-based approach may lead to an improved ability to identify children with PHIV at high risk of lowered ND performance, understand the underlying mechanisms of persistent ND deficits, and suggest potential therapeutic interventions. Specifically, our results suggest that there may be continuous HIV replication in the CNS compartment of children with PHIV with lowered ND performance that could be controlled with targeted antiviral therapy. An alternative mechanism for ND impairment that needs to be further explored is cardiovascular disorder, which has not been studied in children with PHIV. Mitigating interventions could be deployed if this mechanism is confirmed. Finally, a potential role of CCR5 inhibitors in ameliorating ND deficits in children with PHIV needs to be studied.

# Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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**Figure 1: Boxplots of ND Test Results by Week** ND test results at weeks 0 and 48.

KABC-II MPI: Kaufman Assessment Battery for Children, 2nd edition; TOVA D-prime = Tests of Variables of Attention D-prime; BOT-2: Bruininks-Oseretsky Test of Motor Proficiency, 2nd edition; BRIEF GEC: Behavior Rating Inventory of Executive Function, Global Executive Composite;



Figure 2: Boxplots of biomarkers at three time points (start of controlled viremia, week 0, week 48)

Numbers of participants with available specimens by time point: controlled viremia onset (n = 184), week 0 (n = 185), week 48 (n = 91);

CRP = C-Reactive Protein; CX3CL1 = C-X3-C Motif Chemokine Ligand 1; IL-10 = Interleukin 10; IFNa2 = Interferon a2; IFN $\gamma$  - Interferon  $\gamma$ ; IL-1 $\beta$  = Interleukin 1 $\beta$ ; IL-6 = Interleukin 6; IP-10 = interferon- $\gamma$  inducible protein 10; MCP-1 = Monocyte Chemoattractant Protein 1; MIP-1 $\beta$  = Macrophage Inflammatory Protein 1 $\beta$ ; MMP9 = Matrix Metallopeptidase 9; NFL=neurofilament-light; sCD14 = soluble Cluster of Differentiation 14; soluble Cluster of Differentiation 163; sICAM-1 = soluble Intercellular Adhesion Molecule 1; sICAM-5 = soluble Intercellular Adhesion Molecule 5; sVCAM-1 = soluble Vascular Cell Adhesion Molecule 1; sE-selectin = soluble E-selectin; sP-selectin = soluble P-selectin; TNFa = Tissue Necrosis Factor a; VEGF-A = Vascular endothelial growth factor A;

#### Table 1:

Characteristics of children with PHIV (N=213) included in this analysis.

	Value <sup>a</sup>	
Site <sup>b</sup>		
South Africa: Harriet Shezi	25 (12%)	
South Africa: Soweto (PHRU)	37 (17%)	
South Africa: Tygerberg	48 (23%)	
Malawi: Lilongwe	27 (13%)	
Uganda: Kampala	31 (15%)	
Zimbabwe: Harare	45 (21%)	
Sex		
Male	92 (43%)	
Female	121 (57%)	
Age at initiation of ART (yrs)	1.2 (0.5, 2.6)	
P1060 treatment arm		
NVP-based regimen	109 (51%)	
LPV/r-based regimen	104 (49%)	
Age at P1104s entry (yrs)	7.0 (5.7, 8.7)	
ART regimen at P1104s entry		
NVP-based regimen	63 (30%)	
LPV/r-based regimen	143 (67%)	
Other	7 (3%)	
CD4%, nadir	14.8 (7.7, 24.6)	
Weight z-score at P1104s entry	-0.72 (-1.91, 0.47)	
Height z-score at P1104s entry	-1.01 (-2.24, 0.28)	
Years aviremic before P1104s entry	5.0 (0.8, 7.8)	
Caregiver at entry		
Biological mother	181 (85%)	
Other	32 (15%)	
Socio-economic index <sup>C</sup>	6.1 (2.1, 10.0)	
Premature birth (< 37 weeks)	20 (9%)	
Birthweight < 2000 grams	11 (5%)	

 $^{a}$ Value corresponds to median (10th, 90th percentile) for continuous variables and frequency (percentage) for categorical variables.

<sup>b</sup>All participants with PHIV in P1104s were previously enrolled in P1060 from six study sites in : South Africa (Wits RHI Shandukani clinic [Johannesburg], Chris Hani HIV Unit [Soweto], Family Clinical Research Unit [Cape Town]), Malawi (Kamuzu Central Hospital HIV clinic [Lilongwe]), Uganda (Makerere University – Johns Hopkins University Clinic Mulago National Referral Hospital [Kampala]), and Zimbabwe (Parirenyatwa General Hospital [Harare]).

 $^{C}$ The socioeconomic index is a composite of caregiver education, fuel and water sources, sufficiency of household income for basic needs, caregiver education and work status, major source of information, and whether household has a refrigerator. Values range from 0 (lower SES) to 10 (higher SES).

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# Table 2:

Factor loadings from a seven-factor analysis using principal components method and varimax rotation by each time point

				Start of co	ntrolled v	iremia						-	Week 0							We	ek 48			
Marker	F1	F2	F3	F4	FS	F6	F7	Communality	F1	F2	F3	F4	FS	F6	F7 Con	munality	F1	F2	F3	F4	FS	F6	F7 Commu	nality
IFNa.2	0.77	-0.08	0.21	-0.03	0.23	-0.00	0.02	0.69	0.79	0.07	0.03	-0.13	-0.22	0.02	0.04	0.70	0.80	0.04	-0.13	0.22	0.15 -	0.03	0.21	0.77
IFNγ	0.76	-0.01	0.23	-0.21	0.02	-0.09	0.13	0.69	0.82	0.13	-0.12	0.05	-0.03	- 60.0	0.09	0.72	0.88	0.05	0.07	0.06	0.04	0.08	0.04	0.79
IL-10	0.74	0.22	0.00	0.00	0.13	0.03	-0.08	0.62	0.81	-0.01	-0.09	0.04	0.13	0.04	0.12	0.70	0.72	0.24	0.17 -	0.18	0.35 –	0.17 -	0.08	0.80
IL-1β	0.77	0.02	-0.09	0.23	-0.07	0.07	-0.01	0.66	0.66	0.17	0.19	-0.16	-0.46	-0.18 -	0.07	0.77	0.81	-0.08	-0.12	0.23 -	-0.20	0.11	0.11	0.80
IL-6	0.51	0.15	0.61	0.14	-0.16	-0.00	0.03	0.71	0.89	0.06	0.13	-0.09	0.03	0.05 -	0.09	0.84	0.78	-0.06	-0.25 -	- 0.01	-0.22	0.21	0.00	0.76
IP-10	0.89	0.14	0.15	0.04	-0.04	0.07	-0.04	0.84	06.0	0.13	-0.02	0.01	-0.01	0.07 –	0.08	0.84	0.85	0.15	0.12	0.04 -	-0.03	0.22 -	0.01	0.81
TNFa	0.73	0.22	-0.12	0.22	0.27	0.04	0.04	0.73	0.88	0.12	0.06	0.05	-0.03	-0.03	0.07	0.80	0.80	0.06	0.12 -	0.03	0.35 –	0.05	0.07	0.79
sCD163	-0.13	0.70	0.15	0.09	0.05	0.08	0.06	0.55	0.05	0.44	0.16	0.01	0.62	-0.26 -	0.03	0.67	0.21	0.69	0.07	0.18	0.28	0.08	0.08	0.65
sICAM-1	0.27	0.79	0.20	0.00	-0.15	0.05	-0.06	0.76	0.14	0.85	0.12	-0.13	-0.16	-0.04 -	0.09	0.81	0.00	0.86	0.02	0.12 -	-0.21	0.23	0.10	0.86
sVCAM-1	0.39	0.75	-0.10	-0.03	0.00	0.00	-0.04	0.74	0.10	0.89	0.11	-0.08	0.06	-0.04 -	0.04	0.84	-0.05	0.90	-0.10	0.06 -	-0.01	0.05 -	0.03	0.84
CRP	0.16	0.11	0.74	0.04	-0.15	0.12	0.08	0.63	0.17	0.63	-0.12	0.22	-0.02	0.42	0.13	0.68	0.35	0.48	0.16	0.04	-0.52	0.04	0.08	0.65
sCD14	-0.04	0.05	0.84	-0.12	0.11	-0.10	-0.01	0.74	0.10	-0.00	0.16	-0.13	0.04	0.79 -	0.08	0.68	0.27	0.26	0.29 -	-0.02	-0.38	0.32	0.40	0.64
sE-selectin	-0.06	0.23	0.19	-0.24	0.32	0.67	0.17	0.73	0.04	0.17	0.63	0.06	0.08	0.29	0.01	0.52	0.01	0.31	0.40	0.10 -	0.22	0.63	0.09	0.73
sP-selectin	0.05	0.00	-0.11	0.26	-0.07	0.80	-0.06	0.74	-0.07	0.02	0.83	0.16	0.05	- 00.0-	0.03	0.72	0.23	0.03	-0.14	0.08	0.20	0.82	0.03	0.80
MIP-1β	0.17	0.24	-0.12	0.64	0.01	0.14	-0.16	0.56	0.54	-0.12	0.40	0.21	. 60.0	-0.08	0.22	0.57	0.16	0.04	0.20	0.82	0.17	0.02	0.00	0.77
VEGF-A	0.32	-0.10	0.01	0.64	-0.05	0.33	0.11	0.65	0.15	0.17	0.45	0.06	-0.51	-0.22 -	0.05	0.56	0.06	0.00	-0.19	0.81	-0.03	0.14 -	0.06	0.72
CX3CL1	0.09	-0.31	-0.02	0.27	0.66	-0.02	-0.10	0.62	0.02	-0.06	0.14	0.73	-0.20	0.02	0.07	0.60	-0.12	-0.01	0.27	0.22	0.04	0.26 -	0.69	0.68
MCP-1	0.20	0.12	-0.07	-0.01	0.71	0.09	0.09	0.58	0.61	0.06	-0.01	0.44	0.12	0.14	0.09	0.62	0.31	0.06	-0.12	0.23	0.74	0.15	0.03	0.74
Fibrinogen	0.01	0.07	0.25	0.18	-0.34	0.31	-0.52	0.58	0.02	-0.11	0.13	-0.20	0.71	0.07 -	0.03	0.58	-0.02	-0.15	0.79	0.09	0.04	0.12	0.07	0.67
64MM	-0.23	-0.05	0.10	0.70	0.21	-0.18	0.11	0.64	-0.04	-0.03	0.13	0.74	-0.01	-0.16 -	0.18	0.62	0.01	0.13	0.69	0.07	-0.18	0.13 -	0.21	0.59
NFL	0.03	0.02	0.17	0.09	-0.06	0.13	0.87	0.81	0.01	-0.05	-0.00	-0.08	-0.02	-0.06	0.95	0.92	-0.01	0.07	0.03	0.08	0.03	0.06	0.83	0.71
Eigenvalue	4.40	2.08	2.00	1.74	1.44	1.43	1.17		5.52	2.29	1.67	1.54	1.53	1.11	1.09		5.00	2.57	1.72	1.65	1.61	1.54	1.49	
Communality	refers to	o the pro	portion	of variar	ce in the	e bioma	urker acc	ounted for by	the seve	1 factors	at a give	en time l	ooint											

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Factor loadings (0.4 < |F| - 1.0) denoted by light highlighting.

Color corresponds to the factor loading number at each time point (F1=blue, F2=Red, F3=Green, F4=Orange, F5-F7=Grey).

#### Table 3:

Estimated association between biomarker factor scores at Week 0 and corresponding ND outcomes at Weeks 0 and 48 and Change Between Weeks 0 and 48

Consensus Factor* at Week 0	BOT-2	Reverse-Scored BRIEF GEC	KABC-II MPI	TOVA D-prime
		Week 0 ND meas	urements	
А	0.52 (-0.68, 1.71)	0.78 (-1.11, 2.66)	0.31 (-1.18, 1.80)	-0.04 (-0.18, 0.10)
В	-0.65 (-1.87, 0.57)	-1.38 (-3.28, 0.52)	-2.19 (-3.67, -0.71)	-0.22 (-0.36, -0.07)
С	-0.10 (-1.35, 1.14)	-0.36 (-2.32, 1.60)	0.74 (-0.80, 2.27)	-0.09 (-0.23, 0.06)
D	0.40 (-0.80, 1.60)	1.25 (-0.63, 3.13)	-0.07 (-1.57, 1.42)	-0.08 (-0.22, 0.06)
Е	-0.47 (-1.68, 0.74)	-0.06 (-1.97, 1.85)	-0.51 (-2.01, 0.99)	-0.07 (-0.22, 0.07)
F	0.31 (-1.12, 1.73)	0.68 (-1.56, 2.92)	1.09 (-0.69, 2.87)	0.08 (-0.09, 0.25)
		Week 48 ND meas	surements	
А	-0.36 (-1.54, 0.82)	0.91 (-0.79, 2.61)	-0.24 (-1.96, 1.49)	0.05 (-0.10, 0.21)
В	-1.07 (-2.26, 0.12)	-1.53 (-3.23, 0.18)	-3.04 (-4.73, -1.34)	-0.18 (-0.34, -0.03)
С	-0.11 (-1.34, 1.11)	0.58 (-1.19, 2.35)	0.27 (-1.52, 2.06)	-0.05 (-0.21, 0.11)
D	-0.32 (-1.50, 0.86)	0.40 (-1.30, 2.11)	-0.04 (-1.77, 1.69)	0.17 (0.02, 0.33)
Е	-0.49 (-1.68, 0.69)	-0.75 (-2.47, 0.97)	-1.57 (-3.29, 0.16)	0.01 (-0.15, 0.16)
F	-0.41 (-1.80, 0.98)	0.00 (-2.02, 2.02)	-0.02 (-2.08, 2.04)	0.13 (-0.05, 0.32)
		Week 48 - Week 0 ND	measurements	
А	-0.71 (-1.54, 0.12)	0.62 (-0.82, 2.06)	-0.43 (-1.44, 0.59)	0.08 (-0.05, 0.21)
В	-0.59 (-1.44, 0.26)	-0.82 (-2.28, 0.63)	-0.91 (-1.97, 0.14)	-0.04 (-0.18, 0.10)
С	-0.08 (-0.95, 0.79)	0.66 (-0.83, 2.16)	-0.45 (-1.51, 0.61)	-0.02 (-0.15, 0.12)
D	-0.60 (-1.43, 0.24)	-0.11 (-1.55, 1.34)	0.03 (-0.99, 1.04)	0.20 (0.08, 0.33)
Е	-0.19 (-1.03, 0.65)	-0.69 (-2.14, 0.76)	-1.17 (-2.18, -0.16)	0.04 (-0.09, 0.18)
F	-0.60 (-1.59, 0.39)	-0.35 (-2.06, 1.35)	-0.72 (-1.94, 0.50)	0.05 (-0.11, 0.21)

All analyses adjusted for site, sex, age at enrollment, age at ART initiation, ART regimen at time of biomarker specimen collection (LPV/r-based, NVP-based, or other), and whether the participant switched regimens between P1060 entry and biomarker specimen collection. Analyses involving the change between weeks 0 and 48 also adjusted for values at week 0. Estimates correspond to a one standard deviation increase in the factor score.

Significant regression coefficients (95% CI does not contain 0) denoted by grey highlighting.

Only ND measures with statistically significant relationships with the biomarker factors are shown.

KABC-II MPI = Kaufman Assessment Battery for Children,  $2^{nd}$  edition, Mental Processing Index; TOVA D-prime = Tests of Variables of Attention D-prime; BOT-2 = Bruininks-Oseretsky Test of Motor Proficiency,  $2^{nd}$  edition, Total Motor Composite Score