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# **Oxidized lipoproteins are associated with markers of inflammation and immune activation in HIV-1 infection**

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# **Abstract**

**Objective—**The pathogenesis of immune dysfunction in chronic HIV-1 infection is unclear, and a potential role for oxidized lipids has been suggested. We hypothesize that both oxidized low- and high-density lipoproteins (HDL<sub>ox</sub>, LDL<sub>ox</sub>) contribute to HIV-1 related immune dysfunction.

**Study—**In the AIDS Clinical Trials Group (ACTG) A5260, 234 HIV-infected antiretroviral therapy (ART)-naïve participants were randomized to receive tenofovir-emtricitabine plus protease inhibitors or raltegravir and had HIV-1 RNA <50 copies/ml by week 24 and thereafter.

**Methods—**Associations between biomarkers of inflammation (IL-6, hs-CRP, D-Dimer), immune activation (sCD163, sCD14, sIL-2r, CD38, HLA-DR), inflammatory monocytes (CD14+CD16+), T cell senescence (CD28, CD57) and exhaustion (PD1) and  $HDL_{ox}$ , LDL<sub>ox</sub> were assessed at entry and after ART (week 96) with Spearman (partial) correlations.

**Results—**HDLox declined and LDLox increased over 96 weeks of ART. Positive associations were observed at baseline and over time between  $HDL_{ox}$ , (but not consistently for  $LDL_{ox}$ ) and most markers of inflammation and immune activation (but not senescence/exhaustion), even after adjustment for multiple comparisons, demographics, entry CD4 count and HIV-1 RNA.  $HDL_{ox}$ was positively associated with IL-6 (r=0.19–0.29, p<0.01), and sCD163 (r=0.14–0.41 p 0.04) at all timepoints.

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**Authors contributions:** J.C, J.S., G.M., T.B, T.K. were responsible for the study concept and design. N.J, X.W, D.E carried out the statistical analyses. T.K., N.J, J.C, O.Y drafted the manuscript. T.K., O.Y, M.D, J.S., J.C., G.M. T.B. collected the data. All co-authors participated in discussions about the design of the study, interpretation of the findings, and critically reviewed the manuscript.

**Potential conflicts of interest:**

All authors have submitted the ICMJE Form for Disclosure of Potential Conflicts of Interest. Conflicts that the editors consider relevant to the content of the manuscript have been disclosed.

**Conclusions—**These prospective longitudinal data suggest that oxidized lipoproteins may contribute to persistent immune activation on ART.

#### **Keywords**

Oxidized lipoproteins; HIV; inflammation; immune activation

# **INTRODUCTION**

Human Immunodeficiency Virus type 1 (HIV) infection is characterized by a chronic state of systemic inflammation and immune (T cell and macrophage/monocyte; M/M) activation that is an independent predictor of disease progression. Many of these changes do not normalize despite antiretroviral therapy (ART) [1, 2]. This residual immune activation may be largely responsible for the increased morbidity and mortality observed in HIV-infected subjects on ART but the exact mechanisms are unclear [1, 2].

Oxidative stress is involved in pathogenesis of inflammatory diseases [3] and may impair antiviral immune responses [4]. Lipids play key roles in both viral replication [5] and T cell biology [6]. During oxidative stress oxidized phospholipids acquire novel biological activities such as the ability to regulate innate [7] and adaptive immunity [8, 9] and pathogenesis of many diseases including cardiovascular disease (CVD)[10, 11]. Modified lipoproteins such as oxidized low-density lipoprotein (LDL<sub>ox</sub>) carry oxidized lipids and have pleomorphic atherogenic effects [10].

While high-density lipoprotein (HDL) is generally an anti-inflammatory lipoprotein with protective effects against oxidized lipids and CVD and major immunoregulatory function [12], during systemic inflammation it can be oxidized ( $HDL<sub>ox</sub>$ ), becomes dysfunctional [13, 14] and may contribute to CVD in patients with inflammation [13]. We have shown that HIV-infected subjects have HDL<sub>ox</sub>, demonstrated by impaired antioxidant function and increased lipid hydroperoxide content [14–17] that is associated with measures of subclinical atherosclerosis such as carotid intima media thickness (IMT) [17] and percent non-calcified coronary plaque [18]. In addition, modified lipoproteins may also directly accelerate immune dysfunction and senescence/exhaustion [19, 20] that may contribute to multiple pathologies [21] such as CVD[22]. We [23] and others [24] have shown that oxidized lipoproteins directly upregulate immune activation in vitro. However the exact role of oxidized lipoproteins in HIV pathogenesis is unclear. We hypothesize that there is a cycle of HIV-induced immune activation, inflammation, production of oxidized lipoproteins, and further immune activation and senescence. To elucidate this hypothesis the objective of this study is to investigate in vivo whether oxidized lipoproteins  $(HDL<sub>ox</sub>, LDL<sub>ox</sub>)$  are positively associated with markers of inflammation (hsCRP; IL-6, D- dimer), immune activation (such as sCD14, sCD163, HLA-DR, CD38 expression on CD8 + T lymphocytes) and T cell senescence (CD28, CD57) and to evaluate how these relationships change over time during successful ART.

We evaluated  $HDL_{ox}$ ,  $LDL_{ox}$  in samples from the AIDS Clinical Trials Group (ACTG) A5260s, a prospective study that longitudinally evaluated the changes in biomarkers of inflammation, immunosenescence and immune activation among treatment-naïve individuals

undergoing randomized ART initiation with an integrase-based regimen containing raltegravir (RAL) or a protease inhibitor-based regimen containing either atazanavir/ ritonavir (ATV/r) or darunavir/ritonavir (DRV/r) [25]. This prospective study revealed an overall increase in levels of  $LDL_{ox}$  and decrease in levels of  $HDL_{ox}$  markers of inflammation, coagulation, immune activation,  $CD4<sup>+</sup> T$  cell senescence and exhaustion, and CD8+ T cell exhaustion (but not senescence), which were similar between the different ART regimens after 96 weeks of treatment [25]. Following up on these data, we examined potential associations of plasma oxidized lipoprotein levels with markers of systemic inflammation, coagulation, and immune activation in these individuals.

# **METHODS**

#### **Study Design and Participants**

The design of A5260 and the virologic, tolerability and metabolic outcomes of these ART regimens have been previously reported [25–27]. The parent study and substudy were approved by the Institutional Review Boards at all participating institutions, and all subjects provided written informed consent. For this analysis, the A5260s population was restricted to the subset of virologically suppressed individuals with no ART interruptions greater than seven days and who achieved HIV RNA suppression <50 copies/ml by study week 24 and thereafter.

#### **Determination of biomarkers and plasma oxidized lipoproteins**

Blood samples were drawn at study entry prior to ART initiation and at 24, 48, 96, and 144 weeks on treatment from participants who were required to fast for at least eight hours. Plasma biomarkers of systemic inflammation [IL-6, high-sensitivity C-reactive protein (hsCRP), D-dimer), M/M activation (sCD14, sCD163) and T cell activation [soluble interleukin-2 receptor (sIL-2R)], cellular markers of monocyte (CD163) and T cell (HLA-DR, CD38) activation as well as inflammatory monocyte subsets (non-classical CD14<sup>dim</sup>/  $CD16^+$  and intermediate  $CD14^+CD16^+$ ) have previously been described in this cohort [25]. Oxidized LDL was quantified using ELISA (Mercodia) according to the manufacturer instructions. We also determined the  $LDL_{ox}/LDL$  oxidation ratio which may be a better marker of in vivo oxidation of LDL compared to total levels of  $LDL_{ox}$  [28]. Oxidized HDL was determined using a validated fluorometric cell-free biochemical assay that measures HDL lipid peroxidation ( $HDL_{ox}$ ) [17]. To reduce experimental variability [17] and adjust for HDL amount, we normalized the mean fluorescence readout from quadruplicates of each sample ( $HDL_{ox\,sample}$ ) by the mean fluorescence readout from quadruplicates of a pooled plasma control (HDL<sub>OX</sub> control) and by concurrent HDL cholesterol concentration level (HDL-C) using the following calculation: "normalized" oxidized HDL (nHDL<sub>ox</sub>) =  $[HDL_{ox-sample} \times 40 \ (mg/dl)] / [HDL_{ox\_control} \times HDL-C_{sample} \ (mg/dL)]$ , where 40 mg/dL represents HDL-C of the pooled plasma control. This approach has been validated in clinical studies [14, 29–32]. Throughout the results section oxidized HDL is presented as normalized  $[nHDL<sub>ox</sub>]$  measure to reflect the adjustment for experimental variability and HDL-C.

#### **Statistical Analyses**

Biomarkers were analyzed at study entry (baseline) and further examined at weeks 24 (cellular markers), 48 (plasma markers) and 96. Because of substantial skewness in the variables, a log transformation was used. Changes from baseline were reported as foldchanges, with the 95% confidence interval indicating a statistically significant change at the p <0.05 level. Associations of oxidized lipoproteins and markers of inflammation, immune activation, senescence were examined cross-sectionally prior to (entry) and on ART (week 96) and longitudinally (as fold-change from baseline) using Spearman (partial) correlations which adjusted for the covariates of age, sex, race, BMI, smoking status, baseline CD4 count and baseline viral load. A Spearman partial correlation is similar to its Pearson partial correlation analogue, except that all of the variables are represented as their fractional ranks instead of being in their original units. To calculate the Spearman partial correlation, the covariates (as fractional ranks) are partialed out of two variables of interest, also expressed as ranks. The Pearson correlation between the residuals for the two variables is then the Spearman partial correlation coefficient. This value represents the unique independent effects of the oxidized lipoproteins on the makers of interest. For each set of hypotheses in the multivariate analysis (i.e. each table), the false discovery rate (FDR) was controlled at alpha=0.05 using the Benjamini-Hochberg procedure [33]. Statistical hypothesis tests were two-sided with a significance threshold of 0.05 for p values. All analyses were performed with Stata, version 13.1 (Stata Corp LP, College Station, TX, USA).

# **RESULTS**

#### **Baseline characteristics**

Baseline demographic characteristics of the 328 subjects from the A5260s study population and the 234 subjects (71%) included in the virologically suppressed population for this analysis were previously described [25, 27]. There were no differences in baseline demographic characteristics among treatment groups. Median age was 36 years, 90% of subjects were men and 48% white. Median CD4+ cell count and median HIV RNA were 338 cells/mm<sup>3</sup> and 4.6  $log_{10}$  copies/ml, respectively.

#### **Changes over time in plasma levels of oxidized lipoproteins**

Changes over time in plasma levels of lipoproteins and oxidized lipoproteins are shown in Table 1. Briefly, HDL-C and LDL-C levels increased over time in all ART treated individuals. Levels of normalized  $HDL_{ox}$  declined over week 96 of ART. Post-baseline levels of LDLox and LDLox/LDL increased over 24 weeks and remained elevated compared to baseline levels over 96 weeks of ART.

# **Relationship between oxidized lipoproteins with markers of inflammation, coagulation, immune activation in ART naïve individuals**

We found that  $nHDL_{ox}$  was positively associated with plasma biomarkers of inflammation (IL-6, hs-CRP) and both higher  $nHDL_{ox}$  and  $LDL_{ox}/LDL$  were associated with higher coagulation (D-dimer) at entry prior to ART (Supplemental Table 1)(Figure 1). The strongest relationships between oxidized lipoproteins and markers of inflammation/

coagulation were between  $nHDL_{ox}$  and IL-6 (r=0.36, p<0.01),  $nHDL_{ox}$  and hs-CRP (r=0.27,  $p<0.001$ ) and between  $nHDL_{ox}$  and D-dimer (r=0.19, p<0.01) and remained statistically significant ( $p<sup>FDR</sup> < 0.05$ ) even after adjusting for FDR, age, sex, race, BMI, smoking status, baseline CD4 count and viral load (Table 2). Higher  $nHDL_{ox}$  and  $LDL_{ox}/LDL$  (but not  $LDL<sub>ox</sub>$ ) were associated with higher levels of plasma markers of innate immune activation (sCD163 but not sCD14) and T cell activation (sIL-2r) as well as cellular markers of T cell activation such as expression of CD38 and HLA-DR (Supplemental Tables 1, 2). Similar results were observed for CD4+ and CD8+ T cells.  $LDL_{ox}/LDL$  levels correlated with sCD14 levels. We also found an inverse relationship between  $HDL_{ox}$  and CD38-DR+ T cells, the latter having previously been correlated with favorable outcome in the MACS cohort (Supplemental Table 2)[34]. There were no relationships between levels of plasmaoxidized lipoproteins, inflammatory monocytes and cellular monocyte expression of CD163 (Supplemental Table 2). The most notable positive associations between oxidized lipoproteins and markers of immune activation at baseline were between  $\text{HDL}_{ox}$  and CD38 expression on T cells (for both CD4+ and CD8+T cells r=0.34,  $p<0.001$ ) and sCD163  $(r=0.30, p<0.001)$  (Figure 2, Supplemental Figure 1); these associations remained statistically significant ( $p<sup>FDR</sup> < 0.05$ ) even after adjusting for covariates (Tables 2 and 3).

## **Relationship between oxidized lipoproteins with markers of inflammation, coagulation, immune activation in ART-treated individuals with suppressed viremia**

Consistent with the data in viremic ART-naïve individuals, we also found that  $LDL_{ox}$ ,  $LDL_{ox}/LDL$  and nHD $L_{ox}$  were positively associated with plasma biomarkers of inflammation at 96 weeks of effective ART (Supplemental Table 1). Most notably,  $nHDL_{ox}$ had positive associations with IL-6 (r=0.27, <0.001)(Figure 1) even after adjusting for covariates and applying the FDR correction (Table 2). The relationships of  $LDL_{ox}/LDL$  ratio and  $LDL_{ox}$  with hs-CRP and D-dimer were not consistent (Supplemental Table 1, Tables 2 and 3). We found that higher  $HDL_{ox}$  but not  $LDL_{ox}$  or  $LDL_{ox}/LDL$  were associated with higher levels of plasma sCD163 (r=0.14, 0.05)(at 96 weeks of effective ART (Supplemental Table 1)(Figure 2) but this association did not remain significant after covariate adjustment (Table 2).

## **Declines in plasma levels of oxidized HDL over 96 weeks of ART were associated with declines in markers of inflammation and immune activation**

There were consistent positive associations between declines over 96 weeks in levels of nHDL<sub>ox</sub> and IL-6, hs-CRP, sCD14, sCD163, inflammatory monocytes (Supplemental Table 2) that remained significant after adjusting for covariates and controlling the FDR (Tables 2,3). During this same interval  $LDL_{ox}$  and  $LDL_{ox}/LDL$  ratio increased but there were no consistent relationships between changes in  $LDL_{ox}$ ,  $LDL_{ox}/LDL$  ratio and changes in markers of inflammation, coagulation and immune activation. In the subset of  $nHDL_{ox}$ values that declined most over time (first tertile), all significant associations were more notable and there was a largest decline in levels of IL-6 (data not shown).

# **Association between oxidized lipoproteins and markers of T cell senescence and exhaustion in ART naïve and in ART-treated individuals with suppressed viremia**

We found that higher nHDL<sub>ox</sub> and LDL<sub>ox</sub>/LDL (but not LDL<sub>ox</sub>) were associated with higher cellular markers of T cell senescence (% CD57+ of CD8+ T cells) and exhaustion (% PD1+ of CD4+ and CD8+ T cells) in ART naïve individuals (Supplemental Table 3). These associations were not present at 96 weeks of ART. Changes in levels of  $nHDL_{ox}$  and LDL<sub>ox</sub>/LDL over 96 weeks were positively associated with changes in % CD28-CD57+ of CD4+ T cells, and % PD1+ of CD4+ T cells. LDL<sub>ox</sub>/LDL ratio had differential relationships with markers of T cell senescence and exhaustion compared to  $LDL_{ox}$  (Supplemental Table 3). All the noted relationships were weak, not consistent across all timepoints (baseline, week 96, changes over 96 weeks), were attenuated and did not remain statistically significant after adjusting for covariates (Supplemental Table 4).

#### **DISCUSSION**

In this prospective study of ART-naïve subjects initiating RAL, ATV/r or DRV/r with TDF/FTC and successfully achieving virologic suppression, oxidized lipoproteins and primarily  $HDL_{ox}$  were associated with markers of T cell activation, as well as plasma levels of monocyte activation, inflammation and coagulation. We chose to focus our analyses on markers such as hsCRP; D- dimer, sCD14, sCD163 and CD38 expression that have been associated with serious clinical events in HIV infected persons, including CVD and mortality [35, 36]. To our knowledge, this is the most comprehensive prospective study describing changes in oxidized lipoproteins and immune activation (both M/M and lymphocyte) and inflammation after ART initiation and successful immune suppression. The data from this prospective study with in vivo oxidized lipoproteins from HIV infected subjects are consistent with our prior data [23] and data from others [24] that have shown that in vitro oxidized lipoproteins directly upregulate immune activation in vitro. Thus, we hypothesize that oxidized lipoproteins have a central role in HIV pathogenesis that both result from and contribute to systemic inflammation and immune activation of HIV infection [37].

Consistent with the proinflammatory effect of the oxidized lipoproteins, overall, we found positive associations of both  $HDL_{ox}$  and  $LDL_{ox}$  with various plasma biomarkers of inflammation and coagulation in both viremic and ART treated (at 96 weeks) HIV infected persons.  $HDL_{ox}$  showed consistent positive associations over time with IL-6 and hs-CRP, which have been associated with serious clinical events in HIV infected persons, including CVD and mortality [35, 36]. Interestingly we found that successful ART increased rather than decreased levels of  $LDL_{ox}$ . Initiation of ART within HIV-infected patients reduces markers of systemic inflammation but there is incomplete reversal of systemic inflammation [25]. HIV infected individuals receiving ART may have higher oxidative stress compared to HIV infected naïve or healthy subjects due to higher production of reactive oxygen species (ROS), and alterations in antioxidant systems [38, 39]. Thus, ART, different levels of ROS and antioxidant systems in treated compared to ART-naïve HIV infected persons, may explain the discordant relationship between  $LDL_{ox}$  and markers of systemic inflammation before and after ART initiation.

Consistent with our hypothesis that the immunostimulatory effects of oxidized lipoproteins (that carry oxidized lipids) may contribute to HIV-related immune activation, there were consistent positive associations of both  $HDL_{ox}$  and  $LDL_{ox}/LDL$  with several markers of immune activation in ART-naïve viremic persons. The immunostimulatory effects of oxidized lipoproteins have been shown in other inflammatory states [8]. Dysfunctional HDL that is known to be oxidized [14–17], has previously been shown in vitro to directly influence monocyte [40] and dendritic cell [41] function in systemic inflammatory states [40, 41]. Lipoproteins bind endotoxin which is present in viremic HIV-infected individuals, acts synergistically with  $LDL_{ox}$  [42] and may contribute to immune activation [1]. Thus, further studies are needed to elucidate whether oxidized lipoproteins, through effects on immunity and endotoxin, may contribute to immune activation in viremic HIV infected individuals.

In view of our prior data that oxidized lipoproteins directly induce T cell activation in peripheral blood mononuclear cells from HIV infected ART-treated persons [23], we hypothesized that oxidized lipoproteins may contribute to immune activation in ART-treated HIV-infected individuals. Consistent with this hypothesis we found that after 96 weeks of successful ART,  $HDL_{ox}$  was associated with several biomarkers of immune activation and importantly decline in  $HDL_{ox}$  over 96 weeks was also positively associated with decline over 96 weeks in these biomarkers of immune activation even after adjustment for multiple comparisons, demographics, entry CD4 count and HIV-1 RNA. More specifically, HDLox was associated with sCD163 and there were consistent positive associations between changes over 96 weeks in levels of  $HDL_{ox}$  with several markers of M/M (sCD14, sCD163) inflammatory monocytes) and T cell (cellular expression of CD38, HLA-DR) activation. In a prior small study of 54 HIV infected patients where 91% were receiving ART, both LDL and  $LDL_{ox}$  correlated positively with sCD14 and *in vitro* stimulation with  $LDL_{ox}$ , resulted in expansion of inflammatory monocytes [24]. In a randomized trial of rosuvastatin in HIVinfected subjects on ART,  $LDL_{ox}$  levels decreased 24 weeks after statin therapy, was associated with changes in markers of monocyte activation, and independently predicted changes in IMT, supporting our findings of the importance of oxidized lipids in potentially driving immune activation on ART [43]. In our study, there was also a robust positive association of  $HDL_{ox}$  with sCD163, a marker of M/M activation that has been linked to CVD in HIV disease [44]. These data are consistent with our prior data in a cohort of 102 HIV infected treated subjects, where increased HDL redox activity, a measure of HDL function, correlated positively with sCD163 [18]. Interestingly, the higher increases in baseline CD8+ T cell activation in viremic patients were also seen in subjects with the highest  $HDL_{ox}$  and sCD163. CD163 is shed in response to inflammation and is a scavenger receptor for complexes of hemoglobin/haptoglobin, which may alter HDL function [45]. The exact mechanisms that may mediate the interplay between  $HDL_{ox}$  and immune activation remain to be determined.

Senescent T cells may increase the risk of morbidity, such as CVD [46]. Modified lipoproteins have pleomorphic atherogenic effects [47] and may directly accelerate senescence [19, 20, 48] and contribute to cell exhaustion. We found weak and inconsistent relationships between  $HDL_{OX}$  and  $LDL_{OX}/LDL$  with markers of T cell senescence and exhaustion, across all timepoints, which did not remain significant after covariate

adjustment. Further research is needed to determine whether oxidized lipids may contribute to HIV-related T cell dysfunction.

The differential immunoregulatory effects of  $HDL_{ox}$  versus  $LDL_{ox}$  during the first few years of ART treatment remain unknown. HDL lipids are oxidized in preference to those in LDL when human plasma is exposed to ROS [49]. Moreover, our data corroborate prior evidence that  $LDL_{ox}/LDL$  oxidation ratio may be a better marker of *in vivo* oxidation of  $LDL$ compared to total levels of  $LDL_{ox}$  [28]. Thus, further studies to elucidate the role of oxidized lipoproteins in HIV-related pathogenesis should focus on both HDLox and LDL<sub>ox</sub> as well as the ratio of oxidized lipoproteins to their total plasma levels.

Our study has several limitations. This analysis of oxidized lipoproteins was not the primary outcome of the A5260s study and may have limited power to detect complex modifications of lipoproteins that occur in vivo in the context of measuring oxidized lipoproteins i) in a population with an overall low cardiovascular disease risk ii) in the setting of initiation of different ART regimens during a period where there may be major changes in inflammation and oxidative stress that may further increase between subject variability and may compromise the ability to detect differences in measures of oxidized lipoproteins iii) in cryopreserved rather than fresh samples [14, 17] iv) using biochemical assays that have limitations.

It is also recognized that although our in vitro data suggest that in vitro oxidized lipoproteins directly upregulate immune activation [23], oxidized lipoproteins may also reflect and result from [37] systemic inflammation and immune activation that may be driven by several other mechanisms in HIV infection. For example, ART may directly affect oxidative stress and immune activation in addition to its antiviral effects [38, 50] and these effects may lead to increased plasma levels of oxidized lipoproteins.

Despite the above limitations, this is the most comprehensive prospective study describing changes in oxidized lipoproteins with regards to markers of immune activation and inflammation after ART initiation. Our data support the hypothesis that there is a cycle of HIV-induced immune activation, inflammation, production of oxidized lipoproteins that contributes to further immune activation. Further studies are needed to confirm these findings and further elucidate the differential proinflammatory role of  $HDL_{ox}$  vs  $LDL_{ox}$  in HIV infection. A deeper understanding of the specific mechanisms causing chronic immune activation is crucial to target immune activation in HIV-infected individuals.

#### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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#### **Figure 1.**

Spearman correlations of  $nHDL_{ox}$  with higher levels of markers of systemic inflammation IL-6 (Figure 1 A, B), and hs-CRP (Figure 1 C, D), at baseline (A, C), week 96 after ART (B, D). Oxidized HDL represents the normalized [nHDLox] measure as described in Methods. Spearman correlation coefficients (r) are shown.

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#### **Figure 2.**

Spearman correlations of  $nHDL_{ox}$  with higher levels of markers of M/M activation (sCD163; Figure 2 A, B), and CD8+ T cell activation [CD38 expression on CD8+ T cells (Figure 1 C, D) at baseline (A, C), week 96 (B, D). Oxidized HDL represents the normalized [nHDLox] measure as described in Methods. Spearman correlation coefficients (r) are shown.

#### **Table 1**

Lipid Panel Variables: Fold Change from Baseline over Time among On-Treatment A5260s Subjects with Viral Suppression and No ART Interruption



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# **Table 2**

Partial Spearman Correlations adjusted for Age, Sex, Race, BMI, Current Smoking, Baseline CD4 and RNA (log-scale) level: HDL<sub>ox</sub>, LDL<sub>ox</sub> and Soluble Partial Spearman Correlations adjusted for Age, Sex, Race, BMI, Current Smoking, Baseline CD4 and RNA (log-scale) level: HDL<sub>ox</sub>, LDL<sub>ox</sub> and Soluble Markers of Inflammation/Coagulation Measured Concurrently or Current Change from Baseline over Time. Markers of Inflammation/Coagulation Measured Concurrently or Current Change from Baseline over Time.



Significant (p<0.05) correlations are shown in italics and discrepant correlations between LDL<sub>ox</sub> and LDL<sub>ox</sub>/LDL are shown in bold. Nominal p-values presented. P>0.05 unless noted as statistically y/LDL are shown in bold. Nominal p-values presented. P>0.05 unless noted as statistically

 $^{*}$   $<$  0.05,

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\*\*  $< 0.01$ .

Those with false discovery rate (FDR)  $< 0.05$  are underlined. Those with false discovery rate (FDR) < 0.05 are underlined.

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# **Table 3**

Partial Spearman Correlations adjusted for Age, Sex, Race, BMI, Current Smoking, Baseline CD4 and RNA (log-scale) level: HDL<sub>ox</sub>, LDL<sub>ox</sub> and Partial Spearman Correlations adjusted for Age, Sex, Race, BMI, Current Smoking, Baseline CD4 and RNA (log-scale) level: HDL<sub>ox</sub>, LDL<sub>ox</sub> and Cellular Markers of immune Activation Measured Concurrently or Current Change from Baseline over Time. Cellular Markers of immune Activation Measured Concurrently or Current Change from Baseline over Time.



Nominal p-values presented. P>0.05 unless noted as statistically significant: á ຶ

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 $*$   $<$  0.05,

\*\*  $< 0.01$ .

Significant (p<0.05) correlations are shown in italics and discrepant correlations between LDL<sub>OX</sub> and LDL<sub>OX</sub>/LDL are shown in bold. P values with false discovery rate (FDR) < 0.05 are underlined. Significant (p<0.05) correlations are shown in italics and discrepant correlations between LDL<sub>OX</sub> and LDL<sub>OX</sub>/LDL are shown in bold. P values with false discovery rate (FDR) < 0.05 are underlined.

Similar results were observed for CD4+ and CD8+ T cells and: Similar results were observed for CD4+ and CD8+ T cells and:

 $1$ CD14+CD16+MNCs; CD14+CD16+MNCs;

 $2$  MFI CD163 of CD163+ MNCs MFI CD163 of CD163+ MNCs

 $\mathcal{I}_{\%}$  CD38+ CD8+ T cells (not shown) % CD38+ CD8+ T cells (not shown)