Reduced sTWEAK and Increased sCD163 Levels in HIV-Infected Patients: Modulation by Antiretroviral Treatment, HIV Replication and HCV Co-Infection

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Abstract

Background: Patients infected with the human immunodeficiency virus (HIV) have an increased risk of cardiovascular disease due to increased inflammation and persistent immune activation. CD163 is a macrophage scavenger receptor that is involved in monocyte-macrophage activation in HIV-infected patients. CD163 interacts with TWEAK, a member of the TNF superfamily. Circulating levels of sTWEAK and sCD163 have been previously associated with cardiovascular disease, but no previous studies have fully analyzed their association with HIV.

Objective: The aim of this study was to analyze circulating levels of sTWEAK and sCD163 as well as other known markers of inflammation (hsCRP, IL-6 and sTNFRII) and endothelial dysfunction (sVCAM-1 and ADMA) in 26 patients with HIV before and after 48 weeks of antiretroviral treatment (ART) and 23 healthy subjects.

Results: Patients with HIV had reduced sTWEAK levels and increased sCD163, sVCAM-1, ADMA, hsCRP, IL-6 and sTNFRII plasma concentrations, as well as increased sCD163/sTWEAK ratio, compared with healthy subjects. Antiretroviral treatment significantly reduced the concentrations of sCD163, sVCAM-1, hsCRP and sTNFRII, although they remained elevated when compared with healthy subjects. Antiretroviral treatment had no effect on the concentrations of ADMA and sTWEAK, biomarkers associated with endothelial function. The use of protease inhibitors as part of antiretroviral therapy and the presence of HCV-HIV co-infection and/or active HIV replication attenuated the ART-mediated decrease in sCD163 plasma concentrations.

Conclusion: HIV-infected patients showed a proatherogenic profile characterized by increased inflammatory, immune-activation and endothelial-dysfunction biomarkers that partially improved after ART. HCV-HIV co-infection and/or active HIV replication enhanced immune activation despite ART.

Introduction

Patients with human immunodeficiency virus (HIV) infection have a higher risk of developing cardiovascular disease than does the general population [1–3]. Several potential mechanisms may explain the relationship between HIV infection and atherosclerosis. HIV infection induces persistent inflammation and immune activation, endothelial dysfunction, lipid disorders and direct
vascular injury [4]. Furthermore, although antiretroviral therapy (ART) decreases morbidity and mortality in HIV patients [5], a number of studies have also reported long-term adverse effects of ART on vasculature, including oxidative stress and endothelial dysfunction [6]. Conventional risk prediction models based on traditional cardiovascular risk factors may underestimate the incidence of atherosclerotic cardiovascular events in patients with HIV infection [7] because they do not consider the specific incidence of atherosclerotic cardiovascular events in patients with traditional cardiovascular risk factors may underestimate the dysfunction [6]. Conventional risk prediction models based on ART on vasculature, including oxidative stress and endothelial dysfunction [4]. Furthermore, although antiretroviral therapy decreases morbidity and mortality in HIV patients [5], a number of studies have also reported long-term adverse effects of ART on vasculature, including oxidative stress and endothelial dysfunction [6]. Conventional risk prediction models based on traditional cardiovascular risk factors may underestimate the incidence of atherosclerotic cardiovascular events in patients with HIV infection [7] because they do not consider the specific atherosclerotic processes reported by these patients. Identifying biomarkers for HIV patients at higher cardiovascular risk may be of great interest and could improve cardiovascular risk predictions by traditional stratification scales.

Tumor necrosis factor (TNF)-like weak inducer of apoptosis (TWEAK) is a member of the TNF superfamily that is synthesized as a type II transmembrane glycoprotein and circulates in plasma as a soluble form (sTWEAK) [8]. TWEAK may participate in the development of atherosclerosis by promoting the production of proinflammatory cytokines and altering the proliferation and migration of vascular smooth muscle cells and the expression of extracellular matrix-degrading enzymes [9–11]. sTWEAK plasma levels are decreased in patients with peripheral artery disease (PAD) [12], coronary artery disease (CAD) [13], carotid atherosclerosis [14], abdominal aortic aneurysms [15] and atherosclerosis-associated disorders such as type 2 diabetes or chronic kidney disease [16–18]. Moreover, sTWEAK levels are associated with an adverse prognosis in patients with chronic stable heart failure, myocardial infarction, PAD or non-dialysis CKD [19–22].

CD163 has been recently described as a new receptor for TWEAK present on the surface of monocytes/macrophages [23–24]. A soluble form (sCD163) is generated by the proteolytic cleavage of CD163 at the cell surface in response to proinflammatory and oxidative stimuli [25–27]. The plasma levels of sCD163 are increased in patients with CAD [28], carotid atherosclerosis [24], PAD [12] and atherosclerosis-associated disorders such as type 2 diabetes [29]. The plasma levels of sCD163 have been associated with HIV disease activity in early and chronically infected patients [30]. Moreover, sCD163 plasma levels have been independently associated with noncalcified coronary atherosclerotic plaques in HIV-infected men [31] and correlate with arterial inflammation as determined by positron emission tomography [32]. To our knowledge, there are no studies examining sTWEAK plasma levels in patients with HIV infection. In this study, we analyzed sTWEAK and sCD163 plasma levels and other known markers of inflammation [high-sensitivity C-reactive protein (hsCRP), interleukin 6 (IL-6) and soluble tumor necrosis alpha receptor II (sTNFRII)], endothelial dysfunction (soluble vascular cell adhesion molecule 1 [sVCAM-1]) and asymmetric dimethylarginine (ADMA) and thrombotic activity (D-dimer) in healthy subjects and patients with HIV. Furthermore, we assessed the influence of antiretroviral treatment on these biomarkers after 48 weeks of follow-up.

**Materials and Methods**

**Subjects and design**

Twenty-three individuals without HIV infection (control subjects) and 26 patients infected with HIV were included in the study. HIV patients received ART according to regional recommendations for initiating ART [33]. Patients included in the HIV group were 18 years or older, had a documented HIV-1 infection at least 6 months before entering the study and had not received any antiretroviral drug before the baseline date. Regardless of their HIV status, subjects were excluded if they had known cardiovascular disease, diabetes mellitus, chronic kidney disease, pregnancy, active malignancy or any terminal medical condition. Blood samples were obtained from HIV patients at baseline and 48 weeks after initiating ART. Blood samples were obtained only once from the participants in the control group. This study was conducted at Virgen del Rocío University Hospital and the Biomedicine Institute of Seville (IBiS) (Seville, Spain).

**Ethics statement**

The study was conducted according to the principles expressed in the Declaration of Helsinki. Patients and controls provided written informed consent, and the Ethical Committee of the Virgen del Rocío University Hospital approved the study.

**Laboratory investigation**

Venous blood was collected in EDTA tubes. The whole-plasma samples were stored at −80°C until analyses were performed. Laboratory analyses (lipid, glucose, creatinine, CD4 and CD8 cells, HIV copies and RNA levels) were performed according to routine practice in our hospital. sTWEAK and sCD163 plasma concentrations were measured in duplicate with commercially available ELISA kits (Bender MedSystems, Vienna, Austria) and R&D Systems [Abingdon, UK], respectively. hsCRP, sTNFRII, sVCAM-1 and ADMA levels were also determined with commercially available ELISA kits (Immundiagnostik AG [Ben-heim, Germany], R&D Systems [Abingdon, UK], DLD Diagnostika GmbH [Hamburg, Germany] and eBioscience’s [San Diego, USA], respectively). IL-6 was measured by ultrasensitive ELISA (Quantikine HS Human IL-6 Immunoassay; R&D Systems [Abingdon, UK]). D dimer levels were measured in an automated latex-enhanced immunoassay (HemosIL D-Dimer HS 500; Instrumentation Laboratory [Bedford, UK]). Ig G antibodies to cytomegalovirus were determined by enzyme immunoassay (GenWay Biotech [San Diego, USA]).

**Statistical analysis**

The data are presented as the mean ± standard deviation or the median (interquartile range) for continuous variables depending on the normality of the distribution. Categorical variables are presented as proportion (count). To test whether the variables were normally distributed, we performed the Shapiro-Wilk test. Qualitative variables were analyzed using Fisher’s test and the chi-squared test. The comparisons between two groups were performed using Student’s t-test for normally distributed continuous variables. The Mann-Whitney U test and Wilcoxon matched-pairs test were used to determine significant difference for unpaired and paired data, respectively, if the distribution was non-normal. Pearson correlation coefficients were used to assess correlations for normally distributed data. For non-normally distributed end points, data were either log-transformed, or Spearman rho was used to assess correlation. We conducted a multivariate linear regression analysis in order to evaluate the relative impact of different baseline characteristics (age, triglycerides, BMI, antihypertensive treatment and cytomegalovirus IgG positivity) on plasma sTWEAK, sCD163 and sCD163/sTWEAK levels. P<0.05 was considered statistically significant. Statistics were calculated using the Statistics Package for Social Sciences software (SPSS for Mac OS, version 20.0).

**Results**

**HIV case-control study**

The general characteristics of the study subjects are presented in Table 1. The median time from HIV diagnosis was 5 months.
and 3 received maraviroc during their treatment. Eighteen patients received non-nucleoside reverse transcriptase inhibitors (NNRTIs), 18 received nucleoside reverse transcriptase inhibitors (NRTIs), 12 received protease inhibitors (PIs), 25 received nucleoside reverse transcriptase inhibitors (NRTIs), 18 received non-nucleoside reverse transcriptase inhibitors (NNRTIs) and 3 received nucleoside reverse transcriptase inhibitors (NRTIs) and 3 received maraviroc during their treatment.

HIV-infected patients were older than the control individuals (37.5 ± 10.5 vs. 32.1 ± 8 years; \(P = 0.05\)) and had a higher proportion of positive IgG antibodies to cytomegalovirus (95% vs. 47.6%; \(P = 0.001\)). The proportion of males, active smokers and body mass index (BMI) did not significantly differ between groups. At baseline, HIV-infected patients had significantly higher levels of glucose, triglycerides and the LDL/HDL ratio in addition to lower levels of HDL than controls. Despite the use of ART, HIV-infected patients had higher levels of glucose, triglycerides, BMI, antihypertensive treatment and cytomegalovirus IgG positivity (\(P = 0.012\); \(P < 0.001\) and \(P < 0.001\) respectively) (Figure 1). Differences in sTWEAK, sCD163 and sCD163/sTWEAK levels remained significant after adjusting for age, triglycerides, BMI, antihypertensive treatment and cytomegalovirus IgG positivity (\(P = 0.012\); \(P < 0.001\) and \(P < 0.001\) respectively). Age and anti-hypertensive therapy were independently associated with the sCD163 plasma concentration and the sCD163/sTWEAK ratio, but not with sTWEAK levels.

Table 1. Cardiovascular-related, immunovirologic and biomarkers data in control subjects and HIV-infected patients at baseline and 48 weeks after initiating ART.

|                      | Control (n = 23) | Naïve HIV (n = 26) | \(p^a\) | ART HIV (n = 26) | \(p^b\) | \(p^c\) |
|----------------------|-----------------|-------------------|---------|-----------------|---------|---------|
| Age (years)          | 32.1 ± 8        | 37.5 ± 10.3       | 0.05    | -               | -       | -       |
| Sex (males/females)  | 12/11           | 20/6              | 0.065   | -               | -       | -       |
| Smokers (n)          | 7               | 13                | NS      | -               | -       | -       |
| Treatment for hypertension (%) | 0               | 19.2              | 0.026   | -               | -       | -       |
| BMI (kg/m²)          | 23.7 ± 2.3      | 25.8 ± 4.5        | 0.07    | -               | -       | -       |
| Total cholesterol (mg/dL) | 185.7 ± 29.5    | 170.8 ± 45.3      | NS      | 197.1 ± 43.1    | NS      | 0.001   |
| LDL cholesterol (mg/dL) | 115.6 ± 31.7    | 103.3 ± 37.2      | NS      | 109.6 ± 34.8    | NS      | NS      |
| HDL c                | 62 (60–71)      | 42 (36–52)        | <0.001  | 50 (41–68)      | 0.01    | 0.006   |
| LDLc/HDLc ratio      | 1.72 ± 0.58     | 2.45 ± 0.95       | 0.02    | 2.19 ± 1.01     | NS      | NS      |
| Triglycerides (mg/dL) | 60 (50–91)      | 110 (75–142)      | 0.001   | 123 (89–197)    | <0.001  | 0.039   |
| Creatinine (mg/dL)   | 0.82 ± 0.11     | 0.79 ± 0.15       | NS      | 0.83 ± 0.13     | NS      | NS      |
| Fasting glucose (mg/dL) | 83 ± 6.4        | 89.3 ± 10.7       | 0.025   | 93.1 ± 10       | 0.002   | NS      |
| CD4 (cells/μL)       | -               | 283 (151–405)     | -       | 435 (300–652)   | <0.001  | -       |
| CD8 (cells/μL)       | -               | 743 (570–1130)    | -       | 758 (563–1077)  | -       | NS      |
| HIV-RNA <20 (n/total) | -               | NA                | -       | 23/26           | NA      | -       |
| HIV-RNA (copies/mL)  | 32050 (8520–171246) | 248 (173–759) * | NA*     | -               | NA*     | -       |
| CMV IgG%             | 47.6            | 95                | 0.001   | -               | -       | -       |
| sVCAM-1 (ng/mL)      | 415 (307–601)   | 617 (424–832)     | 0.011   | 393 (318–519)   | NS      | <0.001  |
| ADMA (mg/mL)         | 79.5 (43–112)   | 210 (158–290)     | <0.001  | 194 (166–256)   | <0.001  | NS      |
| hsCRP (mg/L)         | 0.4 (0.21–1.33) | 2.41 (1.62–5.43)  | <0.001  | 1.16 (0.59–1.58)| 0.025   | 0.002   |
| IL-6 (pg/mL)         | 0.42 (0.3–0.74) | 1.02 (0.6–1.43)   | 0.001   | 0.58 (0.36–1.2) | NS      | 0.02    |
| sTNFRII (pg/mL)      | 1923 (1586–2300)| 4553 (3051–6671)  | 0.001   | 2363 (1782–3930)| 0.003   | 0.001   |
| D-dimer (μg/L)       | 287 (220–412)   | 306 (212–582)     | NS      | 269 (212–457)   | NS      | NS      |

BMI = Body mass index, CMV = cytomegalovirus, NS = Not significant, NA = Not applicable. a) \(p\) for control subjects vs. naïve HIV patients. b) \(p\) for control subjects vs. HIV patients with 48 weeks of ART. c) \(p\) for comparison of HIV patients at baseline vs. 48 weeks. d) Proportion of HIV patients with undetectable viral load at 48 weeks. e) Values calculated using the only 3 patients with detectable viral loads at 48 weeks (with descriptive aims, no comparison).

doi:10.1371/journal.pone.0090541.t001
Influence of HCV-HIV co-infection and active viral replication

After 48 weeks of ART, 23 of the 26 HIV-infected patients had undetectable viral loads with a median CD4 cell count of 435 cells/ml (IQR 300–622). The median viral load in the 3 patients with active HIV replication was 248 copies/ml (IQR 173–759). Because HCV-HIV co-infection and active HIV replication may maintain immune activation, we analyzed the possible influence of these conditions on all the biomarkers analyzed. Patients with HCV-HIV co-infection and/or active HIV replication after 48 weeks on ART (n = 5) had a significantly increased sCD163 concentration and sCD163/sTWEAK ratio compared with patients without HCV-HIV co-infection and controlled viral replication (n = 21) [1,290 (IQR 997–2,152) vs. 776 (IQR 501–951) ng/ml; P = 0.01 and 4.08 (IQR 3.04–9.01) vs. 2.15 (IQR 1.3–2.71); P < 0.001, respectively]. No differences were found in the sTWEAK or ADMA levels according to the presence of HCV and active HIV replication. However, a trend toward a worse inflammatory profile with higher hsCRP [2.14 (IQR 0.89–9.6) vs. 0.99 (IQR 0.48–1.55) mg/l; P = 0.11] and sTNFRII [4.404 (IQR 2.660–4.825) vs. 2.203 (IQR 1.741–2.977) pg/ml; P = 0.13] was observed between patients with HCV-HIV co-infection and/or active HIV replication compared to those without these conditions.

Differences observed between HIV patients at baseline versus after ART for hsCRP [2.18 (IQR 1.23–3.65) vs. 0.99 (IQR 0.48–1.53) mg/l; P < 0.001, respectively], IL-6 [1.03 (IQR 0.6–2.32) vs. 0.43 (IQR 0.32–0.89) pg/ml; P = 0.041, respectively], sTNFRII [4,433 (IQR 3,003–6,791) vs. 2,170 (IQR 1,737–2,840) pg/ml; P = 0.008, respectively], sCD163 plasma levels [1,079 (IQR 865–2,252) vs. 776 (IQR 501–951) ng/ml; P = 0.005, respectively] and the sCD163/sTWEAK ratio [2.87 (IQR 2.47–3.36) vs. 2.15 (IQR 1.3–2.71); P = 0.005, respectively] were also significant when patients with HCV-HIV co-infection or active HIV replication were excluded from the analysis.

Correlations between sTWEAK and sCD163 with HIV infection markers

Correlations between sTWEAK, sCD163 and sCD163/sTWEAK with immunovirologic and metabolic parameters are presented in Table 2. sCD163 and the sCD163/sTWEAK ratio was inversely correlated with CD4 nadir at baseline (r = −0.473; P = 0.015 and r = −0.437; P = 0.025, respectively) and with CD4 total count after ART (r = −0.562; P = 0.003 and r = −0.527; P = 0.006, respectively). We did not find a significant correlation between these markers and viral load in HIV-infected patients at baseline or after ART. sTWEAK correlated inversely with hsCRP (r = −0.471; P < 0.001), whereas sCD163 and the sCD163/sTWEAK ratio correlated positively with ADMA (r = 0.689; P < 0.001 and r = 0.615; P < 0.001, respectively), sVCAM-1 (r = 0.374; P = 0.002 and r = 0.441; P < 0.001, respectively) hsCRP (r = 0.445; P < 0.001 and r = 0.557; P < 0.001, respectively), IL-6 (r = 0.468; P = 0.038 and r = 0.471; P = 0.036) and sTNFRII (r = 0.568; P = 0.009 and r = 0.567; P = 0.009) at baseline. After antiretroviral treatment sCD163 and sCD163/sTWEAK ratio correlated positively with IL-6 (r = 0.529; P = 0.008 and r = 0.412; P = 0.045) and sTNFRII (r = 0.576; P = 0.003 and r = 0.559; P = 0.005), but not with the other biomarkers.
Table 2. Correlation between sTWEAK, sCD163 and sCD163/sTWEAK with immunovirological, inflammatory and endothelial biomarkers.

|                  | sTWEAK | sCD163 | sCD163/sTWEAK |
|------------------|--------|--------|---------------|
|                  | Baseline | ART | Baseline | ART | Baseline | ART |
| Age              | 0.062; p = 0.762 | −0.145; p = 0.479 | 0.348; p = 0.081 | 0.200; p = 0.327 | 0.277; p = 0.171 | 0.254; p = 0.211 |
| Viral load       | −0.138; p = 0.502 | −0.12; p = 0.955 | 0.216; p = 0.288 | 0.295; p = 0.143 | 0.207; p = 0.311 | 0.258; p = 0.203 |
| CD4 nadir        | −0.112; p = 0.587 | −0.346; p = 0.083 | −0.473; p = 0.015 | −0.346; p = 0.083 | −0.437; p = 0.223; p = 0.254 |
| CD4 count        | −0.136; p = 0.509 | 0.064; p = 0.756 | −0.361; p = 0.07 | −0.562; p = 0.003 | −0.288; p = 0.154 | −0.527; p = 0.006 |
| CD8 count        | −0.061; p = 0.766 | −0.044; p = 0.831 | −0.132; p = 0.522 | −0.181; p = 0.377 | −0.111; p = 0.589 | −0.144; p = 0.483 |
| CD4/CD8          | −0.227; p = 0.265 | −0.030; p = 0.883 | −0.283; p = 0.162 | −0.393; p = 0.047 | −0.169; p = 0.410 | −0.346; p = 0.083 |
| Tchol            | 0.175; p = 0.392 | 0.095; p = 0.644 | −0.108; p = 0.600 | −0.207; p = 0.311 | −0.299; p = 0.137 | −0.268; p = 0.185 |
| LDL              | 0.252; p = 0.224 | 0.112; p = 0.588 | −0.121; p = 0.564 | −0.364; p = 0.067 | −0.334; p = 0.103 | −0.396; p = 0.045 |
| HDL              | −0.454; p = 0.023 | −0.400; p = 0.043 | −0.082; p = 0.698 | 0.313; p = 0.119 | 0.159; p = 0.447 | 0.446; p = 0.022 |
| Triglycerides    | 0.179; p = 0.382 | 0.530; p = 0.005 | −0.001; p = 0.996 | −0.138; p = 0.500 | −0.183; p = 0.371 | −0.352; p = 0.077 |
| Creatinine       | 0.035; p = 0.867 | 0.012; p = 0.952 | −0.438; p = 0.025 | −0.186; p = 0.364 | −0.329; p = 0.101 | −0.151; p = 0.462 |
| Glucose          | 0.499; p = 0.009 | −0.045; p = 0.828 | −0.056; p = 0.785 | 0.200; p = 0.327 | −0.304; p = 0.131 | 0.188; p = 0.357 |
| ADMA             | −0.187; p = 0.113 | 0.304; p = 0.139 | 0.689; p < 0.001 | 0.262; p = 0.206 | 0.615; p < 0.001 | 0.128; p = 0.540 |
| sVCAM-1          | −0.209; p = 0.088 | −0.352; p = 0.118 | 0.374; p = 0.002 | 0.058; p = 0.8 | 0.441; p = 0.001 | 0.188; p = 0.414 |
| hsCRP            | −0.471; p < 0.001 | −0.289; p = 0.229 | 0.445; p < 0.001 | 0.137; p = 0.576 | 0.557; p < 0.001 | 0.254; p = 0.293 |
| IL-6             | 0.122; p = 0.609 | 0.289; p = 0.17 | 0.468; p = 0.038 | 0.529; p < 0.008 | 0.471; p = 0.036 | 0.412; p = 0.045 |
| sTNFRII          | −0.56; p = 0.816 | −0.046; p = 0.831 | 0.568; p = 0.009 | 0.576; p < 0.003 | 0.567; p < 0.009 | 0.559; p < 0.005 |
| D-dimer          | −0.069; p = 0.772 | 0.048; p = 0.824 | 0.349; p = 0.132 | 0.188; p = 0.379 | 0.394; p = 0.086 | 0.139; p = 0.518 |

BMI = Body mass index; TChol: Total cholesterol.
doi:10.1371/journal.pone.0090541.t002

Discussion

In the present study, we investigated the association among sTWEAK and sCD163 plasma concentrations, the sCD163/ sTWEAK ratio and other inflammatory, endothelial dysfunction and thrombotic biomarkers (hsCRP, IL-6, sTNFRII, sVCAM-1, ADMA and D-dimer) with the presence of HIV infection. We also investigated the impact of ART on these biomarkers. We observed that HIV-infected patients had decreased plasma levels of sTWEAK and increased concentrations of sCD163, hsCRP, IL-6, sTNFRII, sVCAM-1 and ADMA compared with healthy subjects. ART decreased the levels of sCD163, sVCAM-1, hsCRP, IL-6 and sTNFRII but did not affect sTWEAK and ADMA concentrations, reflecting a reduction in inflammation and immune activation associated with HIV.

Several pro-inflammatory mediators and adhesion molecules implicated in the progression of atherosclerosis, including hsCRP, IL-6, sVCAM-1, sCAM-1, P-selectin, E-selectin and ADMA, are elevated in HIV-infected patients [34-36]. IL-6, hsCRP, and D-dimer predict risk for CVD and all-cause mortality in HIV-infected patients [37-39]. The soluble TNF receptors levels have been also associated with the HIV-disease progression [39]. The sVCAM-1 levels are associated with carotid intima media thickness [48], and ADMA is associated with coronary artery calcium scores [41]. Both sVCAM-1 and ADMA are surrogate markers of atherosclerosis. In agreement with these observations, our patients showed increased hsCRP, IL-6, sTNFRII, sVCAM-1 and ADMA plasma concentrations. In addition to these traditional biomarkers, we evaluated the sCD163 plasma concentrations and, for the first time, characterized sTWEAK and the sCD163/ sTWEAK ratio in patients with HIV.

Previous studies have demonstrated that sTWEAK and sCD163 levels are independently associated with cardiovascular disease. sTWEAK plasma concentrations are lower and sCD163 levels are higher in patients with carotid, peripheral and coronary artery disease [12-14,28]. Moreover, circulating sCD163 and sTWEAK are both biomarkers of subclinical atherosclerosis in asymptomatic subjects, independent of traditional cardiovascular risk factors [24]. sTWEAK and the sCD163/sTWEAK ratio have also been associated with long-term global and cardiovascular mortality in patients with peripheral artery disease [22]. We observed that treatment-naive HIV patients had decreased sTWEAK plasma levels and increased concentrations of sCD163 compared with healthy subjects. This is the first study analyzing sTWEAK in HIV-infected patients. The mechanisms leading to lower sTWEAK levels in these subjects are not known. However, decreased sTWEAK plasma concentration has been also reported in other inflammatory pathologies, including rheumatoid arthritis, atherosclerosis and chronic kidney disease [16-18]. A number of studies have reported that the pathological effects of TWEAK are mediated by the binding of TWEAK with its receptor Fn14 [42]. Fn14 expression is increased under inflammatory conditions in multiple tissues [43]. Therefore, the decreased sTWEAK levels observed in our study could reflect an enhanced uptake by Fn14 in inflamed tissues. Although it is not entirely clear [44], it seems that sTWEAK binds to CD163 [23]. Therefore, sTWEAK levels may be also decreased by clearance via its scavenger receptor CD163 [24]. Increased CD163 expression has been reported in monocytes from HIV-infected subjects [45]. On this basis, we hypothesized that both mechanisms, an increased expression of Fn14 and an upregulation of CD163 on monocyte/macrophages, should be involved in the decreased sTWEAK levels observed in HIV-infected patients.
Increased sCD163 plasma levels in HIV-infected patients have been reported in previous studies. Hearps et al. demonstrated a higher concentration of sCD163 in 38 young viremic HIV-positive patients compared with HIV-uninfected controls [46]. In another study, sCD163 levels were elevated in 30 patients with chronic HIV infection and 14 patients with early HIV infection compared with 29 HIV-seronegative individuals [50]. In contrast to these studies and our results, a previous report comparing 11 uninfected donors and 38 HIV-infected patients found no differences in the sCD163 levels, although 23 patients were receiving ART [45]. In our study, the sCD163 levels and the sCD163/sTWEAK ratio were inversely associated with CD4 nadir at baseline and with CD4 count and the CD4/CD8 ratio after ART. Tippett et al. described an association between sCD163 and CD4 count [45], and Burdo et al. reported a similar association between sCD163 and the CD4/CD8 ratio [31]. However, although our results have been confirmed in these previous studies, we must be cautious with these associations due to our relatively small sample size.

sCD163 and the sCD163/sTWEAK ratio decreased significantly after ART, but no changes were observed for sTWEAK. Consistent with our results, Burdo et al. reported a decrease in sCD163 plasma levels after 3 months of ART in chronically infected and early diagnosed HIV patients [31]. In chronically infected HIV patients, sCD163 levels remained elevated compared with control subjects. Additionally, in early diagnosed HIV patients, the sCD163 concentration returned to levels observed in HIV-seronegative individuals [30]. Higher levels of sCD163 in HIV-infected patients with ART have also been reported in some but not all studies [31,46].

We observed a reduction in hsCRP, IL-6, sTNFRII and sVCAM-1 after ART, but no reduction was found in ADMA or D-dimer levels. Previous studies have also described an improvement in these inflammatory and/or endothelial dysfunction biomarkers after ART. Thus, a decrease in sVCAM-1, sICAM-1, E-selectin and hsCRP was observed in 115 HIV-infected patients after 2 and 14 months of antiretroviral treatment [47]. In a further study, TNF-α, sTNFRI, sTNFRII sVCAM-1 and sICAM-1 plasma levels decreased significantly after 24 and 96 weeks of ART [48]. Moreover, D-dimer levels were also found decreased with initiation of ART [49,50]. In contrast to our results, a recent report based on patients in the SMART study showed a decrease in ADMA levels after 12 months of ART; differences were only observed in participants randomized to immediate ART but not in patients randomized to deferral of ART [51]. Therefore, differences in ADMA levels due to ART require further research.

In our study, in spite of ART, sCD163 plasma levels remained elevated in HIV-infected patients compared with control subjects. Higher levels of sCD163 in HIV-infected patients with ART have also been reported in other studies [31,32,46,52]. This result suggests that in spite of controlled viral replication via ART, HIV patients may have a chronic immune activation state that cannot be completely abrogated by ART. In this line, it has been recently proposed that HIV infection trigger immune changes which, if persist through time, may not be completely resolved despite controlling the initial stimulus (HIV infection) [53]. In this sense, there are evidences supporting that the timing of initiating ART may be important to control immune activation. Burdo et al. demonstrated that ART initiated within 1 year of infection (early diagnosed HIV patients) returned sCD163 levels to observed in HIV-seronegative individuals, but not if started later (chronically infected HIV patients) [30]. Moreover, our results show that hsCRP, sTNFRII and ADMA remained elevated in spite of ART, reflecting persistent inflammation and endothelial dysfunction, as previously reported in patients receiving ART [34,47,54]. However, we cannot exclude other potential underlying mechanisms responsible of sCD163 elevation in ART treated HIV patients, such as residual undetected HIV replication, increased translocation of bacterial products from the gastrointestinal tract due to mucosal injury, coinfections such as cytomegalovirus or HCV and the presence of established atherosclerotic lesions. Altogether, chronic immune activation, inflammation and endothelial dysfunction may explain the increased cardiovascular risk observed in these patients even when receiving proper treatment [55].

In our study, sCD163 plasma concentrations and the sCD163/sTWEAK ratio correlated with some inflammatory and endothelial biomarkers in treatment-naive HIV patients, but not after 48 weeks of ART. These results suggest a different effect of ART on monocyte activation and inflammation or endothelial dysfunction. The decrease in sCD163 with ART was modulated by the type of drug used. Thus, patients who were taking PIs had a lower decrease in sCD163 concentration compared with those without PIs. This finding has not been reported previously and could suggest that PIs, despite their antiviral efficacy, may have less effect on controlling monocyte-macrophage activation. Consistent with this suggestion, the relationship between PIs and cardiovascular disease in HIV infected patients is well recognized [6,56,57].

HIV-infected patients receiving ART who had HCV-HIV co-infection and/or persistent HIV replication presented a higher plasma concentration of sCD163 and sCD163/sTWEAK ratio. To our knowledge, there are no previous data related to HCV-HIV co-infection and sCD163. Our findings may indicate that the failure of ART to control viral replication or the existence of HCV-HIV co-infection attenuates the beneficial effect of this therapy on inflammation and macrophage activation. Thus, there may be an increased cardiovascular risk in these patients. In support of this hypothesis, previous data have indicated the importance of viral replication control in vascular disease [5]. Moreover, in a cross-sectional study, Sosner et al. reported that the prevalence of subclinical atherosclerosis was significantly higher in HCV-HIV co-infected patients. The increased atherosclerosis was independent of traditional risk factors, such as LDL-cholesterol and blood pressure [58]. Therefore, the presence of either HCV-HIV co-infection and/or active viral replication may justify a more careful cardiovascular risk management.

One limitation of this study is the relatively low number of patients. However, the strict inclusion criteria for patients and the design of our study limited us to include more HIV patients. Moreover, it is important to note that in the present work we could not analyze the effect mediated by HCV coinfection and/or persistence of HIV replication on immune activation in an independent manner. However, these data raise an attractive hypothesis to be explored in future studies.

In conclusion, we demonstrate that sTWEAK levels are reduced, whereas sCD163 and the sCD163/sTWEAK ratio are increased in HIV patients compared with healthy subjects. ART diminishes sCD163 levels in HIV patients, but no effect is observed for sTWEAK. Our study suggests that treatment with PIs or the presence of HCV-HIV co-infection and/or active HIV replication attenuates the beneficial effects of ART on immune activation, thereby facilitating atherosclerosis.

Acknowledgments

We are indebted to María Jesús Domínguez-Simeón (Unidad Clínico-Experimental de Riesgo Vascular/IBIS) for assistance in the collection of samples; We thank Prof. García Puig, and Dr. Torres (Unidad Metabólico-Vascular/IdiPAZ Hospital La Paz) for assistance in improving the quality
of the manuscript. We thank Oliver Shaw (IIS-FJD) and American Journal Experts for their editorial assistance. We thank Ignacio Mahillo (IIS-FJD), José R. Banegas and Juan J. de la Cruz (Department of Preventive Medicine and Public Health, Universitat Autònoma de Madrid/IdiPAZ – CIBER in Epidemiology and Public Health [CIBER-ESP]) for statistical assistance.

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