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RECEIVED 03 May 2023 ACCEPTED 04 September 2023 PUBLISHED 18 September 2023

CITATION

Nana BC, Esemu LF, Besong ME, Atchombat DHN, Ogai K, Sobgui TMP, Nana CMM, Seumko'o RMN, Awanakan H, Ekali GL, Leke RGF, Okamoto S, Ndhlovu LC and Megnekou R (2023) Soluble biomarkers of HIV-1-related systemic immune activation are associated with high plasma levels of growth factors implicated in the pathogenesis of Kaposi sarcoma in adults. *Front. Immunol.* 14:1216480. doi: 10.3389/fimmu.2023.1216480

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Soluble biomarkers of HIV-1related systemic immune activation are associated with high plasma levels of growth factors implicated in the pathogenesis of Kaposi sarcoma in adults

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Background: Human Herpesvirus-8 (HHV-8) is the etiologic agent of Kaposi's sarcoma (KS), a multicentric angio-proliferative cancer commonly associated with Human Immunodeficiency Virus (HIV) infection. KS pathogenesis is a multifactorial condition hinged on immune dysfunction yet the mechanisms underlying the risk of developing KS in HHV-8 seropositive adults remains unclear. Here we explored whether soluble markers of HIV-1-related systemic immune activation (SIA) and angiogenesis (VEGF and FGF acidic) are involved in the pathogenesis of KS in adults with HHV8.

Methodology: Blood samples from 99 HIV-1 infected and 60 HIV-1 uninfected adults were collected in Yaoundé, Cameroon. CD3+/CD4+ T cell counts and HIV-1 plasma viral load were determined using the Pima Analyzer and the RT-PCR technique, respectively. Plasma levels of SIA biomarkers (sCD163, sCD25/IL-2R α , and sCD40/TNFRSF5) and biomarkers of progression to KS (VEGF and FGF acidic) were measured using the Luminex assay. Seropositivity (IgG) for HHV-8 was determined using the ELISA method.

Results: Overall, 20.2% (20/99) of HIV-1 infected and 20% (12/60) of HIV-1 uninfected participants were seropositive for HHV8. Levels of sCD163, sCD25/IL-2R α , sCD40/TNFRSF5, and FGF acidic were higher in the HIV-1 and HHV8 co-infection groups compared to the HIV-1 and HHV8 uninfected groups (all P <0.05). In addition, Higher plasma levels of VEGF correlated with sCD163 (r_s = 0.58, P =0.0067) and sCD40/TNFRSF5 (r_s = 0.59, P = 0.0064), while FGF acidic levels correlated with sCD40/TNFRSF5 (r_s = 0.51, P = 0.022) in co-infected. In HIV-1 mono-infected donors, VEGF and FGF acidic levels correlated with sCD163 (r_s = 0.25, P = 0.03 and r_s = 0.30, P = 0.006 respectively), sCD25/IL-2R α (r_s = 0.5, P <0.0001 and r_s = 0.55, P <0.0001 respectively) and sCD40/TNFRSF5 (r_s = 0.59, P = 0.0001 respectively) and even in patients that were virally suppressed sCD25/IL-2R α (r_s = 0.39, P = 0.012 and r_s = 0.53, P = 0.0004 respectively) and sCD40/TNFRSF5 (r_s = 0.44, P = 0.0045 respectively).

Conclusion: Our findings suggest that although the development of KS in PLWH is multifactorial, HIV-associated SIA might be among the key drivers in coinfections with HHV8 and is independent of the patients' viremic status.

KEYWORDS

HIV-1, systemic immune activation, IgG anti HHV8, VEGF, FGF acidic

Introduction

Human Herpesvirus-8 (HHV-8) or Kaposi's sarcomaassociated herpesvirus (KSHV) is the etiologic agent of Kaposi's sarcoma (KS), the most common HIV/AIDS-associated cancer (1– 3). Until the advent of the AIDS epidemic, KS was found in specific populations, notably individuals from equatorial Africa (endemic KS), older men from Eastern Europe and the Mediterranean (classic KS), and individuals who received an organ transplant or were immune compromised (iatrogenic KS) (2). Despite being a relatively infrequent malignancy worldwide, KS remains the most common cancer in People Living with HIV (PLWH) in the African continent.

Although the widespread use of antiretroviral therapy (ART) has led to a remarkable decline in KS incidence in PLWH (4), the risk of developing KS in PLWH remains substantially higher than in the general population (5). KS has a viral etiology and multifactorial pathogenesis hinged on immune dysfunction (6). HHV-8 infection can be considered a necessary but not a sufficient condition for the development of KS.

Persistent systemic immune activation (SIA) and inflammation is a hallmark of HIV infection and is strongly associated with AIDS and non-AIDS morbidity and mortality. Several studies demonstrated that plasma levels of immune activation and inflammation biomarkers, as well as those of their mediators, have improved with the introduction of ART (7), but the levels are still higher in PLWH compared to HIV-uninfected populations (8–11). T-cell activation (12), and monocyte/macrophage (13, 14) biomarkers are persistently higher among PLWH, and are implicated in the immunopathogenesis of a wide range of infections in PLWH (15, 16). It has been demonstrated that circulating levels of sCD40/TNFRSF5 and sCD25/IL-2R alpha, both lymphocyte activation biomarkers are elevated in patients with chronic renal failure, chronic liver diseases, and hematological malignancies (17–20) which also affect people with chronic HIV. But their role in other cancers such as KS, an angio-proliferative cancer commonly associated with HIV infection remains unclear.

Angiogenesis is a fundamental event in the process of tumor growth and metastatic dissemination. Angiogenic mediator like Vascular Endothelial Growth Factor (VEGF) has recently been postulated as a major angiogenic and growth factor in several cancer including KS (21). In synergy with Fibroblast Growth Factor (bFGF), VEGF induces endothelial cell growth and vascular permeability in HIV-associated KS (22–24). The question that arises is whether HIV-associated SIA potentially drives angiogenesis in the pathogenesis of KS in co-infections with HHV8? This study therefore aims to evaluate the modulation of soluble biomaker plasma levels of HIV-1-related SIA and that of growth factors (VEGF and FGF acidic) in HIV-1 and HHV8 co-infected adults, in order to define the relationships between these markers across infection status. This may help to preconize a better management and reduce opportunistic infection in adult living with HIV.

Methods

Study design and population

This cross-sectional study was carried out between June 2018 and July 2019 at the HIV clinic of the Efoulan District Hospital and

Central Hospital in Yaoundé, Cameroon. These Hospitals were chosen due to an existing collaboration with the hospitals and University research center and of the high number of people living with HIV visiting these treatment centers. A total of 159 out-patients and in-patients aged 21 to 55 years were enrolled. Sociodemographic and clinical information was recorded. HIVuninfected participants were defined by having negative HIV rapid diagnostic test (RDT) (Alere Determine HIV-1/2, Matsudo, Japan and Oraquick HIV-1/2, Orasure Technologies, Inc) while HIVinfected participants were defined by a positive HIV RDT result. Pregnant women and individuals with malaria infection were excluded from this study to minimize the effect of these conditions on the plasma concentration of biomarkers used in this study. In fact, we wanted to avoid biases due to malaria specific immune responses and pregnancy status, given the fact that pregnancy is characterised by physiological changes that can lead to immunosuppression. Peripheral blood samples were collected in EDTA tubes from all participants. A portion of the blood was used to quantify CD3+/CD4+ Lymphocyte T (LT) cell count and the leftover was centrifuged to collect plasma for storage at -80° C, viral load quantification, and the measurement of immune markers of interest.

CD3+/CD4+ T cell count quantification

Freshly collected EDTA whole blood $(25\mu L)$ was used to determine CD3+/CD4+ T cell counts using the Pima Analyser (Alere Technologies GmbH; Loebstedter Str. 103-105 D-07749 Jena; Germany), as described by the manufacturer. The whole blood samples were charged and inserted into the Pima CD4 test cartridge in the Analyzer, then transported in the incubation compartment where it interacted with the following specific antibodies: an antihuman CD3 monoclonal antibody conjugated to dye 1 and an antihuman CD4 monoclonal antibody conjugated to dye 2. Fluorescence signals were detected by an integrated camera and analyzed using the embedded proprietary software algorithms on the built-in computer. T-cell auxiliaries present both CD3 and CD4 surface antigens and emit a signal light at specific wavelengths to the two antibody-dye conjugates. The results were automatically generated by the Pima Analyzer and the cell counts were expressed per μ L.

HIV-1 serology determination and plasma viral load quantification by real-time PCR

HIV-1 serology was determined according to the National algorithm for HIV Rapid Diagnosis Test (RDT) (25) in Cameroon as described in Figure 1. Whole blood samples were first tested with the RDT1, a highly sensitive test (Alere Determine HIV-1/2, Matsudo, Japan). If the test was reactive for RDT1, a highly specific test RDT2 (Oraquick HIV-1/2, Orasure Technologies, Inc) was used for the confirmation. When reactive with RDT2, the sample was considered positive for HIV but if the RDT2 was nonreactive, the sample was considered indeterminate and an ELISA test (AiDTM anti-HIV 1 + 2 ELISA, Beijing Wantai Biological Pharmacy Enterprise Co., Ltd) was performed to confirm the result. A sample was considered negative for HIV if the RDT1 was non-reactive. Viral load was determined as described by Livo et al. (26). Undetectable viral load threshold was set at <40 copies/mL.



National algorithm for HIV Rapid Test in Cameroon, RDT1, highly sensitive (Alere Determine HIV-1/2, Matsudo, Japan); and RDT2, highly specific (Oraquick HIV-1/2, Orasure Technologies, Inc).

To quantify HIV-1 plasma viral load, the Reverse Transcriptase-Polymerase Chain Reaction (RT-PCR) technique was performed using a RealTime HIV-1 kit on an Abbott m2000rt instrument (Abbott Molecular Inc. 1300 East Tauhy Avenue Des Plaines, IL 60018 USA), according to the manufacturer's instructions. In brief, an RNA sequence that is unrelated to the HIV-1 target sequence was introduced into each sample at the beginning of sample preparation. The amount of HIV-1 target sequence that was present at each amplification cycle was measured through the use of fluorescent-labeled oligonucleotide probes on the Abbott m2000rt instrument. The amplification cycle at which fluorescent signal is detected by the Abbott m2000rt was proportional to the log of the HIV-1 RNA concentration present in the original sample.

Serology test for HHV8

A Human herpes virus-8 IgG Antibody ELISA Kit was used to measure Herpes virus-8 IgG antibodies following the manufacturer's protocols (MyBioSource.com, USA). Briefly, after adding positive control, negative control, and diluted samples in the corresponding well of the micro-ELISA Strip plate, HRP-conjugate Anti-Human IgG antibody was added, and the plates were incubated at 37°C for 60 minutes. After washing plates five times with washing buffer, the substrate was added, and the plate was incubated for 15 minutes at 37°C. Thereafter, the stop solution was added and the Optical Densities (O.D.) were read within 15 minutes at 450 nm using a microplate reader (Spectra Max plus microplate Spectrophotometer, Sunnyvale, CA94089, USA). The results were determined according to the protocol by calculating the critical CUT OFF (Critical CUT OFF= the average of Negative control well + 0.15). Results were considered positive if sample OD was \geq calculated critical CUT-OFF OFF and negative if sample OD was < calculated critical CUT-OFF. This test provided qualitative, but not a quantitative determination of IgG antibodies.

Measurement of soluble biomarkers of systemic immune activation and growth factors

Plasma levels of soluble biomarkers of SIA (sCD163, sCD25/IL-2R alpha, sCD40/TNFRSF5) and two growth factors (VEGF, FGF acidic) were measured by magnetic Luminex screening assay, using Human Premixed Multi-Analyte Kit (R&D Systems, Inc. Minneapolis, MN, USA). The assay was carried out following the protocol and instructions of the manufacturer. Briefly, plates were incubated with 50 μ L/well of each diluted sample and different concentrations of standard cytokines and with 50 μ L/well of Human Magnetic Premixed Microparticle Cocktail containing antibodies specific for each of the analytes. After washing with wash buffer three times using a magnetic plate separator (Luminex, Austin, TX, USA, Cat# CN-0269-01), plates were incubated with 50 μ L/well of Human Premixed Biotin-antibodies cocktail specific for each of the analytes. The second washing step was performed, and antibody-analyte complexes were revealed using 50 μ L/well of Streptavidin-PE. Plates were then incubated for 30 minutes and following the last washing step, 100 μ L/well of wash buffer was added to resuspend the microparticles. Plates were read after 2 minutes of incubation using a Luminex MAGPix Analyzer (XMAP Technology, SN, USA). All the incubation steps were at room temperature on the horizontal shaker. Results were expressed as Median Fluorescence Intensity (MFI). A standard curve was generated for each analyte to convert MFI into corresponding analyte relative concentration.

Statistical analyses

The GraphPad Prism 8.4.3 software was used for statistical analyses. Results were reported as medians with the interquartile range for continuous variables such as age, educational levels, LT CD4 count, and Duration on ARTs, or n (%) where appropriate. Mann–Whitney rank sum test was used to evaluate inter-group differences. Spearman rank order correlation (r_s) was used to evaluate parameter association between the different groups. Proportions were compared using Fisher's exact test. P-values <0.05 were considered statistically significant.

Results

Study population

The baseline characteristics of the study participants are summarized in Table 1. In total, 159 participants were recruited in this study among which, 99 were living with type 1 HIV and 60 were HIV-1 uninfected. The median age was higher in HIV-1 infected (P<0.0001) compared to uninfected group while participants in the non-infected group had a higher level of education than those in the infected group (P<0.0001). More women were living with HIV than men (P = 0.0012). The seroprevalence to HHV8 was 20.12% in this study and was not affected by HIV status (P >0.9999)

In the HIV-1 group, 44 (44,4%) had undetectable viral load and 55 (55,6%) had a detectable viral load. The median CD4 T cell count was significantly higher among the HIV-1-infected group with undetectable viral load compared to those with detectable viral load (P < 0.0001). The percentage of HIV-1-infected participants without immune suppression (CD4 T cell count ≥500 cells/µL) was significantly higher among those with an undetectable viral load, while the percentage of HIV-1-infected individuals with severe immune suppression (CD4 T cell count < 200 cells/µL) was significantly higher among those with detectable viral load (P < 0.0001 for the both). Concerning HHV8, its seroprevalence was higher among HIV-1 infected participants with detectable viral loads compared to those with undetectable viral loads (PVL< 40 cp/ml; P = 0.0219). The median length of duration on ART use was significantly higher among HIV-1 infected participants with an undetectable viral load (UVL< 40 cp/ml) compared to those with detectable viral load (P = 0.0028). A similar trend but not significant (p=0.0638) was observed with the percentage of patients who took ARTs drugs.

Variables		All participants (N = 159)	HIV-1 uninfected (n = 60)	HIV-1 infected (n = 99)		P-values
Age (years) [Median (IQR)]		38 (28 - 45)	27.5 (24.25 - 34.5)	40 (35 - 48)		<0.0001
Sex	Male (%) Female (%)	47 (29.6) 112 (70.4)	27 (45) 33 (55)	20 (20.2) 79 (79.8)		0.0012
Education level [Median (IQR)]		2 (2-4)	4 (1-2)	2 (2-4)		< 0.0001
HHV-8 Seropositivity (n/%)		32 (20.12)	12 (20)	20 (20.2)		>0.9999
		·		Undetectable Viral load (n = 44)	Detectable Viral loal (n = 55)	
CD4 LT count (cells/µL) [Median (IQR)]		-	-	566 (343.8 - 724.5)	184 (70 – 431)	< 0.0001
CD4 LT count \geq 500) (cells/µL) [n/%]	-	-	28 (63.64) 10 (18.18)		<0.0001
350 - 499 (cells/µL)	CD4 LT count [n/%]	-	-	5 (11.36)	8 (14.55)	0.7684
200 – 349 (cells/µL) CD4 LT count [n/%]		-	-	7 (15.91)	9 (16.36)	>0.9999
CD4 LT count <200 (cells/µL) [n/%]		-	-	4 (9.09)	28 (50.91)	< 0.0001
HHV-8 seropositivity (n/%)		-	-	4 (9.09)	16 (29.09)	0.0219
Duration of ARTs taking (month) [Median (IQR)]		-	NA	54 (22-132)	24 (1-84)	0.0028
ART taking (n/%)		-	NA	44 (100)	50 (90.91)	0.0638

TABLE 1 General characteristics of the population.

IQR, Inter Quartile Range; HIV, Human Immunodeficiency Virus; ART, Antiretroviral Tritherapie: P values represent comparisons between HIV negative and HIV positive; and comparison in HIV positive group between undetectable and detectable viral load.

Soluble biomarkers of SIA is Higher in PLWH

To evaluate the impact of HIV infection on the levels of sCD163, sCD25/IL-2R alpha, and sCD40/TNFRSF5, plasma levels of those biomarkers were compared between HIV-1 infected (HIV +) and HIV-1 uninfected adults (HIV-). Results from Figure 2 showed that plasma levels of sCD163 and sCD25/IL-2R alpha were higher among adults living with HIV (ALWH) compared to HIV-1 uninfected adults (P < 0.0031 and P = 0.016 respectively). In

addition, the Spearman rank correlations were used to explore associations between plasma levels of those biomarkers with CD4 T cell count, Viral load, and duration of ARTs taken in ALWH (Table 2). The results showed that plasma levels of sCD163 ($r_s = -0.2294$, P = 0.0420), sCD25/IL-2R alpha ($r_s = -0.3580$; P = 0.0012) and sCD40/TNFRSF5 ($r_s = -0.2978$, P = 0.0077) correlated negatively with CD4 T cell count. Plasma levels of sCD163 ($r_s = -0.3704$, P = 0.0013) and sCD25/IL-2Ralpha ($r_s = -0.2111$, P = 0.0730) inversely correlated with the duration of ART use. Levels of sCD163 ($r_s = 0.2387$, P = 0.0342), sCD25/IL-2R alpha ($r_s = 0.6280$,



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P<0.0001), and sCD40/TNFRSF5 (r_s = 0.4399, P<0.0001) correlated with plasma viral load. These results suggest that HIV infection is associated with high levels of SIA.

Plasma levels of sCD163, sCD25/IL-2R alpha, sCD40/TNFRSF5, VEGF, and FGFacidic are higher in donors with detectable viral loads (DVL)

To determine whether the levels of soluble biomarkers of systemic immune activation (sCD163, sCD25/IL-2R alpha, sCD40/TNFRSF5) and that of growth factors (VEGF, FGF-acidic) varied with viral load in HIV mono-infected participants (HIV-1 positive/HHV-8 negative), plasma levels of these biomarkers were compared between ALWH with undetectable viral load (UVL), those with detectable viral load (DVL) and those uninfected to HIV-1. Results from Figure 3 showed that plasma levels of sCD163 (P = 0.006), sCD25/IL-2R alpha (P < 0.0001), sCD40/TNFRSF5 (P = 0.0008), and VEGF (P = 0.008) were significantly higher among ALWH with DVL compared to those with UVL. Comparing plasma levels of soluble markers between ALWH co-infected with HHV-8 with a DVL (DVL & HHV-8 positive) to those mono-infected to HIV with a DVL (DVL & HHV-8 negative) showed no significant differences (Supplementary Table 1).

Growth factors (VEGF and FGF acidic) levels correlated with 2 SIA biomarkers (sCD25/IL-2R alpha, and sCD40/TNFRSF5) in ALWH with UDL but not with sCD163

To investigate whether SIA in ALWH with UVL could drive the development of Kaposi sarcoma, Spearman rank correlations were used to explore associations between the growth factors and SIA in UVL and DVL donors as shown in Table 3. In ALWH with DVL the levels of VEGF and FGF acidic were positively and significantly correlated with the levels of all SIA biomarkers tested: sCD163 ($r_s = 0.33$, P = 0.038, and $r_s = 0.48$, P = 0.002); sCD25/IL-2R alpha ($r_s = 0.37$, P = 0.018 and $r_s = 0.57$, P = 0.0001); sCD40/TNFRSF5 ($r_s = 0.55$, P = 0.0002 and $r_s = 0.68$, P < 0.0001). Among ALWH with UVL, plasma levels of VEGF and FGF acidic were positively and significantly correlated with the levels of two SIA biomarkers tested: sCD25/IL-2R alpha ($r_s = 0.39$, P = 0.012 and $r_s = 0.53$, P = 0.0004)

and sCD40/TNFRSF5 (r_s = 0.81, P< 0.0001 and r_s = 0.44, P = 0.0045).

Plasma levels of sCD163, sCD25/IL-2R alpha, sCD40/TNFRSF5, VEGF and FGFacidic are elevated in HIV-1 infected adults with seropositivity to HHV8

To determine whether the levels of soluble biomarkers of systemic immune activation (sCD163, sCD25/IL-2R alpha, sCD40/TNFRSF5) and that of growth factors (VEGF, FGF-acidic) varied with HIV-1 and HHV8 co-infection, plasma levels of these biomarkers were compared between ALWH with seropositivity to HHV8 (HIV+&HHV8+) and those uninfected to HIV-1 without seropositivity to HHV8 (HIV-&HHV8-); those uninfected to HIV-1 with seropositivity to HHV8 (HIV-&HHV8+) and those infected to HIV-1 without seropositivity to HHV8 (HIV+&HHV8-). As can be seen in Figure 4 and Table 4 plasma levels of sCD163 (P = 0.003), sCD25/IL-2R alpha (P < 0.0001), sCD40/TNFRSF5 (P = 0.041), and FGF acidic (P = 0.003) were significantly higher among HIV +&HHV8+ compared to double negative HIV-&HHV8-. In addition, plasma levels of sCD25/IL-2R alpha were significantly higher among HIV+&HHV8+ compared to HIV+/HHV8- (P = 0.024). A similar trend was observed when comparing plasma levels of FGF acidic between HIV- and HHV8+ mono-infected and HIVand HHV8- uninfected (P = 0.023) groups. No significant difference has been observed between HIV+&HHV8+ and HIV-&HHV8+ for sCD163 (P = 0.307); sCD25/IL-2R alpha (P = 0.157) and sCD40/ TNFRSF5 (P = 0.7445) and between HIV+ and HHV8+ coinfection and HIV+ and HHV8- mono-infection for sCD163 (P = 0.520); sCD40/TNFRSF5 (P = 0.529).

High plasma levels of VEGF and FGF acidic correlated with sCD163, sCD25/IL-2R alpha, and sCD40/TNFRSF5 levels in HIV-1 and HHV8 co-infected adults

Spearman rank correlations were used to explore associations between plasma levels of growth factors tested (VEGF, FGF-acidic) and soluble biomarkers of systemic immune activation (sCD163, sCD25/IL-2R alpha, sCD40/TNFRSF5) in HIV+&HHV8+, HIV+&HHV8- and HIV-&HHV8+ adults (Table 5). Results

TABLE 2 Correlation between plasma levels of solubles markers of systemic immune activation and CD4 LT count, viral load, and duration of ARTs taken in HIV-1 infected adults.

	LT CD4 count		Viral Load		Duration of ARVs taking	
sCD163	$r_s = -0.2294$	P = 0.0420	$r_{s} = 0.2387$	P = 0.0342	$r_s = -0.3704$	P = 0.0013
sCD25/IL-2R alpha	$r_s = -0.3580$	P = 0.0012	$r_{s} = 0.6280$	P < 0.0001	$r_s = -0.2111$	P = 0.0730
sCD40/TNFRSF5	$r_s = -0.2978$	P = 0.0077	$r_{s} = 0.4399$	P < 0.0001	$r_s = -0.0369$	P = 0.7563

The Sperman's rank Order correlation test was used.



TABLE 3 Correlation between plasma levels of growth factors and soluble biomarkers of SIA stratified by Viral load.

UVL							
	sCD163	sCD25/IL-2R alpha	sCD40/TNFRSF5				
VEGF	$r_s = 0.05 P = 0.74$	$r_s = 0.39 \ P = 0.012$	$r_s = 0.81 \ P < 0.0001$				
FGF acidic	$r_s = 0.05 P = 0.72$	$r_s = 0.53 \ P = 0.0004$	$r_s = 0.44 \ P = 0.0045$				
DVL							
	sCD163	sCD25/IL-2R alpha	sCD40/TNFRSF5				
VEGF	$r_s = 0.33 P = 0.038$	$r_s = 0.37 P = 0.018$	$r_s = 0.55 \ P = 0.0002$				
FGF acidic	$r_s = 0.48 P = 0.002$	$r_s = 0.57 \ P = 0.0001$	$r_s = 0.68 \ P < 0.0001$				

UVL, Undetectable Viral Load; DVL, Detectable Viral Load; VEGF, Vascular Endothelium Growth Factor; FGF-acidic, Fibroblast Growth Factor acidic.

The Spearman's rank Order correlation test was used: $r_s = coefficient$ of correlation; P = coefficient of significance.

showed that plasma levels of VEGF and FGF acidic were positively and significantly correlated with the levels of sCD163 ($r_s = 0.58$, P = 0.0067 and $r_s = 0.43$, P = 0.058); sCD40/TNFRSF5 ($r_s = 0.59$, P = 0.0064 and $r_s = 0.51$, P = 0.022). although positive, this was not significant with sCD25/IL-2R alpha ($r_s = 0.31$, P = 0.17 and $r_s =$ 0.37, P = 0.102) in HIV+/HHV8+ adults. Similar trends were observed in the levels of VEGF and FGF acidic with sCD163 ($r_s =$ 0.25, P = 0.03 and $r_s = 0.30$, P = 0.006), sCD25/IL-2R alpha ($r_s = 0.5$, P<0.0001 and $r_s = 0.55$, P < 0.0001), sCD40/TNFRSF5 ($r_s = 0.7$, P<0.0001 and $r_s = 0.59$, P < 0.0001) among HIV+&HHV8- adults. Also, FGF acidic strongly correlated with sCD40/TNFRSF5 ($r_s =$ 0.65 P = 0.025) but not with sCD163 and sCD25/IL-2R alpha in HIV-&HHV8+. Although positive, no significant correlation has been observed between VEGF and sCD163, sCD40/TNFRSF5, sCD25/IL-2R alpha in HIV-&HHV8+ adults (r_s = 0.25, P = 0.430; r_s = 0.41, P = 0.184; and r_s = 0.13, P = 0.696 respectively).

Discussion

Kaposi's sarcoma (KS) is the most common neoplasm that arises in HIV-infected patients. A better understanding of the relationship between HIV-1-associated immune activation and biomarkers of angiogenesis involve in the pathogenesis of KS is crucial for the identification of potential immunopathogenesis mechanisms linked to the risk of developing KS in HHV-8 seropositive adults. To date, the impact of HIV-1-associated systemic immune activation on the plasma level of some growth



FIGURE 4

Plasma levels of solubles biomarkers of systemic immune activation and growth factors in HIV-1 and HHV8 in adults. Plasma levels of sCD163, sCD25/IL-2R alpha, sCD40/TNFSF5, FGF-acidic and VEGF were compared between HIV+&HHV8+ (N = 20) and HIV-&HHV8- (N = 48) adults using Mann-Whitney Rank Sum test

TABLE 4 Median levels of markers of SIA and angiogenesis in HIV-1 and HH	V-8.
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	HIV+&HHV8+	HIV-&HHV8-	Р	HIV+&HHV8-	HIV-&HHV8-	Р	HIV-&HHV8+	HIV-&HHV8-	Р
sCD163	244612 (186310 - 565526)	107643 (73805 -246830)	0.0031	240588 (118952 -426568)	107643 (73805 -246830)	<0.0001	156672 (43183 - 452162)	107643 (73805 -246830)	0.8165
sCD25/IL- 2R alpha	1262 (730.2 - 2354)	595.0 (422.8 - 907.4)	< 0.0001	722.1 (487.4 - 1333)	595.0 (422.8 - 907.4)	0.0160	834.2 (457.1 - 1487)	595.0 (422.8 - 907.4)	0.1345
sCD40/ TNFRSF5	438.9 (235.6 - 622.9)	327.1 (232.5 - 411.1)	0.0405	383.8 (236.3 - 550.5)	327.1 (232.5 - 411.1)	0.0602	368.4 (257.3 - 649.0)	327.1 (232.5 - 411.1)	0.2428
VEGF	34.33 (6.39 - 78.63)	28.5 (10.68 - 52.42)	0.6912	27.67 (10.77 - 86.24)	28.5 (10.68 - 52.42)	0.4056	23.4 (8.077 - 55.08)	28.5 (10.68 - 52.42)	0.6060
FGF-acidic	7.558 (4.535 - 11.77)	4.535 (1.901 - 6.984)	0.0035	6.084 (4.535 - 10.58)	4.535 (1.901 - 6.984)	0.0035	6.084 (3.802 - 11.77)	4.535 (1.901 - 6.984)	0.0230

HIV+&HHV8+, positive to HIV-1 and seropositive to HHV8; HIV+&HHV8-, positive to HIV-1 and seronegative to HHV8; HIV-&HHV8+, negative to HHV1 and seropositive to HHV8; HIV-&HHV8-, negative to HIV-1 and seronegative to HHV8. Values outside the brackets correspond to the median while values in brackets correspond to the inter-quartile range (25 and 75 percentiles).

factors known as biomarkers of KS progression has not been extensively studied. This study aimed to investigate the relationship between soluble markers of HIV-1-related immune activation and the level of growth factors in HHV8 seropositive adults about KS pathogenesis. The seroprevalence of HHV-8 was 20.12% in the present study and was not affected by HIV status. This result is in accord with those from other studies in sub-Saharan Africa (27-29) and can be explained by the geographical variation

associated with this oncogenic virus. We found that plasma concentrations of soluble biomarkers sCD163 and sCD25/IL-2R alpha were higher among HIV-1 infected compared to HIV-1 uninfected adults. In addition, plasma concentrations of all these soluble biomarkers of immune activation correlated negatively with LT CD4 count but positively with viral load. This is consistent with findings that suggest plasma levels of sCD163, a monocyte/ macrophage activation biomarker, are elevated in HIV-infected TABLE 5 Correlation between plasma levels of growth factors and soluble biomarkers of systemic immune activation in relation to HIV-1 and HHV8 status.

HIV+&HHV8+							
	sCD163	sCD25/IL-2R alpha	sCD40/TNFRSF5				
VEGF	$r_s = 0.58 P = 0.006$	$r_s = 0.31 P = 0.177$	$r_s = 0.59 \ P = 0.006$				
FGF acidic	$r_s = 0.43 P = 0.058$	$r_s = 0.37 \ P = 0.102$	$r_s = 0.51 \ P = 0.022$				
HIV+&HHV8-							
	sCD163	sCD25/IL-2R alpha	sCD40/TNFRSF5				
VEGF	$r_s = 0.24 P = 0.03$	$r_s = 0.50 \ P < 0.0001$	$r_s = 0.71 \ P < 0.0001$				
FGF acidic	$r_s = 0.30 P = 0.006$	$r_s = 0.55 \ P < 0.0001$	$r_s = 0.59 \ P < 0.0001$				
HIV-&HHV8+							
	sCD163	sCD25/IL-2R alpha	sCD40/TNFRSF5				
VEGF	$r_s = 0.25 P = 0.430$	$r_s = 0.13 P = 0.696$	$r_s = 0.41 \ P = 0.184$				
FGF acidic	$r_s = 0.46 P = 0.137$	$r_s = 0.49 \ P = 0.106$	$r_s = 0.65 \ P = 0.025$				

HIV+&HHV8+, positive to HIV-1 and seropositive to HHV8; HIV+&HHV8-, positive to HIV-1 and seronegative to HHV8; HIV-&HHV8+, negative to HIV-1 and seropositive to HHV8; HIV-&HHV8-, negative to HIV-1 and seronegative to HHV8; VEGF, Vascular Endothelium Growth Factor; FGF-acidic, Fibroblast Growth Factor acidic. The Spearman's rank Order correlation test was used: r_s = coefficient of correlation; P = coefficient of significance.

individuals. Even under treatment with ART, abnormal immune activation persists (30-32). In addition, it has been demonstrated that HIV gene products, such as gp120 and Nef, directly stimulate the activation of lymphocytes and macrophages, resulting in the secretion of proinflammatory cytokines and chemokines (33). This study, is among the pioneers that show elevated plasma levels of sCD25/IL-2R alpha and sCD40/TNFRSF5 in PLWH compared to HIV-uninfected adults. We also show that sCD25/IL-2R alpha and sCD40/TNFRSF5 levels correlate positively with viral load but negatively with CD4 LT count. These data indicate that HIV infection is associated with a greater degree of immune activation. Although ART helps reduce inflammation it remains higher than in HIV-1-negative individuals Systemic immune activation has emerged as an essential component of the immunopathogenesis of HIV, it leads to faster disease progression and an accelerated decline of several immune competencies. It is a strong predictor of the development of comorbidity and even mortality in PLWH. This remains true but to a lesser degree in patients who are virally suppressed (34). We demonstrated in this study that in ALWH with UVL, two soluble biomarkers of SIA (sCD25/IL-2R alpha and sCD40/TNFRSF5) were positively associated with plasma levels of growth factors (VEGF and FGF acidic) implicated in the pathogenesis of Kaposi sarcoma. This further highlights the role of SIA in the pathogenesis of KS and shows an increased risk in HIV patients even when they are virally suppressed. We found no difference between SIA and angiogenesis markers in DVL/ HHV8 + vs DVL/HHV8-, and, in UVL/HHV8+ vs UVL/HHV8participants. These suggest that soluble markers of SIA and angiogenesis are not influenced by the infection with HHV-8. However, our lack of statistical power in the DVL/HHV8 + and UVL/HHV8+ pushes us to ascertain these findings in subsequent studies.

Our results reveal that co-infection to HIV-1 and HHV-8 is associated with increased immune activation, as reflected by higher levels of sCD163, sCD25/IL-2R alpha, and sCD40/TNFRSF5 and angiogenesis as assessed by the high levels of FGF acidic, also we find high level of FGF acidic in HHV-8 mono-infected adults. These results suggest that persons with co-infection have higher levels of systemic immune activation markers and may be implicated in the development of KS, and FGF acidic showed potential to be a biomarker of HHV-8 infection in HIV-uninfected adults. Given the evidence that HHV-8 is not sufficient for the development of KS, there is a great need to identify additional factors and mechanisms that may be implicated in the pathogenesis of KS in ALWH. It is worth noting that elevated levels of soluble biomarkers are associated with the severity of other diseases. Several studies have shown elevated sCD163 in the serum of tumor patients and that increased levels were linked to poor prognosis in several cancers (35-38). About sCD25/IL-2R alpha, the levels of this biomarker have been observed to increase in hepatocellular carcinoma and were considered a mediator of T-cell suppression and tumor progression (39, 40). Sebastian and colleagues (41) demonstrated that a high circulating level of sCD40/TNFRSF5 is a marker of liver metastasis risk in rectal cancer. Although the roles of sCD163, sCD25/IL-2R alpha, and sCD40/TNFRSF5 in the control of other diseases have been proven, to our knowledge, this study is the first to demonstrate the association of HIV-1 and HHV-8 coinfection with high levels of these three systemic immune activation biomarkers tested.

The biomarkers of angiogenesis and progression in our study reveal new findings and suggest that HIV-1-related systemic immune activation is associated with high VEGF and FGF acidic levels in HHV8 seropositive adults. In fact, VEGF, an angiogenic and vascular permeability factor, has been postulated as a major angiogenic and growth factor in KS and some previous data indicate that VEGF in synergy with basic FGF induces endothelial cell growth and angiogenesis (25–27). In addition, Felipe and colleagues (42) reported that KS lesions and KS-derived spindle cell cultures co-express high levels of basic FGF and VEGF, which is promoted by the inflammatory cytokines such as tumor necrosis factor (TNF), interleukin-1(IL-1), and interferon- γ (IFN- γ) in PLWH. Although high levels of VEGF and basic FGF have previously been associated with pro-inflammatory cytokines, this study demonstrated for the first time that high levels of VEGF and acidic FGF are associated with soluble biomarkers of systemic immune activation, namely soluble CD163, soluble CD 25/ Inerleukine-2R alpha and soluble CD40/Tumor necrosis factor receptor-SF5 in HIV-1 and HHV8 co-infected adults

The limitation of this study is that it was a cross-sectional study, more data on systemic immune activation biomarkers at different time points are necessary to make a good conclusion on the implication of HIV-1- associated immune activation in the risk of developing KS among HHV-8 seropositive adults. Also, the lack of HHV8 viral load data in this study limited the evaluation of the viral activity of HHV8. The data of this study will serve as preliminary data, in-depth study needs to be conducted for the identification of biomarkers linked to the risk of developing Kaposi sarcoma among people co-infected with HIV-1 and HHV8.

Conclusion

In HIV-1 infection, soluble biomarkers of systemic immune activation are associated with high levels of biomarkers of angiogenesis involves in the pathogenesis of KS in HHV-8 seropositive adults. This may contribute to an increase in the risk among HIV-1 and HHV-8 coinfected adults to develop KS even under controlled viral loads.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding author.

Ethics statement

This study was reviewed and approved by Professor Lazare Kaptué, President of the National Ethics Committee of Cameroon (Ethical Clearance 2019/07/N°38/CE/CNERSH/SP) and by Dr Casimir Doho Beye, Chair Person of the Regional Ethics Committee, Centre Region (Ethical Clearance No CE2023/CRERSHC/2022) for Human Health Research. The studies were conducted in accordance with the local legislation and institutional requirements. The participants provided their written informed consent to participate in this study.

Author contributions

BN and RM conceived and designed the study. BN, DA, TS, HA, and GE help with data acquisition. BN, CN, DA, TS, and RS

performed most of the experiments. BN, LE, MB, and CN analysed and interpreted the data. BN and LE drafted the manuscript. RM, LN, SO, MB, and KO substantively reviewed the final manuscript for important intellectual content. RM, LN, and RL supervised the study. All authors contributed to the article and approved the submitted version.

Funding

This work was supported by the following grants: National Cancer Institute (NCI), Fogarty International Center through Northern Pacific Global Health Fellow Program, National Institutes of Health (NIH) (D43TW009345-10S6), and by Scholarship of the Ministry of Higher Education of Cameroon (21–00134).

Acknowledgments

We thank the Directors (Dr. Mohamadou Guemse) of the Efoulan District Hospital and Hôpital du jour of the Central Hospital (Dr. Kouanfack) for their collaborations. We greatly appreciate the contribution of all the participants who voluntarily accepted to participate in this study. We acknowledge the National Cancer Institute (NCI), Fogarty International Center through the Northern Pacific Global Health Fellow Program, National Institutes of Health (NIH), and the Ministry of Higher Education of Cameroon for the funds that allow for this work. We also thank Dr. Djontu Jean Claude from the Immunology laboratory of the Biotechnology Centre of the University of Yaoundé 1, Cameroon for his help in data analyses.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fimmu.2023.1216480/ full#supplementary-material

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