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## ORIGINAL ARTICLE

**Title:** Elevated liver stiffness is linked to increased biomarkers of inflammation and immune activation in HIV/HCV-coinfected patients

**Running head:** Liver and immune system

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## Abstract

**Objectives:** Immune dysregulation is a hallmark of HIV and HCV infections. To evaluate the relationship between liver stiffness measure (LSM) and biomarkers of T cell activation, bacterial translocation, inflammation, endothelial dysfunction, and coagulopathy in HIV/HCV-coinfected patients.

**Design:** Cross-sectional study.

**Methods:** We studied 238 HIV/HCV-coinfected patients, 32 healthy controls and 39 HIV-monoinfected patients. Patients were stratified according to LSM into four groups: <12.5 kPa, 12.5 to 25 kPa, 25 to 40 kPa, and >40 kPa. T-cell subsets were measured using flow cytometry and plasma biomarkers using immunoassays.

**Results:** HIV/HCV-coinfected patients had higher biomarker levels of immune activation in peripheral blood [T cell activation (CD4<sup>+</sup>CD38<sup>+</sup> and CD8<sup>+</sup>CD38<sup>+</sup>), bacterial translocation (sCD14), inflammation (IL-1b, IL-6, IL-8, IL-18, IP-10) endothelial dysfunction (sVCAM1, sICAM1, and sTNFR1), and coagulopathy (PAI-1)] than healthy controls and HIV-monoinfected patients. Moreover, in HIV/HCV-coinfected patients, a direct relationship between LSM and immune activation [T cell activation (CD8<sup>+</sup>CD38<sup>+</sup> bacterial translocation (LPS), inflammation (IL-8, IP-10), endothelial dysfunction (sVCAM1, sICAM1, and sTNFR1), and coagulopathy (D-dimer)] was found. Subsequently, patients were stratified into different fibrosis stages, finding that patients with cirrhosis who had LSM $\geq$ 40 kPa showed higher biomarker values of immune activation [T cell activation (CD4<sup>+</sup>CD38<sup>+</sup> and CD8<sup>+</sup>CD38<sup>+</sup>), bacterial translocation (LPS), inflammation (IL-8, IL-6, IP-10), endothelial dysfunction (sVCAM1, sICAM1 and sTNFR1), and coagulopathy (D-dimer)] than patients from the other three groups (<12.5 kPa, 12.5-25 kPa, and 25-40 kPa).

**Conclusion:** T cell activation, bacterial translocation, inflammation, endothelial dysfunction, and coagulopathy increased with the severity of liver fibrosis in HIV/HCV-coinfected patients, particularly in patients who had LSM $\geq$ 40 KPa.

## Key Words

Chronic hepatitis C; HIV; cirrhosis; T cell activation; bacterial translocation; inflammation; coagulopathy

## Introduction

Hepatitis C virus (HCV) infection is common among human immunodeficiency virus (HIV) - infected people due to similar routes of transmission [1]. Chronic hepatitis C (CHC) has been a leading comorbidity in HIV-infected patients since the introduction of combination antiretroviral therapy (cART) [2, 3]. The course of CHC may be accelerated in patients coinfecting with HIV, resulting in higher rates of fibrosis progression, cirrhosis, and end-stage liver disease than HCV-monoinfected patients [4]. HIV infection leads to a gradual CD4<sup>+</sup> T cell count decline and causes persistent innate and acquired immune activation, which contributes to the pathogenesis of both AIDS and non-AIDS related diseases [5, 6]. Additionally, despite suppressive cART, HIV-infected patients show abnormally high levels of plasma biomarkers related to immune activation, inflammatory and coagulation markers that may predict increased morbidity and mortality [5]. This immune activation is related to several situations, such as the persistence of HIV replication, HCV coinfection, and bacterial translocation defined as the passage of bacteria or microbial products from the intestinal lumen to mesenteric lymph nodes or other extra-intestinal sites caused by HIV itself and CHC [6, 7].

Bacterial translocation is a key factor in the pathogenesis of both HIV and HCV infection, especially in the advanced stages of disease when bacterial translocation increases (AIDS [8] and cirrhosis [9]). During HIV infection, the depletion of CD4<sup>+</sup> T cells in gut-associated lymphoid tissue (GALT) compromises gut mucosal integrity, promoting increased intestinal permeability that leads to bacterial translocation [8]. In HCV infection, there is a pathological increase in bacterial translocation in liver cirrhosis due to changes and overgrowth of intestinal microbiota, an increase in intestinal permeability and the dysregulation of the immune response in GALT [9]. During this process, host immune cells are stimulated by bacterial pathogen-associated molecular patterns (PAMPs), such as LPS [10], which bind to the CD14/TLR4 complex activating the NF- $\kappa$ B pathway and inducing the synthesis of proinflammatory cytokines such as TNF- $\alpha$ , IL-1, IL-6, etc., and overexpressing chronic activation markers [11-13].

Moreover, during CHC the massive destruction of liver cells leads to dramatic physiological and pathophysiological changes during advanced fibrosis and cirrhosis [14]. Furthermore, cirrhosis is linked to a dysregulation in the balance between activation and homeostasis of the immune system, leading to a state characterized by immune activation and inflammation [13]. In addition, most blood proteins are produced by hepatocytes and their concentration may be

altered by CHC progression, leading to an increased thrombotic risk [15].

The aim of our study was to evaluate the relationship between LSM and biomarkers of T cell activation, bacterial translocation, inflammation, endothelial dysfunction, and coagulopathy in HIV/HCV-coinfected patients.

# Patients and methods

## Patients

We carried out a cross-sectional study in 238 HIV/HCV-coinfected patients, who were selected from the cohort of “Grupo de Estudio del SIDA” (GESIDA 3603b study; see **Appendix**), which is composed of patients enrolled between February 2012 and February 2016 at 19 institutions in Spain. The GESIDA 3603b study is a cohort study that has the aim of evaluating the effect of HCV eradication on the immune system and the associated clinical outcomes. We selected HIV/HCV-coinfected patients at baseline of study.

The cohort included both naïve and anti-HCV therapy experienced patients, who were candidates to receive treatment with HCV therapy (peg-IFN- $\alpha$ /ribavirin or peg-IFN- $\alpha$ /ribavirin/direct-acting antivirals (DAAs)). Anti-HCV therapy in Spain is provided by hospital pharmacies and is covered by the National Health System. The selection criteria were: 1) detectable HCV RNA and HIV RNA by polymerase chain reaction (PCR); 2) availability of a valid baseline LSM; 3) availability of a valid sample of fresh blood to carry out immunological assays; 4) CD4<sup>+</sup> T cell count higher than 200 cells/ $\mu$ L; 5) stable cART for at least 6 months or no need for cART according to guidelines used in the study period. We excluded patients with acute hepatitis C, co-infection with hepatitis B virus, decompensated liver disease, or a prior diagnosis of hepatocellular carcinoma at the time of the transient elastography study.

We selected two control groups for evaluating differences of HIV/HCV-coinfected patients with respect to the normality in peripheral blood biomarkers. On the one hand, the healthy controls are negative subjects for HIV, HCV, and HBV infection. On the other hand, HIV-monoinfected patients with undetectable HIV viral load and CD4<sup>+</sup>>500 cells/mm<sup>3</sup> represent the standard normality of HIV patient without HCV and HBV infection, and other severe comorbidities. The characteristics of both control groups are shown in **Supplemental Table 1**.

The study cohort received the approval of the ethics committees of the participating centers for analysis of anonymized routine clinical data with a view to scientific publication. This work was conducted in accordance with the Declaration of Helsinki. The Institutional Review Board and the Research Ethic Committee of the Instituto de Salud Carlos III approved the study. All patients gave their informed consent for the study.

## **Clinical data**

All the information was recorded at each institution using a common database via an online form, which satisfied local requirements of data confidentiality. This database included all demographic, clinical, virological, and laboratory data. All the centers included in the cohort were monitored to verify that all the information in the database was consistent with the patient's medical records.

We extracted the following baseline data from hospital records: (1) demographics; (2) HIV-related data (HIV transmission category, Centers for Disease Control and Prevention [CDC] clinical category, nadir CD4<sup>+</sup> T cell count, the most recent CD4<sup>+</sup> T cell count, the most recent HIV RNA load, and whether or not patients were receiving cART); (3) liver disease-related data (HCV genotype, HCV RNA load, hepatitis B surface antigen [HBsAg], and anti-HCV therapy); and (4) history of substance abuse including alcohol consumption >50 g/d.

The duration of HCV infections for patients with a history of intravenous drug use (IDU) was estimated starting from the first year they shared needles and other injection paraphernalia [16]. For non-IDU patients, we only calculated the time of infection for those patients that the initiation of HCV infection could be determined with certainty (acute hepatitis C, use of blood and blood products, needle piercing, identified sexual contact, etc.). The time of HIV infections was calculated from the date of HIV diagnosis.

Liver stiffness measurement (LSM) was assessed by transient elastography (FibroScan®, Echosens, Paris, France) using a single machine. Results were expressed in kilopascals (kPa) with a range of 2.5 to 75 kPa. Trained operators performed all FibroScan® examinations. We considered 10 acquisitions with a success rate ≥60% and an interquartile range <30% of the median value as representative measurements of liver stiffness [17]. Fasting was not routinely required prior to the examination.

From these values of LSM, patients were stratified according to the following clinically relevant LSM cutoffs previously used: <12.5 kPa (non-cirrhosis, [17]), 12.5 to 25 kPa (non-risk of bleeding varices, [18]), 25 to 40 kPa (risk of bleeding varices, [18]), and >40 kPa (risk of hepatic decompensation, [19]).

## **Flow cytometry**

The expression of CD38 was evaluated in CD4<sup>+</sup> and CD8<sup>+</sup> T-cell subsets by flow cytometry in 100µL fresh anticoagulated whole blood. The cells were labeled with the following antibodies: anti-CD38-APC-Cyanine 5.5 (APC-Cy5.5, clone HIT2, Invitrogen, Frederick, MD), anti-CD4-APC-

Cyanine 7 (APC-Cy7, clone OKT4, BioLegend, San Diego, CA), anti-CD8-Pacific Blue (PB, clone SK1, BioLegend, San Diego, CA), anti-CD3-Pacific Orange (PO, clone VCHT1, Invitrogen, Frederick, MD) and incubated for 20 min at room temperature in the dark. Next, the IMMUNOPREP Reagent System (Beckman Coulter, Mervue Galway, Ireland) was added to each sample using a Coulter MULTI-Q-PREP Lysing Workstation (Beckman Coulter, Miami, FL) to lyse and fixate them. Fluorescence was measured with a Gallios™ flow cytometer (Beckman Coulter, Miami, FL). The number of events was stopped at a minimum of 200,000 cells in the lymphocyte gate for each sample and flow cytometry data were analyzed using Kaluza™ acquisition software (version 1.5; Beckman Coulter, Miami, FL).

## **Multiplex assay and ELISA**

We selected several biomarkers for each of the objectives we wanted to analyze: a) bacterial translocation [soluble CD14 (sCD14), lipopolysaccharide (LPS), fatty acid-binding protein 2 (FABP2), lipopolysaccharide binding protein (LBP)]; b) inflammation [interleukin (IL)-1b, IL-8, IL-6, IL-18, IFN- $\gamma$ -inducible protein 10 (IP-10)]; c) endothelial dysfunction: soluble vascular cell adhesion molecule 1 (sVCAM1), soluble intercellular cell adhesion molecule 1 (sICAM1), soluble tumor necrosis factor receptor 1 (sTNFR1), monocyte chemoattractant protein-1 (MCP1)]; d) coagulopathy [D-Dimer, plasminogen activator inhibitor-1 (PAI-1)].

Plasma biomarkers were measured by ProcartaPlex™ multiplex immunoassay (ThermoFisher, USA) according to the manufacturer's specifications using a Luminex 200™ analyzer (Luminex Corporation, Austin, TX, United States) with the exception of sCD14 (Raybiotech, Georgia, USA), LPS (HycultBiotech, Uden, The Netherlands), LBP (R&D Systems, Minneapolis, USA) and FABP2 (Raybiotech, Georgia, USA), which were performed according to the manufacturer's procedure for each specific commercial ELISA.

## **Statistical analysis**

The statistical analysis was performed with the Statistical Package for the Social Sciences (SPSS) 21.0 (SPSS INC, Chicago, IL, USA). Statistical significance was defined as  $p < 0.05$ . All  $p$ -values were two-tailed.

For the descriptive study, values were expressed as absolute number (percentage) and median [25th; 75th percentile]. Categorical data and proportions were analyzed using the chi-squared test or Fisher's exact test as required. Kruskal-Wallis and Mann-Whitney tests were used to compare data among independent groups.



Generalized Linear Models (GLM), with a gamma distribution (log-link), was used to evaluate the association between LSM values (continuous variable and ordinal variable) and levels of biomarkers in peripheral blood. This test gives the differences between groups and the arithmetic mean ratio (AMR) or the ratio by which the arithmetic mean of the original outcome is multiplied. Each regression test was adjusted by age, gender, nadir CD4<sup>+</sup> T cells, baseline CD4<sup>+</sup> T cells, HIV viral load (>50 cp/mL), high alcohol intake, diabetes, log<sub>10</sub> HCV RNA, HCV-GT1, previous HCV therapy (IFN $\alpha$ +ribavirin), and prior AIDS.

# Results

## Patients

The characteristics of the 238 HIV/HCV-coinfected patients are shown in **Table 1**. Overall, the median age was 49 years, 78.6% were males, 49.1% had high alcohol intake, 78.1% acquired HIV by IVDU, 26.9% had had prior aids-defining conditions, and 98% were on cART.

Furthermore, the mean CD4<sup>+</sup> T cell count was 574 cells/mm<sup>3</sup>, 12.6% had values of HIV RNA >50 copies/mL, 72.3% were HCV-GT1 and 71.4% had HCV RNA > 850,000 IU/mL. When patients were stratified by LSM values, we only found significant differences between groups in CD4<sup>+</sup> T cells ( $p=0.005$ ) and HCV-GT4 ( $p=0.005$ ).

## HIV/HCV-coinfected patients vs. control groups

HIV/HCV-coinfected patients had higher values of markers of T cell activation [CD4<sup>+</sup>CD38<sup>+</sup> ( $p\leq 0.001$ ), CD8<sup>+</sup>CD38<sup>+</sup> ( $p\leq 0.001$ )] and plasma biomarkers of bacterial translocation [sCD14 ( $p\leq 0.010$ )], inflammation [IL-1b ( $p\leq 0.006$ ), IL-8 ( $p\leq 0.001$ ), IL-6 ( $p\leq 0.001$ ), IL-18 ( $p\leq 0.001$ ), IP-10 ( $p\leq 0.001$ ), sVCAM1 ( $p\leq 0.001$ ), sICAM1 ( $p\leq 0.001$ ), sTNFR1 ( $p\leq 0.001$ )], and coagulopathy [PAI-1 ( $p\leq 0.001$ )] than healthy controls and HIV-monoinfected patients (**Table 2**). Additionally, HIV/HCV-coinfected patients had higher values of MCP1 than healthy controls ( $p=0.005$ ).

## Biomarkers of liver fibrosis

The relationship of liver stiffness measurements and biomarkers of activation, translocation, inflammation, and coagulopathy are shown in **Table 3**. In multivariate analysis, we found that the high LSM values were associated with high levels of CD8<sup>+</sup>CD38<sup>+</sup> (aAMR=1.15;  $p=0.037$ ), LPS (aAMR=1.18;  $p=0.009$ ), IL-8 (aAMR=2.02;  $p<0.001$ ), IP-10 (aAMR=1.27;  $p=0.006$ ), sVCAM1 (aAMR=1.39;  $p<0.001$ ), sICAM1 (aAMR=1.29;  $p=0.045$ ), sTNFR1 (aAMR=1.48;  $p<0.001$ ), and D-Dimer (aAMR=1.71;  $p<0.001$ ).

## Biomarkers of severe cirrhosis

**Table 4** shows the univariate analysis of the biomarker values of activation, translocation, inflammation, and coagulopathy stratified by fibrosis/cirrhosis stages. Patients with >40 kPa had higher values of CD8<sup>+</sup>CD38<sup>+</sup>, IL-8, IL-6, and D-Dimer than patients with <12.5 kPa ( $p\leq 0.005$ ), 12.5-25 kPa ( $p\leq 0.011$ ), and 25-40 kPa ( $p\leq 0.027$ ). Furthermore, patients with >40 kPa had higher values of IP-10, sTNFR1, and PAI-1 than patients with <12.5 kPa ( $p\leq 0.003$ )

and 12.5-25 kPa ( $p \leq 0.004$ ). Additionally, patients with  $>40$  kPa had higher values of sVCAM1 than patients with  $<12.5$  kPa ( $p \leq 0.001$ ).

These differences among groups were also analyzed by multivariate analysis with GLM tests, which showed significant values of aAMR  $>1$  (patients with  $>40$  kPa had higher values than other groups) for several biomarkers (**Table 5**). Patients with  $>40$  kPa had higher values of CD8<sup>+</sup>CD38<sup>+</sup>, IL-6, sICAM1, and D-Dimer than patients with  $<12.5$  kPa ( $p \leq 0.05$ ), 12.5-25 kPa ( $p \leq 0.05$ ), and 25-40 kPa ( $p \leq 0.05$ ). Furthermore, patients with  $>40$  kPa had higher values of LPS, IL-8, IP-10, and sTNFR1 than patients with  $<12.5$  kPa ( $p \leq 0.05$ ) and 12.5-25 kPa ( $p \leq 0.05$ ). Additionally, patients with  $>40$  kPa had higher values of sVCAM1 than patients with  $<12.5$  kPa ( $p \leq 0.001$ ) and CD4<sup>+</sup>CD38<sup>+</sup> than patients with 25-40 kPa ( $p = 0.038$ ).

## Discussion

In this study, HIV/HCV-coinfected patients were found to have higher biomarker levels of immune activation in peripheral blood [T cell activation (CD4<sup>+</sup>CD38<sup>+</sup> and CD8<sup>+</sup>CD38<sup>+</sup>), bacterial translocation (sCD14), inflammation (IL-1b, IL-6, IL-8, IL-18, IP-10), endothelial dysfunction (sVCAM1, sICAM1, and sTNFR1), and coagulopathy (PAI-1)] than healthy controls and HIV-monoinfected patients. Moreover, in HIV/HCV-coinfected patients, we found a direct relationship between LSM values and biomarker values of immune activation [T cell activation (CD8<sup>+</sup>CD38<sup>+</sup>), bacterial translocation (LPS), inflammation (IL-8, IP-10), endothelial dysfunction (sVCAM1, sICAM1, and sTNFR1) and coagulopathy (D-dimer)]. Subsequently, patients were stratified at different fibrosis stages according to their LSM values. The resulting analysis found that patients with cirrhosis who had LSM $\geq$ 40 kPa showed higher biomarker values of immune activation [T cell activation (CD4<sup>+</sup>CD38<sup>+</sup> and CD8<sup>+</sup>CD38<sup>+</sup>), bacterial translocation (LPS), inflammation (IL-8, IL-6, IP-10), endothelial dysfunction (sVCAM1, sICAM1 and sTNFR1), and coagulopathy (D-dimer)] than patients from other groups (<12.5 kPa, 12.5-25 kPa, and 25-40 kPa). We also stratified by established cut-off points of LSM, such as <7.1 kPa (F0-F1), 7.1-9.4 kPa (F2; significant fibrosis), 9.5-12.4 kPa (F3; advanced fibrosis), and  $\geq$ 12.5 kPa (F4; cirrhosis) [20]. However, although some significant differences were found, we did not find a clear trend in the relationship of LSM stages (ordinal variable) with markers of T cell activation and plasma biomarkers of bacterial translocation, inflammation, and coagulopathy in our HIV/HCV-coinfected patients (*data not shown*). To our knowledge, this is the first time that very high levels of these biomarkers of immune activation have been found in HIV/HCV-coinfected patients with compensated cirrhosis. Our results could shed light on the importance of hyperactivation of the immune system in the pathophysiology of CHC in HIV/HCV-coinfected patients with compensated cirrhosis. However, longitudinal studies are necessary to confirm this hypothesis.

In our study, HIV/HCV-coinfected patients had higher plasma sCD14 values than control groups (healthy controls and HIV-monoinfected patients), which may indicate increased bacterial translocation in these subjects, which is usually accompanied by increased inflammation and liver disease severity [21-24], increasing the risk of developing non-AIDS-related complications and death [25]. However, we did not find any association between sCD14 values and LSM in HIV/HCV-coinfected patients. It is possible that the characteristics of our cohort may have played a part in this lack of association, resulting in the loss of a linear relationship between sCD14 and LSM. Moreover, plasma LPS levels were similar in HIV/HCV-coinfected patients and both control

groups, although elevated LPS levels have been previously observed in HIV/HCV-coinfected patients [26]. It is possible that the absence of significant differences may be influenced by the highest sCD14 values in our HIV/HCV-coinfected patients, which may capture and remove LPS from systemic circulation [27], reducing the plasma LPS concentration. However, LPS levels were related to LSM values and cirrhotic patients with  $\text{LSM} \geq 25$  kPa had the highest LPS values. In accordance with our data, an association between plasma LPS levels and advanced stages of liver disease has been found in CHC patients [21, 22]. Thus, LPS may indicate liver disease severity and short-term survival of cirrhotic patients [28, 29]. However, there are also studies that did not find any such association [30, 31].

Regarding T cell immune activation, HIV/HCV-coinfected patients had higher percentages of  $\text{CD4}^+\text{CD38}^+$  and  $\text{CD8}^+\text{CD38}^+$  than healthy controls and HIV-monoinfected patients. Previous reports have shown that HCV infection may increase the percentage of  $\text{CD8}^+\text{CD38}^+$  to above levels of healthy controls [32, 33], with a return to normal values after HCV elimination [33]. In addition, HIV/HCV-coinfected patients have higher levels of immune activation than HIV-monoinfected patients [34]. This immune activation may accelerate CHC [35, 36], and it has been linked to AIDS progression, onset of comorbidities and death in HIV-infected patients [25]. Moreover, high LSM values were linked to high  $\text{CD8}^+\text{CD38}^+$  percentages, and cirrhotic patients with  $\text{LSM} \geq 40$  kPa had the highest  $\text{CD8}^+\text{CD38}^+$  percentages. The activation of the immune system is one of the main pathogenic mechanisms of liver disease, including CHC [9, 13]. Thus, the increased CD38 expression in T cells may indicate an increased risk of CHC progression in HIV/HCV-coinfected patients, as well as an increased risk of progression to AIDS.

Inflammation is characteristic of both HIV and HCV infection, and it plays a key role in the pathogenesis of liver disease in HIV/HCV-coinfected patients [25, 35, 36]. Additionally, inflammation is directly related to endothelial dysfunction, which is implicated in the development of liver diseases and increased cardiovascular risk [37]. In our study, among all biomarkers analyzed, the most relevant were IL-6, IL-8, IP-10, sVCAM1, sICAM1, and sTNFR1. These biomarkers showed higher values in HIV/HCV-coinfected patients and increased as liver stiffness increased, particularly in patients with advanced cirrhosis ( $\text{LSM} \geq 40$  kPa). Overall, our results are in accordance with previous results described in the literature in HCV-infected patients. Firstly, plasma IL-6 levels are higher in patients with HCV infection, particularly in patients with advanced fibrosis or cirrhosis [23, 24, 38, 39]. Secondly, increased circulating IL-8 levels have been linked to CHC progression [40-45]. Thirdly, increased plasma IP-10 levels are related to liver disease severity in HCV-infected patients [26, 44, 46-49]. Fourthly, plasma sVCAM1 and sICAM1 are linked to liver disease severity in HCV-infected

patients [46, 50, 51] and HIV/HCV-coinfected patients [52, 53]. Fifthly, plasma sTNFR1 levels are increased in CHC patients with cirrhosis [54], are related to the influx of portal endotoxin [55] and liver disease severity [56-58], and predict death in patients with cirrhosis [59]. However, the excessive increase of inflammatory biomarkers in our compensated cirrhotic patients with  $\text{LSM} \geq 40$  kPa should be highlighted, because this could indicate an increased risk of AIDS progression [25], development of decompensated cirrhosis and death [12, 35, 36].

Coagulopathy has been linked to an increased risk of progression and death in HIV-infected people [60] and CHC patients [61]. In our study, HIV/HCV-coinfected patients had higher PAI-1 values than healthy controls and HIV-monoinfected patients; whereas high LSM values were directly related to increased levels of D-Dimer with cirrhotic patients with  $\text{LSM} \geq 40$  kPa having the highest D-Dimer levels. Previous studies have reported plasma PAI-1 values to be associated with liver disease severity in patients with non-alcoholic fatty liver disease (NAFLD) [62, 63], and D-Dimer levels are elevated in cirrhotic patients and gradually increase with increasing hepatic dysfunction [64, 65]. Additionally, D-Dimer is related to the development of portal venous thrombosis [66] and bleeding of esophageal varices [67, 68]. D-dimer also predicts in-hospital mortality in patients with hepatic cirrhosis [64]. Thus, coagulopathy biomarkers were also altered in HIV/HCV-coinfected patients, indicating a risk of clinical progression and death, particularly in cirrhotic patients with  $\text{LSM} \geq 40$  kPa.

## **Limitations of study**

Several aspects must be taken into consideration for the correct interpretation of our results. Firstly, this report has a cross-sectional design, which may entail a lack of uniformity, and the study has a limited number of patients in some of the study groups, which could limit the possibility of finding significance. Secondly, all selected patients met a set of criteria for starting HCV treatment (e.g., no alcohol abuse,  $\text{CD4}^+$  cell counts  $>200$  cells/ $\text{mm}^3$ , controlled HIV replication, and good treatment adherence), and this may have introduced a selection bias. Thirdly, we did not have a control group of HCV-monoinfected patients to provide information of possible differential biomarkers from HIV/HCV-coinfected patients. We also did not have any patients with decompensated cirrhosis to study the biomarker profile in end-stage liver disease. Fourthly, our results were not adjusted by multiple comparisons. In this regard, when clinical-orientated studies are not a random search of a meaningful result, it is not recommended adjusting the “p-value” after multiple tests because it can penalize significantly relevant results [67, 68]. Note that our hypothesis is supported by theory and previous reports in patients infected with HIV, HCV, or both, as discussed previously. Additionally, our data had a

clear interpretation since always pointed out in the same direction and the analyzed biomarkers cannot be considered completely independent. Fifthly, we have not used a fixable Live/Dead dye in our freshly whole blood samples, which may influence the results of flow cytometry. However, it is unlikely that there was a bias with respect to a group, since all the samples were processed in the same way. Finally, another limitation of the study is that TE was not always performed in the fasting state, as is currently recommended, since food intake increases liver stiffness in patients with HCV infection [71].

## **Conclusions**

In conclusion, biomarker levels of T cell activation, bacterial translocation, inflammation, endothelial dysfunction, and coagulopathy increased with the severity of liver fibrosis in HIV/HCV-coinfected patients, particularly in patients who had  $LSM \geq 40$  KPa. Further studies are needed to confirm our findings and evaluate whether these biomarkers in HIV/HCV-coinfected patients with compensated cirrhosis can serve as prognostic factors.

## List of abbreviations

Human immunodeficiency virus (HIV)

Hepatitis C virus (HCV)

Liver stiffness measures (LSM)

Chronic hepatitis C (CHC)

Combination antiretroviral therapy (cART)

Gut associated lymphoid tissue (GALT)

Direct-acting antivirals (DAAs)

Soluble CD14 (sCD14)

Lipopolysaccharide (LPS)

Fatty acid-binding protein 2 (FABP2)

Lipopolysaccharide binding protein (LBP)

Interleukin (IL)

IFN- $\gamma$ -inducible protein 10 (IP-10)

Soluble vascular cell adhesion molecule 1 (sVCAM1)

Soluble intercellular cell adhesion molecule 1 (sICAM1)

Soluble tumor necrosis factor receptor 1 (sTNFR1)

Monocyte chemoattractant protein-1 (MCP1)

Plasminogen activator inhibitor-1 (PAI-1)

Generalized Linear Models (GLM)

Arithmetic mean ratio (AMR)

Non-alcoholic fatty liver disease (NAFLD)

Acquired immune deficiency syndrome (AIDS)

Non-nucleoside analogue HIV reverse transcriptase inhibitor (NNRTI)

Nucleoside analogue HIV reverse transcriptase inhibitor (NRTI)

Protease inhibitor (PI)

Integrase inhibitor (II)



# **Declarations**

## **Ethics approval and consent to participate**

The study was conducted in accordance with the Declaration of Helsinki and patients gave their written consent. The Institutional Review Board and the Research Ethic Committee of the Instituto de Salud Carlos III (ISCIII) approved the study.

## **Consent for publication**

Not applicable

## **Availability of data and materials**

The datasets used and/or analyzed during the current study may be available from the corresponding author upon reasonable request.

## **Competing interests**

The authors declare that they have no competing interests.

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## **Author contributions**

Conceptualization: SR, JB, and JGG.

Data curation: JB, JGG, JMG, MC, CQ, JS.

Formal analysis: SR, LMM.

Funding acquisition: JB, JGG, and SR.

Investigation and methodology: LMM, PGB, and IC.

Project Administration: JB.

Supervision and visualization: SR.

Writing – original draft preparation: LMM, PGB, and SR.

Writing – Review & Editing: MAJS, JB.

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## **Authors' information**

Not applicable

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**Table 1.** Clinical and epidemiological characteristics of HIV/HCV-coinfected patients.

	Patients stratified by LSM					<i>p</i>
	All patients	<12.5 kPa	12.5-25 kPa	25-40 kPa	>40 kPa	
<b>No.</b>	238	119	73	28	18	-
<b>Age (years)</b>	49 (46; 52)	48 (45; 52)	49 (46; 51)	49.5 (46; 53)	50 (47; 52)	.482
<b>Gender (male)</b>	187 (78.6%)	96 (80.7%)	54 (74%)	23 (82.1%)	14 (77.8%)	.694
<b>BMI (kg/m<sup>2</sup>)</b>	24.4 (21.8; 26.9)	23.8 (21.4; 26.3)	24.6 (22.7; 28.1)	24.7 (21.9; 26.5)	24.6 (21.6; 25.9)	.247
<b>BMI ≥25 (kg/m<sup>2</sup>)</b>	98 (41.2%)	46 (38.6%)	33 (45.2%)	12 (42.8%)	7 (38.9%)	.693
<b>Diabetes</b>	20 (8.4%)	8 (6.7%)	6 (8.2%)	4 (14.3%)	2 (11.1%)	.600
<b>High alcohol intake</b>	117 (49.1%)	56 (47.1%)	36 (49.3%)	16 (57.1%)	9 (50%)	.876
<b>HIV acquired by IVDU</b>	186 (78.1%)	92 (77.3%)	56 (76.7%)	24(85.7%)	14 (77.7%)	.773
<b>Prior AIDS</b>	64 (26.9%)	27 (22.7%)	23 (31.5%)	8 (28.6%)	6 (33.3%)	.374
<b>Years since HIV diagnosis</b>	23 (18; 26)	22 (17; 26)	24 (20; 27)	21 (18; 26)	23 (19; 27)	.194
<b>Years since HCV infection</b>	21 (16; 24)	21 (13; 24)	21 (18; 25)	19 (17; 21)	22 (15; 26)	.575
<b>Previous HCV therapy (IFNα+rib)</b>	114 (47.9%)	38 (31.9%)	53 (72.6%)	17 (60.7%)	6 (33%)	<b>.001</b>
<b>Antiretroviral therapy</b>						
<b>Non-treated</b>	5 (2.1%)	2 (1.7%)	1 (1.4%)	2 (7.1%)	0 (0%)	.242
<b>PI-based</b>	35 (14.7%)	19 (15.9%)	12 (16.4%)	2 (7.1%)	2 (11.8%)	.627
<b>2NRTI+II-based</b>	59 (24.8%)	31 (26.1%)	18 (24.7%)	6 (21.4%)	4 (23.5%)	.944
<b>2NRTI+PI-based</b>	47 (19.7%)	25 (21%)	10 (13.7%)	6 (21.4%)	6 (35.3%)	.220
<b>2NRTI+NNRTI-based</b>	70 (29.5%)	31 (26.1%)	26 (35.6%)	10 (35.7%)	3 (17.6%)	.344
<b>Others</b>	22 (9.2%)	11 (9.2%)	6 (8.2%)	2 (7.2%)	2 (11.8%)	.274
<b>HIV markers</b>						
<b>Nadir CD4+ T cells</b>	172 (84; 254)	196 (78; 277)	167 (87; 234)	160 (85; 251)	116 (95; 198)	.345
<b>Nadir CD4+ T cells&lt;200 cells/mm<sup>3</sup></b>	131 (55.1%)	58 (48.7%)	42 (57.5%)	17 (60.7%)	14 (77.7%)	.100
<b>CD4+ T cells</b>	547 (394; 803)	603 (436; 832)	511 (339; 736)	570 (395; 828)	364 (243; 520)	<b>.005</b>
<b>CD4+ T cells&lt;500 cells/mm<sup>3</sup></b>	100 (42%)	41 (34.4%)	35 (47.9%)	13 (46.4%)	11 (61.1%)	.065
<b>HIV-RNA &gt;50 cp/mL</b>	30 (12.6%)	16 (13.4%)	7 (9.6%)	4 (14.3%)	3 (16.7%)	.808
<b>HCV markers</b>						
<b>HCV genotype (n=235)</b>						
<b>1</b>	170 (72.3%)	78 (66.7%)	58 (80.6%)	20 (71.4%)	14 (77.8%)	.205
<b>2</b>	5 (2.1%)	3 (2.6%)	1 (1.4%)	1 (3.6%)	0 (0%)	.915
<b>3</b>	39 (16.6%)	18 (15.4%)	12 (16.7%)	7 (25%)	2 (11.1%)	.599

4	21 (8.9%)	18 (15.4%)	1 (1.4%)	0 (0%)	2 (11.1%)	.005
<b>Log<sub>10</sub> HCV-RNA (IU/mL)</b>	6.3 (5.87; 6.74)	6.32 (5.83; 6.83)	6.36 (6.04; 6.69)	6.36 (5.79; 6.52)	6.16 (5.78; 6.59)	.692
<b>HCV-RNA &gt; 500,000 IU/mL</b>	191 (80.2%)	92 (77.3%)	63 (86.3%)	23 (82.1%)	13 (72.2%)	.495
<b>HCV-RNA &gt; 850,000 IU/mL</b>	170 (71.4%)	80 (97.2%)	58 (79.4%)	19 (67.9%)	13 (72.4%)	.608

**Statistics:** Values expressed as absolute number (percentage) and median (interquartile range). *P*-values were calculated by Chi-square tests and Mann-Whitney tests in HIV/HCV-coinfected patients stratified by LSM (<12.5 kPa, 12.5-25 kPa, 25-40 kPa, and >40 kPa).

**Abbreviations:** HCV, hepatitis C virus; HCV-RNA, HCV plasma viral load; HIV-1, human immunodeficiency virus type 1; LSM, liver stiffness measure; HIV-RNA, HIV plasma viral load; IVDU, intravenous drug user; AIDS, acquired immune deficiency syndrome; IFN $\alpha$ +rib, interferon-alpha plus ribavirin; NNRTI, non-nucleoside analogue HIV reverse transcriptase inhibitor; NRTI, nucleoside analogue HIV reverse transcriptase inhibitor; PI, protease inhibitor; II, integrase inhibitor; FIB-4, noninvasive test for liver fibrosis based on AST/ALT ratio and platelet count.

**Table 2.** Summary of markers of T cell activation and plasma biomarkers of bacterial translocation, inflammation, and coagulopathy in healthy controls, HIV-monoinfected and HIV/HCV-coinfected patients.

	<b>Healthy controls (0)</b>	<b>HIV-monoinfected (1)</b>	<b>HIV/HCV-coinfected (2)</b>	<b>p (0-1)</b>	<b>p (0-2)</b>	<b>p (1-2)</b>
<b>T cells (%)</b>						
CD4 <sup>+</sup> CD38 <sup>+</sup>	3.7 (2.5; 5.8)	2.8 (2; 5.6)	6.8 (3.8; 14.7)	.206	<b>.000</b>	<b>.000</b>
CD8 <sup>+</sup> CD38 <sup>+</sup>	6.3 (5.1; 8.5)	7.1 (4.6; 10.7)	11.6 (7.1; 20)	.504	<b>.000</b>	<b>.001</b>
<b>Bacterial translocation</b>						
sCD14 (µg/mL)	3.3 (2.3; 3.9)	3.7 (1.9; 5.4)	5.1 (3.3; 7.4)	.370	<b>.000</b>	<b>.010</b>
FABP2 (ng/mL)	0.5 (0.3; 0.7)	0.6 (0.4; 1.3)	0.7 (0.3; 1.6)	.051	.050	.822
LPS (UE/mL)	1.3 (0.9; 4.7)	1.2 (0.9; 2.1)	1.4 (1; 1.9)	.654	.765	.483
LBP (µg/mL)	0.7 (0.2; 1.4)	0.8 (0.4; 1.4)	0.9 (0.6; 1.5)	.599	.088	.294
<b>Inflammation</b>						
IL-1b (pg/mL)	0.6 (0.2; 1.5)	0.6 (0.3; 1)	1.2 (0.5; 2.3)	.705	<b>.006</b>	<b>.000</b>
IL-8 (pg/mL)	1.2 (1.2; 2.8)	1.9 (1.2; 3.5)	5.2 (3.6; 11.3)	.197	<b>.000</b>	<b>.000</b>
IL-6 (pg/mL)	2 (1.4; 3.6)	3.4 (2.4; 4)	5.4 (3.7; 8.1)	<b>.014</b>	<b>.000</b>	<b>.000</b>
IL-18 (pg/mL)	87.4 (55.9; 147.3)	122.8 (87.9; 194.4)	247.7 (128.2; 507.2)	.077	<b>.000</b>	<b>.000</b>
IP-10 (pg/mL)	26.6 (20.1; 50.5)	28.4 (17.5; 35.4)	203.7 (111.3; 350.6)	.704	<b>.000</b>	<b>.000</b>
<b>Endothelial dysfunction</b>						
sVCAM1 (µg/mL)	0.3 (0.2; 0.5)	0.3 (0.2; 0.6)	1.6 (0.8; 3.2)	.154	<b>.000</b>	<b>.000</b>
sICAM1 (µg/mL)	0.4 (0.1; 0.9)	0.6 (0.3; 1.2)	2.1 (1.1; 3.9)	.084	<b>.000</b>	<b>.000</b>
sTNFR1 (ng/mL)	1.5 (0.2; 2.2)	1.4 (0.3; 2.1)	2.3 (1.3; 3.6)	.917	<b>.000</b>	<b>.000</b>
MCP1 (pg/mL)	12.5 (9.2; 21.7)	29.5 (17.7; 38.4)	20.7 (11; 41.2)	<b>.000</b>	<b>.005</b>	.173
<b>Coagulopathy</b>						
D-Dimer (ng/mL)	23.5 (10.8; 62.2)	28.2 (12.7; 50)	33.3 (12.4; 77.8)	.898	.293	.253
PAI-1 (ng/mL)	4.9 (3.7; 8.6)	6.8 (5.3; 8.9)	10 (7; 12.8)	<b>.035</b>	<b>.000</b>	<b>.000</b>

**Statistics:** Values expressed as median (interquartile range). *P*-values were calculated by Mann-Whitney tests.

**Abbreviations:** HCV, hepatitis C virus; HIV-1, human immunodeficiency virus type 1; CDXX, cluster of differentiation; sCD14, soluble CD14; LPS, lipopolysaccharide; FABP2, fatty acid-binding protein 2; LBP, lipopolysaccharide binding protein; IL, interleukin; IP-10, IFN-γ-inducible protein 10; sVCAM1, soluble vascular cell adhesion molecule 1; sICAM1, soluble intercellular cell adhesion molecule 1; sTNFR1, soluble tumor necrosis factor receptor 1; MCP1, monocyte chemoattractant protein-1, PAI-1, plasminogen activator inhibitor-1.

**Table 3.** Association of liver stiffness measurements (continuous variable) with markers of T cell activation and plasma biomarkers of bacterial translocation, inflammation, and coagulopathy in HIV/HCV-coinfected patients.

	AMR (95%CI)	<i>p</i>	aAMR (95%CI)	<i>p</i>
<b>T cells (%)</b>				
CD4 <sup>+</sup> CD38 <sup>+</sup>	0.9 (0.78;1.04)	.150	0.97 (0.83;1.14)	.734
CD8 <sup>+</sup> CD38 <sup>+</sup>	1.14 (1;1.29)	<b>.048</b>	1.15 (1.01;1.31)	<b>.037</b>
<b>Bacterial translocation</b>				
sCD14 (µg/mL)	1.04 (0.89;1.22)	.628	1.12 (0.93;1.3)	.226
FABP2 (ng/mL)	1.09 (0.9;1.3)	.376	1.14 (0.93;1.40)	.205
LPS (UE/mL)	1.19 (1.06;1.34)	<b>.003</b>	1.18 (1.04;1.34)	<b>.009</b>
LBP (µg/mL)	1.02 (0.89;1.18)	.742	1.03 (0.88;1.21)	.679
<b>Inflammation</b>				
IL-1b (pg/mL)	0.91 (0.75;1.11)	.355	1.01(0.81;1.24)	.975
IL-8 (pg/mL)	1.99 (1.72;2.3)	<b>.000</b>	2.02 (1.74;2.35)	<b>.000</b>
IL-6 (pg/mL)	1.06 (0.92;1.22)	.415	1.14 (0.98;1.34)	.089
IL-18 (pg/mL)	0.96 (0.78;1.2)	.743	1.04 (0.83;1.29)	.748
IP-10 (pg/mL)	1.34 (1.14;1.57)	<b>.000</b>	1.27 (1.07; 1.51)	<b>.006</b>
<b>Endothelial dysfunction</b>				
sVCAM1 (µg/mL)	1.5 (1.29;1.75)	<b>.000</b>	1.39 (1.18;1.65)	<b>.000</b>
sICAM1 (µg/mL)	1.14 (0.9;1.43)	.275	1.29 (1.01;1.66)	<b>.045</b>
sTNFR1 (ng/mL)	1.51 (1.3;1.75)	<b>.000</b>	1.48 (1.27;1.74)	<b>.000</b>
MCP1 (pg/mL)	1.15 (0.98;1.35)	.098	1.14 (0.96;1.36)	.139
<b>Coagulopathy</b>				
D-Dimer (ng/mL)	1.88 (1.54;2.3)	<b>.000</b>	1.71 (1.38; 2.12)	<b>.000</b>
PAI-1 (ng/mL)	1.05 (0.94;1.18)	.365	1.05 (0.93;1.18)	.429

**Statistics:** *P*-values were calculated by Generalized Linear Models (GLM) with a gamma distribution (log-link).

**Abbreviations:** HCV, hepatitis C virus; HIV-1, human immunodeficiency virus type 1; AMR, arithmetic mean ratio; aAMR, adjusted AMR; CDXX, cluster of differentiation; sCD14, soluble CD14; LPS, lipopolysaccharide; FABP2, fatty acid-binding protein 2; LBP, lipopolysaccharide binding protein; IL, interleukin; IP-10, IFN-γ-inducible protein 10; sVCAM1, soluble vascular cell adhesion molecule 1; sICAM1, soluble intercellular cell adhesion molecule 1; sTNFR1, soluble tumor necrosis factor receptor 1; MCP1, monocyte chemoattractant protein-1, PAI-1, plasminogen activator inhibitor-1.

**Table 4.** Summary of T cell activation and plasma biomarkers of bacterial translocation, inflammation, and coagulopathy in HIV/HCV-coinfected patients according to fibrosis/cirrhosis stages.

	<12.5 kPa (0)	12.5-25 kPa (1)	25-40 kPa (2)	>40 kPa (3)	<i>p</i> (0-3)	<i>p</i> (1-3)	<i>p</i> (2-3)
<b>T cells (%)</b>							
CD4 <sup>+</sup> CD38 <sup>+</sup>	7.5 (4.5; 17.6)	6 (3.3; 14.1)	5.3 (2.6; 7.6)	10.9 (5.5; 24.8)	.268	.080	<b>.015</b>
CD8 <sup>+</sup> CD38 <sup>+</sup>	11.2 (6.3; 18.7)	10.9 (7.1; 19.2)	12.7 (6.3; 17.7)	21 (10; 39.9)	<b>.005</b>	<b>.011</b>	<b>.013</b>
<b>Bacterial translocation</b>							
sCD14 (µg/mL)	4.8 (3.2; 7.1)	5 (3.2; 7.5)	4.9 (3.5; 6.6)	7 (4.4; 10.6)	.081	.154	.100
FABP2 (ng/mL)	0.6 (0.3; 1.4)	0.6 (0.3; 1.5)	1 (0.4; 2.2)	1.2 (0.4; 2.5)	.082	.149	.529
LPS (UE/mL)	1.3 (1; 1.7)	1.3 (0.9; 2)	1.7 (1.3; 2.2)	1.5 (1.1; 2.3)	.121	.245	.761
LBP (µg/mL)	0.9 (0.5; 1.4)	0.9 (0.5; 1.8)	0.8 (0.5; 1.3)	1.1 (0.8; 1.4)	.412	.914	.184
<b>Inflammation</b>							
IL-1b (pg/mL)	1.3 (0.7; 2.5)	0.9 (0.5; 1.9)	1.6 (0.5; 2.8)	0.5 (0.2; 2.1)	.087	.375	.192
IL-8 (pg/mL)	4.2 (2.8; 6.4)	5.6 (4.1; 10)	11.9 (6.3; 20.9)	18.8 (9.1; 31.5)	<b>.000</b>	<b>.000</b>	<b>.027</b>
IL-6 (pg/mL)	4.6 (3.3; 7.7)	5 (3.6; 6.9)	6.7 (4.6; 12.1)	9.1 (6; 20)	<b>.000</b>	<b>.000</b>	<b>.025</b>
IL-18 (pg/mL)	231.1 (121.2; 569.9)	301.4 (145.6; 612.4)	254.3 (129.5; 465.4)	254 (155.1; 377.7)	.828	.508	.787
IP-10 (pg/mL)	171.3 (110.4; 258.3)	189.2 (79.1; 329)	336.1 (146.2; 455)	407 (203.7; 689.5)	<b>.000</b>	<b>.003</b>	.283
<b>Endothelial dysfunction</b>							
sVCAM1 (µg/mL)	1.2 (0.7; 2.1)	2.1 (0.9; 4.7)	1.4 (0.9; 4.2)	3.5 (1.1; 5.5)	<b>.000</b>	.214	.079
sICAM1 (µg/mL)	1.9 (1; 3.3)	2.2 (1.1; 4.1)	2.4 (1.2; 4.2)	3.9 (1.1; 12.1)	.070	.126	.188
sTNFR1 (ng/mL)	1.9 (1.2; 3.3)	2 (1.2; 3.1)	2.6 (1.3; 5.4)	3.5 (2.3; 4.5)	<b>.003</b>	<b>.003</b>	.498
MCP1 (pg/mL)	18.2 (10; 39.3)	23.2 (12.6; 39.2)	25.3 (11.3; 43.5)	27.9 (10.6; 52)	.234	.750	.787
<b>Coagulopathy</b>							
D-Dimer (ng/mL)	26.6 (11.4; 64.7)	32.9 (13.6; 79.8)	23.5 (8.4; 68.2)	132.2 (59.5; 280.1)	<b>.000</b>	<b>.000</b>	<b>.002</b>
PAI-1 (ng/mL)	9.7 (7; 12.1)	8.8 (5.8; 12.8)	10.7 (7.7; 14.4)	13.2 (10.1; 15.5)	<b>.003</b>	<b>.004</b>	.121

**Statistics:** Values expressed as median (interquartile range). *P*-values were calculated by Mann-Whitney test.

**Abbreviations:** HCV, hepatitis C virus; HIV-1, human immunodeficiency virus type 1; kPa, kilopascal; CDXX, cluster of differentiation; sCD14, soluble CD14; LPS, lipopolysaccharide; FABP2, fatty acid-binding protein 2; LBP, lipopolysaccharide binding protein; IL, interleukin; IP-10, IFN-γ-inducible protein 10; sVCAM1, soluble vascular cell adhesion molecule 1; sICAM1, soluble intercellular cell adhesion molecule 1; sTNFR1, soluble tumor necrosis factor receptor 1; MCP1, monocyte chemoattractant protein-1, PAI-1, plasminogen activator inhibitor-1.

**Table 5.** Summary of values of adjusted arithmetic mean ratio (aAMR) for <12.5kPa, 12.5-25kPa and 25-40kPa, compared to HIV/HCV-coinfected patients with >40 kPa, for T-cell subsets and plasma biomarkers.

	<12.5 kPa		12.5-25 kPa		25-40 kPa	
	aAMR (95%CI)	<i>p</i>	aAMR (95%CI)	<i>p</i>	aAMR (95%CI)	<i>p</i>
<b>T cells (%)</b>						
CD4 <sup>+</sup> CD38 <sup>+</sup>	1.35 (0.89;2.07)	.159	1.55 (1;2.4)	.051	1.68 (1.03;2.75)	<b>.038</b>
CD8 <sup>+</sup> CD38 <sup>+</sup>	1.66 (1.16;2.36)	<b>.005</b>	1.49 (1.03;2.16)	<b>.036</b>	1.62 (1.06;2.49)	<b>.026</b>
<b>Bacterial translocation</b>						
sCD14 (µg/mL)	1.18 (0.77;1.81)	.449	1.05 (0.67;1.66)	.830	1.16 (0.71;1.9)	.548
FABP2 (ng/mL)	1.36 (0.82;2.26)	.230	1.47 (0.86;2.52)	.155	1.07 (0.6;1.91)	.824
LPS (UE/mL)	1.61 (1.18;2.17)	<b>.003</b>	1.45 (1.04;2.04)	<b>.027</b>	1.05 (0.74;1.49)	.788
LBP (µg/mL)	1.09 (0.76;1.56)	.653	1.02 (0.69;1.5)	.937	1.23 (0.8;1.89)	.338
<b>Inflammation</b>						
IL-1b (pg/mL)	0.74 (0.45;1.23)	.245	0.91 (0.53;1.58)	.745	0.52 (0.29;1.01)	.073
IL-8 (pg/mL)	3.35 (2.31;4.86)	<b>.000</b>	2.34 (1.55;3.54)	<b>.000</b>	1.33 (0.86;2.07)	.201
IL-6 (pg/mL)	1.65 (1.12;2.44)	<b>.012</b>	2.23 (1.46;3.4)	<b>.000</b>	1.69 (1.08;2.64)	<b>.022</b>
IL-18 (pg/mL)	0.77 (0.48;1.24)	.280	0.67 (0.4;1.13)	.135	0.74 (0.42;1.29)	.284
IP-10 (pg/mL)	1.78 (1.15;2.75)	<b>.010</b>	1.72 (1.06;2.8)	<b>.029</b>	1.04 (0.62;1.75)	.885
<b>Endothelial dysfunction</b>						
sVCAM1 (µg/mL)	2.14 (1.45;3.17)	<b>.000</b>	1.15 (0.75;1.77)	.515	1.3 (0.81;2.07)	.276
sICAM1 (µg/mL)	1.84 (1;3.35)	<b>.048</b>	2.26 (1.17;4.36)	<b>.015</b>	2.06 (1.02;4.19)	<b>.045</b>
sTNFR1 (ng/mL)	1.56 (1.08;2.26)	<b>.017</b>	1.64 (1.1;2.45)	<b>.015</b>	0.98 (0.63;1.52)	.913
MCP1 (pg/mL)	1.52 (0.99;2.32)	.056	1.23 (0.76;1.98)	.399	1.38 (0.82;2.31)	.228
<b>Coagulopathy</b>						
D-Dimer (ng/mL)	4.06 (2.32;7.11)	<b>.000</b>	4.34 (2.36;7.97)	<b>.000</b>	3 (1.55;5.8)	<b>.001</b>
PAI-1 (ng/mL)	1.23 (0.9;1.67)	.189	1.31 (0.94;1.84)	.112	1.13 (0.79;1.63)	.503

**Statistics:** *P*-values were calculated by Generalized Linear Models (GLM) with a gamma distribution (log-link).

**Abbreviations:** HCV, hepatitis C virus; HIV-1, human immunodeficiency virus type 1; aAMR, adjusted AMR; kPa, kilopascal; CDXX, cluster of differentiation; sCD14, soluble CD14; LPS, lipopolysaccharide; FABP2, fatty acid-binding protein 2; LBP, lipopolysaccharide binding protein; IL, interleukin; IP-10, IFN-γ-inducible protein 10; sVCAM1, soluble vascular cell adhesion molecule 1; sICAM1, soluble intercellular cell adhesion molecule 1; sTNFR1, soluble tumor necrosis factor receptor 1; MCP1, monocyte chemoattractant protein-1, PAI-1, plasminogen activator inhibitor-1.

**Supplemental Table 1.** Characteristics of HIV/HCV co-infected patients.

	<b>Healthy control</b>	<b>HIV</b>	<b>HIV/HCV</b>
<b>No.*</b>	32	39	238
<b>Gender (male)</b>	17 (53.1%)	24 (61.5%)	187 (78.6%)
<b>Age (years) #</b>	49.5 (47; 53)	51 (46; 53)	49 (46; 52)
<b>BMI (kg/m<sup>2</sup>) #</b>	24.9 (23.1; 27.1)	25.3 (23.5; 26.6)	24.4 (21.8; 26.9)
<b>BMI ≥25 (kg/m<sup>2</sup>)*</b>	14 (43.7%)	21 (55.3%)	98 (41.2%)
<b>Diabetes</b>	-	6 (15.8%)	20 (8.4%)
<b>High alcohol intake</b>	-	1 (3.1%)	117 (89.1%)
<b>HIV acquired by IVDU*</b>	-	-	186 (78.1%)
<b>Prior AIDS*</b>	-	13 (33.3%)	64 (26.9%)
<b>Years since HIV infection #</b>	-	-	23 (18; 26)
<b>Years since HCV infection #</b>	-	-	21 (16; 24)
<b>Antiretroviral therapy *</b>			
<b>Non treated</b>	-	-	5 (2.1%)
<b>PI-based#</b>	-	10 (25.6%)	35 (14.7%)
<b>2NRTI+II-based *</b>	-	4 (10.2%)	59 (24.8%)
<b>2NRTI+PI-based *</b>	-	-	47 (19.7%)
<b>2NRTI+NNRTI-based *</b>	-	23 (64.1%)	70 (29.5%)
<b>Others</b>	-	2 (5.1%)	22 (9.2%)
<b>HIV markers</b>			
<b>Nadir CD4+ T-cells #</b>	-	215 (107; 343)	172 (84; 254)
<b>Nadir CD4+ T-cells&lt;200 cells/mm<sup>3</sup>*</b>	-	14 (38.9%)	131 (55.1%)
<b>CD4+ T-cells #</b>	-	832 (685; 1036)	547 (394; 803)
<b>CD4+ T-cells&lt;500 cells/mm<sup>3</sup>*</b>	-	0 (0%)	100 (42%)
<b>HIV-RNA &gt;50 cp/mL*</b>	-	0 (0%)	30 (12.6%)
<b>Non-invasive indexes</b>			
<b>APRI #</b>	-	-	0.98 (0.57; 1.70)
<b>APRI &gt;1.5*</b>	-	-	65 (27.3%)
<b>FIB-4 #</b>	-	-	2.33 (1.45; 3.59)
<b>FIB-4 &gt;3.25*</b>	-	-	64 (26.9%)
<b>Forns index #</b>	-	-	4.94 (3.67; 6.31)
<b>Forns index &gt;6.9*</b>	-	-	36 (15.1%)
<b>HCV markers</b>			
<b>HCV genotype (n=235)*</b>			
<b>1</b>	-	-	170 (72.3%)
<b>2</b>	-	-	5 (2.1%)
<b>3</b>	-	-	39 (16.6%)
<b>4</b>	-	-	21 (8.9%)
<b>Log<sub>10</sub> HCV-RNA (IU/ml) #</b>	-	-	6.3 (5.87; 6.74)
<b>HCV-RNA &gt; 850,000 IU/ml *</b>	-	-	170 (71.4%)

\*Absolute number (percentage).

# Mean and mean standard error.

HCV: Hepatitis C virus; HCV-RNA: HCV plasma viral load; HIV-1: Human immunodeficiency virus type 1; HIV-RNA: HIV plasma viral load; IVDU: intravenous drug user; NNRTI: non-nucleoside analogue HIV reverse transcriptase inhibitor; NRTI: nucleoside analogue HIV reverse transcriptase inhibitor; PI: protease inhibitor; II: integrase inhibitor; APRI: Aminotransferase-to-platelet ratio index; FIB-4: noninvasive test for liver fibrosis based on AST/ALT ratio and platelet count; FORNS: test for liver fibrosis based on age, GGT and platelet count.