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Kupffer Cells are Depleted with HIV Immunodeficiency and Partially Recovered with Antiretroviral Immunereconstitution:

HIV, Kupffer Cells and Antiretroviral Therapy

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Introduction

In the modern era of highly active anti-retroviral therapy (HAART), a leading cause of death in HIV-infected persons is liver disease, most often due to chronic hepatitis C virus (HCV) infection [1]. Approximately one quarter of HIV infected persons have chronic hepatitis C [2], and HIV-HCV co-infected persons have demonstrated a more rapid progression of liver disease compared with HCV mono-infected persons. Although the mechanism(s) are not fully known, liver disease is more severe in those with more profound CD4+ lymphocyte depletion compared to those with preserved peripheral blood CD4+ lymphocyte counts [3– 8].

HIV induced depletion of intestinal CD4+ lymphocytes [9–12] has recently been linked to microbial translocation and HIV progression [13]. Moreover, in a cohort of HCV-infected persons at different stages of liver disease we have shown that microbial translocation in HIV co-infected persons was strongly associated with liver disease progression [14]. In that study we also found that HIV seroconversion amongst HCV-infected persons was associated with an increase in microbial translocation over time. It is, therefore, compelling to consider microbial translocation as a common mechanism that contributes to both HIV and HCV progression. Interestingly, animal models of other forms of liver disease have demonstrated a critical role of intestinal microbial translocation in promoting fibrosis [15–20].

Hepatic macrophages, or Kupffer cells, are responsible for clearing microbial translocation products and play a role in liver disease. Kupffer cells, however, can be infected by HIV, and this may result in their impaired ability to clear these potentially fibrogenic microbial translocation products [21–27]. In this investigation, we tested the hypotheses that Kupffer cell quantities are associated with peripheral CD4+ lymphocyte count in HIV-HCV co-infection, and that changes in CD4+ due to antiretroviral therapy are associated with corresponding alterations in Kupffer cell quantities. In addition, since in chronic viral hepatitis fibrosis begins in the portal and periportal regions where microbial translocation

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products first enter the liver, we tested the hypothesis that Kupffer cells would be most abundant in these regions.

Methods

The study population derives from the HIV-HCV co-infected members of the Johns Hopkins University clinical cohort (Baltimore, MD) [3;28]. Seventy-six individuals were identified who had at least two archived liver tissue samples and correlated clinical data characterizing HIV and HCV stage between January, 1997 and February, 2005. All subjects provided written informed consent for testing through a protocol approved by the Committees on Human Research of the Johns Hopkins School of Medicine or Bloomberg School of Public Health.

Data on clinical and lab parameters were abstracted from the clinical and laboratory databases. Transcutaneous liver biopsies were obtained using an 18-gauge needle. Liver tissue was fixed in 10% formalin and paraffin-embedded. Tissues were stained with hematoxylin and eosin (H&E) as well as trichrome. As previously described, tissues were scored by an experienced liver pathologist (M.T.) for fibrosis according to the Metavir scoring system for HCV infection and were graded for the degree of inflammation by using the Ishak modified hepatic activity index (MHAI) [29]. Hepatic fat (steatosis) was assessed as an average percentage of fat (0, 1-30, 31-60, > 60%) on H&E section. The pathologist was blinded to the subjects' clinical history and laboratory values. In a select group of subjects who were studied longitudinally, slides obtained from the same subject were separately encoded and de-identified before handling by the pathologist. Adequacy of tissue size was determined by the pathologist and subjects with inadequate tissues were excluded from the investigation. The median length of tissues was 12 mm. To detect Kupffer cells, tissue sections were immunostained following heat antigen retrieval with mouse monoclonal anti-CD68 antibodies (Dako, Carpinteria, CA), used at a 1:100 dilution. The DAKO EnVision+ Peroxidase kit was used for immunostaining. Kupffer cells were identified by their strong cytoplasmic staining. Kupffer cell density (KCD) was determined as the arithmetic mean number of Kupffer cells per 5 high power fields. Portal and periportal Kupffer cells were further quantified and compared with centrilobular Kupffer cells. [To detect CD68+/HLA-DRa+ Kupffer cells, paraffin-embedded tissue sections from 20 available liver blocks were selected from a subset of subjects matched for MHAI score who had liver biopsies obtained at a second time point. Blocks were deparaffinized and sequentially immunostained with the anti-CD68+ antibody followed by a rabbit polyclonal anti-HLA-DRa IgG antibody (Santa Cruz, Santa Cruz, CA), both at 1:100 dilution. Dual sequential immunostaining was performed using the EnVision G|2 Doublestain System (Dako, Carpinteria, CA). CD68+/ HLA-DRα-, and CD68-/ HLA-DRα+, and CD68+/HLA- $DR\alpha$ + cells were enumerated in 5–10 parenchymal high powered fields and averaged.]

Regional differences in KCD were characterized by quantifying all CD68+ Kupffer cells within a 50 µm radius of selected portal and centrilobular veins. A Zeiss PALM^R MicroLaser system was used to accurately determine a 50 µm radius around vessels. While this distance extends from portal to periportal regions depending on the section, it allows for precise comparisons between portal and centrilobular Kupffer cells. Cases were selected for analysis if they had at least one suitable complete cross section of both a well oriented portal tract and central vein. To avoid skewed results on the basis of oversampling regions of abundant inflammation, only 1 representative portal or centrilobular vein was used from a cluster of branching vessels. Multiple sequential sections were only used to confirm findings, and not used for counting to avoid oversampling the same cells. Kupffer cell number was normalized per µm² to account for differences in vessel size using the adapted area of an elliptical ring, $\pi * (\frac{1}{2} a + 50) * (\frac{1}{2} b + 50) - \pi * (\frac{1}{2} a) * (\frac{1}{2} b)$, where *a* and *b* were

the measured long and short axes of the vessel in μ m (Fig. 1). Kupffer cell counts of zero were conservatively assigned values of 0.5 per given area as the lower limit of detection. Results are reported in mm².

Liver biopsies were obtained a median (range) of 1.25 (0.1 - 139.6) months from the assessment of CD4+ lymphocyte count and clinical data, which were also ascertained using structured instruments before and after the initial liver biopsy, as described elsewhere [28]. KCD in all subjects was approximately normally distributed. Wilcoxon rank-sum and Kruskal-Wallis tests were used to compare median KCD values across different covariates of interest. Linear regression models were used to model the association between KCD and both current and nadir CD4+ lymphocyte count. Kupffer cell quantities per vessel were not normally distributed and were compared using Wilcoxon rank-sum tests. When log-transformed, Kupffer cell quantities per vessel yielded normally distributed data that was similarly found to be statistically significant using paired *t* tests.

Results

Seventy-six HIV-HCV co-infected persons with archived liver biopsies were identified. The mean age of the participants was 44.8 years at the time of the first biopsy, 59 (77.6%) were male, and 67 (88.2%) were African-American (Table 1). Peripheral blood CD4+ lymphocyte counts ranged from 9 to $1277/\text{mm}^3$ and serum HIV RNA ranged from undetectable to > 400,000 copies per mL. Antiretroviral therapy use was reported in 57 (75%) persons. All persons had chronic HCV infection with a median RNA level of 596,000 IU/mL. HCV infection was predominantly genotype 1a (66.2%) or 1b (25.4%).

Baseline liver fibrosis stage ranged from zero to cirrhosis (Metavir 0–4). Liver inflammation ranged from MHAI 0 to 9, and steatosis from 0 to 60%. KCD was normally distributed over the cohort, with a median (IQR) of 23 cells/HPF (17.7–27.8). No differences were detected in the KCD distribution according to age, gender, or HCV genotype (Table 1). Importantly, peripheral blood monocyte (Kupffer cell precursors) quantity was not associated with KCD (p=0.25). KCD was also not associated with liver disease fibrosis stage on the concurrent or subsequent biopsy, and was not associated with grade of hepatic inflammation or steatosis.

Mean KCD was significantly lower in subjects with lower peripheral blood CD4+ lymphocyte counts compared to those with higher levels (p<0.05, by Wilcoxon rank-sum test). KCD was associated with both contemporaneous CD4+ lymphocyte count and the lowest result recorded before histologic sampling (nadir CD4+ count, Table 2).

Since microbial translocation products enter the liver through the portal vein, we examined whether Kupffer cells were distributed more heavily in portal and periportal regions than centrilobular regions, as has been seen in rats [30;31]. From the 76 archived liver biopsies, 51 cases had sufficient tissue for additional analysis. There were no significant differences in the clinical characteristics between subjects with adequate liver tissue for the regional analysis and those with insufficient tissue. In total, Kupffer cells from 122 portal and 95 centrilobular regions were counted. The median (range) number of portal and periportal Kupffer cells, 328.6 (41.4 - 3252.8) /mm², was markedly higher than for centrilobular Kupffer cells, 90.5 (18.2 - 731.9) /mm² (Fig. 2A, B).

[Quiescent Kupffer cells are normally immunotolerant to physiologic amounts of microbial translocation. In order to determine if progression of AIDS was associated with Kupffer cell immune activation, 20 available liver blocks from equal numbers of subjects with high and low MHAI scores were dually immunostained for CD68 and the activation marker HLA-DR α . Single- and double-positive cells/HPF were enumerated in the hepatic parenchyma. Median KCD in this subgroup was lower than in the larger cohort (7.25 cells/HPF, range 1.8

-20 cells/HPF), and the median quantity of CD68+/HLA-DR α + was 4.15 cells/HPF(range 0 – 11.2 cells/HPF). HLA-DR α staining was found almost entirely on CD68+ cells; only a median of 0.1 (range 0 – 2.3) cells/HPF were CD68–/HLA-DR+, and this staining was confined to sinusoidal endothelial cells. There was a reciprocal trend between the number of dually-stained Kupffer cells per HPF and peripheral CD4+ lymphocyte count, but this was not statistically significant. Interestingly, the percentage of total Kupffer cells that were HLA-DR+ followed the same trend.]

The relationship between CD4+ lymphocyte count and KCD was examined further in a select subgroup of individuals who had immunosuppression (CD4+ lymphocyte count < $350/\text{mm}^3$) at the time of the initial biopsy, began HAART and achieved immune restoration (≥ 2 -fold increase in CD4+ lymphocyte count), then had a second biopsy with sufficient achieved tissue. In the eight evaluable subjects, biopsies were obtained at a median interval of 36.8 months (range 28.1–58.4 months). All 8 subjects had exposure to nucleoside reverse transcriptase inhibitors and protease inhibitors, and 3 subjects also had exposure to non-nucleoside reverse transcriptase inhibitors. The median (IQR) increase in CD4+ lymphocyte count was 246.5 (203–317) cells/mm³ and all subjects had virologic suppression on HAART to < 400 copies/mL. (Fig. 3) KCD went up in all subjects, from a median (IQR) of 16.6 (11 – 23.1) before HAART to a median (IQR) of 32 (24.6 – 39.2) during HAART exposure (*p*=0.007).

Discussion

Kupffer cells are the first line of defense against microbial translocation products that have been associated with liver fibrosis. In this investigation of HIV-HCV co-infected persons we have demonstrated that KCD is lowest in persons with the most AIDS-related immunosuppression compared to those with preserved CD4+ lymphocytes. In a subset of subjects who were given HAART and had immune reconstitution, KCD increased along with absolute CD4+ lymphocyte counts. These data suggest that Kupffer cell loss may contribute to liver fibrosis progression in persons with HIV-HCV co-infection and support observational studies linking HAART with slowed liver disease progression.

Immunosuppression due to HIV and other causes has been associated with HCV progression in numerous studies [3;6–8;32–35], and liver disease progression is slowed with the initiation of HAART [36;37]. There have been several explanations suggested for how HIV worsens HCV progression, and we propose that Kupffer cells may also play a role. Our data showing lower Kupffer cells in persons with the lowest CD4+ lymphocyte count, and the partial recovery of Kupffer cells with HAART and immune reconstitution suggest that the association between liver disease progression and HIV-related immunosuppression may be facilitated by Kupffer cells.

Activated Kupffer cells produce pro-inflammatory and pro-fibrogenic cytokines, such as TNF α and TGF β that in turn activate hepatic stellate cells, the proven precursors of liver fibrosis [38]. Animal models of liver injury of different causes have revealed from macrophage-depletion experiments that liver fibrosis is Kupffer cell-dependent [16;19;39–41]. Interestingly, these experiments have shown a critical role for microbial translocation in Kupffer cell-dependent fibrosis: portal vein-derived lipopolysaccharide (LPS) and its sensing apparatus on Kupffer cells (CD14, LPS-binding protein, and TLR4) are required for the development of liver disease [16;20]. While [activated] Kupffer cells can promote fibrosis in the setting of liver injury, in the quiescent host Kupffer cells are responsible for clearing microbial translocation products [42]. [Activation of Kupffer cells, as measured by increased HLA-DR expression, has been found in HCV mono-infection [43]. In this study we noted a trend suggesting that Kupffer cells are more frequently activated in persons with

the lowest CD4+ lymphocyte counts, though further work is necessary to evaluate differences between HIV-HCV co-infected and HCV mono-infected persons.] Our findings provide insights into how HIV might worsen chronic viral hepatitis.

Though it is not understood how Kupffer cell function is affected by HIV infection, earlier work has shown that HIV preferentially infects Kupffer cells over other hepatic cells [21–27]. Indeed, biopsies from HIV-infected humans and SIV-infected macaques have shown that Kupffer cells are enriched for HIV and SIV proteins, respectively [21–23;25]. Moreover, human and macaque Kupffer cells support productive HIV infection *in vitro* [24;26;27]. As a consequence, Kupffer cell loss may be due to the direct cytotoxic effects of HIV on the Kupffer cell, or as the result of soluble viral or host factors that induce programmed cell death. HIV infection may also alter the trafficking and migration of Kupffer cells and their precursors to target sites in the liver. Although we found no association between KCD and circulating monocyte quantities, recent data from SIV-infected macaques demonstrate that while circulating monocyte levels are unchanged from baseline in animals with AIDS, high monocyte turnover correlated tightly with AIDS progression and mortality [44].

This study also provides an explanation for the early development of periportal fibrosis in HCV infection despite clusters of HCV replication being diffusely found in the hepatic lobules. While injury and inflammation in chronic hepatitis C infection are found throughout the lobules [45], patterns of fibrosis in chronic viral hepatitis have long been identified as beginning in portal and periportal regions, supporting a role for intestinally-derived substances in enhancing fibrosis. In this study we found that Kupffer cells clustered next to portal veins where microbial translocation products first engage the liver, further strengthening an association between Kupffer cells, microbial translocation, and fibrosis and providing clues to the discordant locations of HCV replication and fibrosis in chronic HCV infection. Similar portal and periportal predominance of Kupffer cell shave been found in normal and diseased rats, but the implications of an intervessel Kupffer cell gradient are poorly understood [30;31]. In our study, since the majority of study subjects had low Metavir and MHAI scores, it is unlikely that portal and periportal Kupffer cell clustering is simply a reflection of non-specific infiltrating inflammatory cells.

In this study we did not include HCV mono-infected persons, a group that will be of importance in future studies of Kupffer cell dynamics in various mono- and co-infected populations. In addition, further studies are needed to show that HIV infection of Kupffer cells is directly responsible for Kupffer cell loss. Though we did not find an association between KCD and fibrosis, our study was enriched for persons at earlier stages of liver disease, with only 9 persons who had Metavir scores ≥ 2 at baseline and only 10 persons who had significantly progressed fibrosis between the two biopsies. Of note, this cohort derived from a larger cohort of HIV-HCV co-infected persons who had limited amounts of fibrosis at early time points [3]. Also, because of the cross-sectional nature of our study, the impact of past Kupffer cell activity on contemporaneous liver pathology may not be captured at the instant of biopsy. As a further limitation, we did not have more than one person independently assess KCD on all slides, although ten percent of slides were scored by a second investigator with high correlation between results.

Too little is known about the relationship of KCD and their physiologic function to interpret the magnitude of differences that were observed in this study. Although the dynamic range of Kupffer cell density was limited compared to CD4+ lymphocyte counts, the LPSclearance capacity of Kupffer cells is thought to be quite large. Indeed, after intravenous injection of LPS into rats, most was substantially cleared by the liver and nearly all of it was Our results suggest that Kupffer cells form a dynamic cellular population that is closely tied to peripheral CD4+ lymphocyte counts and localizes to portals of microbial translocation. Hepatic macrophages are a filter of intestinally-derived products, and dysregulated macrophage survival and/or trafficking in the HIV-HCV-coinfected host may be one mechanism by which liver fibrosis rapidly progresses. Further studies into the mechanisms and consequences of HIV infection of Kupffer cells as well as the physiologic effect of Kupffer cell depletion on microbial translocation will need to be explored in *in vitro* and animal models.

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Abbreviations

| BD | bile duct |
|-------|--|
| CLV | centrilobular vein |
| HA | hepatic artery |
| HAART | highly active anti-retroviral therapy |
| HCV | hepatitis C virus |
| HIV | human immunodeficiency virus |
| IQR | interquartile range |
| KCD | Kupffer cell density |
| LPS | lipopolysaccharide |
| NNRTI | non-nucleoside reverse transcriptase inhibitor |
| NRTI | nucleoside reverse transcriptase inhibitors |
| PI | protease inhibitor |
| PV | portal vein |
| SIV | simian immunodeficiency virus |
| TGFβ | transforming growth factor beta |
| TLR4 | toll-like receptor-4 |

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Fig.1.

PV - portal vein; BD - bile duct; HA - hepatic artery; KC - Kupffer cell. Shown is a representative picture of a portal triad, taken with a 20X light microscope. Kupffer cells are CD68 stained and appear brown on a counterstained blue background. A 50 µm designation was used to quantify Kupffer cells in an elliptical donut surrounding an hepatic vessel. Kupffer cells/mm² were defined by (No. of Kupffer cells*1000) / [π (½**a**+50)(½**b**+50)- π (½**a**)(½**b**)].



Fig. 2.

A. PV - portal vein; BD - bile duct; HA - hepatic artery; CLV - centrilobular vein; KC - Kupffer cell. Shown are representative pictures of a portal triad (upper panel) and a centrilobular vein (lower panel) from the same individual, taken with a 20X light microscope. Kupffer cells are CD68 stained and appear brown on a counterstained blue background. *B*. Biopsies from 51/76 subjects with \geq 1 postal vein and \geq 1 centrilobular vein were evaluated for a total of 122 portal and 95 centrilobular veins. Kupffer cells were quantified as described. Groups were compared using Wilcoxon rank-sum tests.



Fig. 3.

Eight HIV-HCV co-infected persons with at least 2 liver biopsies were selected for their initiation and response to HAART between biopsies. Biopsies were obtained at a median interval of 36.8 months apart (range 28.1 – 58.4 months). Response to HAART was defined as having a CD4+ lymphocyte count at the time of the second biopsy that was $\geq 2 \times$ the value at the time of the first biopsy (left upper panel). All subjects had virologic suppression of serum HIV RNA (right upper panel hashed reference line indicates 400 copies viral RNA/mL). KCD was measured as described on the first and second biopsies (lower panel). Symbols denote the same individuals over time and are consistent between the upper and lower panels. Because persons were preselected for increases in CD4+ lymphocyte count on HAART, a p-value was not generated for the top panel.

*comparison of mean KCD on HAART with mean KCD pre-HAART using Student's *t* test. † This individual, distinct in having the smallest increase in KCD during HAART, also had the highest pre-HAART CD4+ lymphocyte count and HIV RNA.

Table 1

Baseline characteristics of 76 HIV-HCV co-infected subjects.

| Characteristics | N (%) | Median KCD | IQR | p value ^f |
|-----------------------|-----------------|------------|-------------|----------------------|
| Age ^a | | | | |
| <40 years | 17 (22.4) | 24.4 | 17.4 - 30.2 | 0.48 |
| 40 - 50 years | 40 (52.6) | 23 | 18 - 26.8 | - |
| \geq 50 years | 19 (25) | 21.2 | 16. – 26.2 | - |
| Gender | | | | |
| Male | 59 (77.6) | 22.8 | 17.4 – 27.6 | 0.48 |
| Female | 17 (22.4) | 24.4 | 18.4 – 29.6 | - |
| Race | | | | |
| Black | 67 (88.2) | 23.4 | 18 - 28.6 | 0.25 |
| All others | 9 (11.8) | 20.8 | 14.2 - 23 | - |
| HCV Genotype | | | | |
| 1a | 47 (66.2) | 23 | 17.4 – 29.4 | 0.76 |
| 1b | 18 (25.2) | 23.2 | 19 - 24.8 | - |
| CD4+ T cell cou | nt ^b | | | |
| < 350/mm ³ | 39 (51.3) | 20.8 | 14.2 - 24.7 | 0.03 |
| ≥ 350/mm3 | 37 (48.7) | 24.4 | 21.2 - 28.6 | - |
| HIV Viral RNA | 2 | | | |
| < 400 cp/mL | 42 (55.3) | 23 | 18 - 26.2 | 0.51 |
| ≥ 400 cp/mL | 34 (44.7) | 23.1 | 14.4 - 30.4 | |
| HAART ^d | | | | |
| Yes | 53 (69.7) | 22 | 14.4 - 27.6 | 0.10 |
| No | 23 (30.3) | 24.4 | 19 – 31 | - |
| Metavir ^e | | | | |
| < 2 | 67 (88.2) | 23.2 | 18-28 | 0.63 |
| ≥ 2 | 9 (11.8) | 20.8 | 16.2 - 27.4 | - |
| MHAI ^e | | | | |
| ≤ 5 | 61 (80.3) | 22.4 | 18 - 25.2 | 0.28 |
| > 5 | 9 (11.8) | 25.7 | 18.4 - 32.2 | - |

HAART - highly active anti-retroviral therapy.

 a Age is given at the time of the first biopsy.

 b CD4+ lymphocyte determinations were made within a median (range) of 1.25 (0.1 – 139.6) months of the first biopsy.

^cHIV viral RNA was determined within a median (range) of 1.50 (0.1 – 68.9) months of the first biopsy.

 d HAART was defined as use of at least 2 nucleoside reverse transcriptase inhibitors (NRTI) + 1 protease inhibitor (PI), or 2 NRTIs +1 non-nucleoside reverse transcriptase inhibitor (NNRTI), or 2 NRTIs + 1 NNRTI.

 e Metavir and MHAI staging were determined by liver biopsy by a trained pathologist (M.T.)

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 ${}^f_{\ p}$ values were generated from Wilcoxon rank-sum and Kruskal-Wallis tests.

Table 2

Independent association of Kupffer cell density with CD4+ lymphocyte count.^a

| | Parameter Estimate | p Value |
|--------------------------------------|--------------------|---------|
| CD4+ T cell count b | .0114 | 0.027 |
| Nadir CD4+ T cell count ^C | .0164 | 0.006 |

 a Shown are univariate models of Kupffer cell density.

 b CD4+ lymphocyte determinations were made within a median (range) of 1.3 (0.1 – 139.6) months of the first biopsy.

 c Nadir CD4+ lymphocyte count was determined within a median (range) of 17.3 (0.1 – 139.6) months before the first biopsy, and was defined as the lowest recorded CD4+ lymphocyte count available.