Clinical and Virologic Characteristics of HIV-1 Positive Patients with Delta Hepatitis

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Abstract

Background and Aim: Hepatitis Delta Virus (HDV) infection has been mainly studied in HIV negative patients, while data on HIV-1 positive patients are limited. We investigated the virological pattern and as well as biochemical and clinical features of liver disease and immune status in HIV-1 positive patients with delta hepatitis. Their clinical characteristics were compared with those of anti-HDV negative, hepatitis B surface antigen (HBsAg) positive/HIV+ patients.

Methods: This retrospective study included HBsAg positive subjects with anti-HDV serology available, during the period 2010-2017. Biochemical and virological parameters were obtained at last visit in 2017 for each patient. Potential determinants for HDV positivity were examined by applying multivariate regression model.

Results: Of 78 HBsAg positive patients 19 (24.4%) were found anti-HDV+. Anti-HDV+ patients were more frequently intra venous drug users, anti-HCV positive and HBV e antigen (HBeAg) negative. Additionally, the patients had more severe liver disease and necro inflammatory activity (assessed by transient elastography and transaminase levels, respectively) than the counterpart of anti-HDV- patients. A suppressive effect of HDV over HCV was also revealed in anti-HDV+ subjects. By multivariate analysis, years of ART (OR 1.22; CI 0.986-1.43, p=0.014) and sexual exposure vs. IVDU (OR 0.08; CI 0.556-0.986, p=0.004) were independently associated with anti-HDV positivity.

Conclusion: Our data underlines the need for continuing prevention program that includes HBV vaccination, screening and monitoring in population at high risk, as well as development of an alternative treatment option for HDV.

Keywords: HBV; HDV; HCV; HIV-1; Replication; Fibrosis degree, Interaction

Abbreviations: HDV: Hepatitis Delta Virus; HBsAg: Hepatitis B Surface Antigen; HBeAg: Hepatitis B e Antigen; ART: Antiretroviral Therapy; Anti HCV: Antibodies Against Hepatitis C Virus; 3TC: Lamivudine; FTC: Emtricitabine; TDF: Tenofovir; IQR: Interquartile Range; CI: Confidence Interval; ALT: Alanine Aminotransferase; AST: Aspartate Aminotransferase; OR: Odds Ratio; MSM: Men Who have Sex with Men; IVDU: Intravenous Drug Users

Introduction

Hepatitis D virus (HDV) or delta hepatitis is a defective single stranded RNA virus that requires hepatitis B surface antigen envelope (HBsAg) for replication and transmission [1]. HDV infection is more frequent in HBsAg positive HIV infected patients, than in the counterpart of HIV negative patients. Additionally, HDV infected patients are more likely persons who inject drugs and a positive hepatitis C virus (HCV) serology compared to HDV-uninfected ones [2,3]. Studies [4,5] show that patients with double HBV and HDV infection have more severe liver disease, more rapid progression to cirrhosis and increased risk of hepatocellular carcinoma and decompensated liver cirrhosis. Thus, HDV infection represents an important health burden especially in high risk group of patients. However, there is only limited information on the clinical and virologic characteristic of Delta hepatitis in HIV-1 infected patients. Therefore, the aim of the present study was to investigate possible factors associated with delta hepatitis by evaluation of clinical and virologic characteristics in HBsAg+/HIV-1+ patient with or without serological markers of delta hepatitis.

Methods

This retrospective observational study examined demographic, clinical, therapeutic information and laboratory data of HIV-1/HBsAg positive patients who were tested for anti-HDV between January 2010 and December 2017. Clinical data including biochemical parameters, immunological status, HIV-RNA, HBV-DNA, and serological markers of HBV infection were collected for each HIV-1 infected patient at last visit available in 2017. The degree of fibrosis was assessed by transient elastography and was considered the last measurement performed in 2017. HBV-DNA and HCV-RNA quantitative assays were performed by real time PCR [Abbott diagnostics Illinois, USA] limit of sensitivity 10 IU/mL, and 12 IU/mL, respectively]. HDV-RNA qualitative assay was performed by home-made RT-PCR [6]. Briefly, total RNA was extracted from 200 μl of each plasma sample using QIAamp Viral RNA kit (Qiagen, Hilden, Germany). After reverse transcription with outer antisense primer D3, 5 μL cDNA was amplified with10 pmoles of outer primers encompassing the HDVag of the HDV genome: outer sense primer D0, 5' AGTGGCTCTCCCTTAGCCAT-3'(nt 813-832); outer antisense primer D3, 5'- TGAACCCCCTCGAAGGTGGA-3'(nt 1147-1128); inner sense primer D1, 5'- GTCCTCCTGGATGCCCCAG-3'(nt 847-866); and inner antisense D2, 5'- GAGTCCCCGAGTCCCCCCT-3'(nt

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Results for continuous variables were reported as median [interquartile range (IQR)]. Characteristics of HIV-1 positive patients were compared using the Pearson's chi-square or Fisher's exact test for categorical variables and the Mann-Whitney test for continuous variables. Potential determinants for HDV positivity were examined by applying multivariate regression model. Independent variables have been included in the stepwise log-linear model when p<0.10 and results were reported as odds ratios (OR) with their corresponding 95% confidence interval (CI). All statistical tests were two-sided at the 5% level (p ≤ 0.05) and were performed using SPSS statistical software, version 22 (IBM SPSS Statistics, IBM Corporation, Chicago, IL, USA).

### Results

Among 78 HBsAg+ HIV-1 infected patients tested for anti-HDV for whom were available clinical data, 59 were anti-HDV negative (HDV-), 7.5% were anti-HDV positive (HDV+), 22% were anti-HDV positive (HDV+). Clinical characteristics of HDV+ patients and HDV- patients are depicted in Table 1. By univariate analysis it was shown that the male gender was a risk factor for HDV infection (p=0.001). Similarly, a higher degree of fibrosis (assessed by transient elastography according to metavir score) was found in HDV+ patients when compared to HDV- patients (p=0.020). HDV+ patients were less often HBeAg positive than HDV- patients (5.9% and 43.6%, respectively, p=0.004). No difference in HIV-RNA and HBV-DNA (evaluated by qualitative and quantitative assays) was found between these two groups. Of note, all patients except one were exposed to ART including drugs against HIV/HBV [lamivudine (3TC), emtricitabine (FTC) or FTC plus tenofovir (TDF)]. Antibodies against HCV (anti-HCV) were more frequently detected in HDV+ than in HDV- patients (68.4% and 22.0%, of anti-HCV positivity respectively, p<0.001). On the contrary, HCV-RNA was more frequently found positive in HDV- than in HDV+ patients (HCV-RNA was detectable in 53.8% of anti-HCV+/anti-HDV- and 7.7% of anti-HCV+/anti-HDV+ patients, p=0.030). In anti-HCV+ patients with undetectable HCV-RNA, data on HCV-RNA were retrieved from clinical records and confirmed negative in a mean of 6 (range 2-9) consecutive specimens.

HDV-RNA qualitative assay was available in 7/19 anti-HDV+ patients and was positive in 4/7 (57.1%) patients. In 3 patients found HDV-RNA positive, the assay was repeated in 3 subsequent serum samples and negative results were obtained.

### Table 1: Characteristics of HIV-1/HBsAg positive patients with or without HDV.

<table>
<thead>
<tr>
<th>Variables</th>
<th>Overall</th>
<th>HDV+ No pts=19</th>
<th>HDV- No pts=59</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gender males/females</td>
<td>71/7</td>
<td>17/2</td>
<td>54/5</td>
<td>1.000</td>
</tr>
<tr>
<td>Age</td>
<td>51.0 (48.0-55.3)</td>
<td>53.0 (50.0-55.0)</td>
<td>51.0 (46.0-56.0)</td>
<td>0.151</td>
</tr>
<tr>
<td>Risk factor for HIV-1</td>
<td>17/39/22</td>
<td>11/4/4</td>
<td>6/35/18</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Years of HIV infection</td>
<td>21.2 (14.4-27.8)</td>
<td>25.5 (23.0-30.5)</td>
<td>18.5 (11.5-25.7)</td>
<td>0.002</td>
</tr>
<tr>
<td>Years of ART</td>
<td>17.0 (9.1-20.7)</td>
<td>20.6 (16.9-23.7)</td>
<td>14.7 (7.3-19.4)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>CD4 T cells number/μm³</td>
<td>698 (426-857)</td>
<td>535 (245-854)</td>
<td>750 (482-867)</td>
<td>0.151</td>
</tr>
<tr>
<td>ALT* IU/L</td>
<td>34 (26-56)</td>
<td>51 (26-88)</td>
<td>32 (22-46)</td>
<td>0.021</td>
</tr>
<tr>
<td>Transient elastography (KPa)</td>
<td>No. pts=58</td>
<td>9.6 (7.2-14.1)</td>
<td>9.5 (3.8-8.3)</td>
<td>0.001</td>
</tr>
<tr>
<td>Fibrosis degree ^</td>
<td>4/1/17</td>
<td>9/9</td>
<td>3/28</td>
<td>0.020</td>
</tr>
<tr>
<td>F0-F2 vs F3-F4</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>HBeAg-pos/neg (N72)</td>
<td>25/47</td>
<td>1/16</td>
<td>24/31</td>
<td>0.004</td>
</tr>
<tr>
<td>HIV-RNA* Positive/Negative</td>
<td>7/71</td>
<td>3/16</td>
<td>4/55</td>
<td>0.352</td>
</tr>
<tr>
<td>HIV-RNA load, copies/mL</td>
<td>164 (60-5569)</td>
<td>352 (-)</td>
<td>112 (59-4218)</td>
<td>0.400</td>
</tr>
<tr>
<td>HBV-DNA pos/neg (N74)</td>
<td>19/55</td>
<td>No pts.=17</td>
<td>No pts.=57</td>
<td>1.000</td>
</tr>
<tr>
<td>HBV-DNA load IU/mL</td>
<td>10 (10-15)</td>
<td>10 (10-42)</td>
<td>10 (10-19)</td>
<td>0.754</td>
</tr>
<tr>
<td>Anti-HCV pos/neg</td>
<td>26/52</td>
<td>13/6</td>
<td>13/46</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>HCV-RNA pos/neg</td>
<td>8/18</td>
<td>1/12</td>
<td>7/6</td>
<td>0.030</td>
</tr>
<tr>
<td>HCV-RNA load IU/mL</td>
<td>114527 (401-446430)</td>
<td>173575 (-)</td>
<td>55478 (362-1830624)</td>
<td>1.000</td>
</tr>
<tr>
<td>FTC or 3TC/TDF+FTC**</td>
<td>12/65</td>
<td>8/10</td>
<td>4/55</td>
<td>0.001</td>
</tr>
</tbody>
</table>

Abbreviations: IVDU: Intra Venous Drug Users; Pos: Positive; Neg: Negative; ALT: Alanine Aminotransferase; FTC: Emtricitabine; 3TC: Lamivudine; TDF: Tenofovir. Results are expressed as median (interquartile range, IQR). P-values according to Mann Whitney test or Chi-square/Fisher’s exact test, as appropriate

^ALT (normal value < 35 IU/L)

^Fibrosis degree was evaluated in patients for whom was available transient elastography

^HIV-RNA positive > 50 copies/mL

^IQR was not calculated because only 3 patients had quantifiable HIV load among anti-HDV+ patients.

**IQR was not calculated because only 1 HDV+ patient had a quantifiable HCV load.

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1084-1066). Nested PCR was performed on 2 μL of the first amplification products as a template with 10 pmol of the inner primer set.

In some cases the test was repeated at regular intervals. Patients previously treated with interferon or direct acting antivirals for HCV were excluded from this analysis. The data were retrieved from the database of the Division of Infectious Diseases of the San Raffaele Hospital (CSLHIV Cohort) Milan, Italy. The CSLHIV Cohort was approved by the ethics committee of the San Raffaele Hospital. At their first visit the patients provide written informed consent for scientific analyses of their data.
samples with an alternance of HDV-RNA positivity or negativity. Finally, HDV- patients were more frequently under ART regimen including 2 drugs against HIV/HBV than the counterpart of HDV+ patients (6.8% vs. 44.4%; p=0.001). By multivariate analysis, years of ART (OR 1.22; CI 0.986-1.43, p=0.014) and sexual exposure vs. IVDU (OR 0.08; CI 0.536-0.986, p=0.004) were independently associated with the risk to be anti-HDV positive. Finally, we evaluated the impact of HCV co-infection on the severity of liver disease in HDV+ patients. We found that anti-HCV positivity was not associated with a higher fibrosis score in HDV+ patients (among HDV+/anti-HCV+, 4 patients had F0-F2 and 8 patients had F3-F4; among HDV+/anti-HCV-, 3 patients had F0-F2 and 3 other patients had F3-F4; p=0.626).

Discussion

We found a high frequency (24.4%) of anti-HDV positivity in HIV/HBsAg+ patients attending our Division of Infectious Diseases in Milan, Italy. By univariate analysis, we found that HDV+ patients were more frequently IVDU respect to anti-HDV- patients and had a longer duration of HIV infection as well as longer duration of ART respect to the counterpart of HDV-. Reports from Europe [7,8] showed a lower prevalence of HDV infection in HIV-1 positive patients, while studies from Taiwan [9,10] reported dramatically high prevalence of HDV among HIV-1 patients/IVDU. So, our data likely reflects the high proportion of IVDU as main risk factor among HDV+ subjects.

A previous study [11] showed that HDV+ patients had more severe liver disease when compared to HDV uninfected persons. Accordingly, we found that among individuals with available data concerning the fibrosis degree, 9/18 (50.0%) HDV+ subjects had evidence of severe liver disease; while in HDV- patients only 8/40 (20.0%) had fibrosis score between F3-F4. However, these patients had a longer duration of HIV infection that could justify the higher degree of fibrosis with respect to HDV- patients. However, these patients had also a longer duration of ART, when compared to those not infected by HDV that could be of benefit at least on double HIV/HBV infection.

Studies [12-14] showed that treatment with nucleoside or nucleotide analogues (NA) were not effective against HDV in HIV-1 negative/ HBsAg positive patients. Soriano et al. [15] showed that ART treatment including tenofovir (TDF) worked well in HIV/HBV/HDV triple infected patients. More recently, Beguelin et al. [16] showed no activity of TDF on HDV replication in HIV/HBV/HDV triple infection. In the present study we did not specifically investigate the effect of treatment with NA on HDV. However, our findings of more advanced liver disease and higher necroinflammatory activity (assessed by transient elastography and ALT levels, respectively) in HDV+ patients respect to HDV- patients, suggested that ART including NA was not effective on HDV. We found that in HDV+ patients, (who were more frequently anti-HCV positive) the fibrosis score was similar in HDV+/anti-HCV+ and HDV+/anti-HCV- patients, suggesting that the severity of liver disease was likely consequent to HDV infection “per se” rather than to HDV/HCV co-infection.

Of note, the majority of HDV- patients had positivity for HBeAg, while near all HDV+ patients were HBsAg negative. It is well known that the majority of HDV+ subjects in Europe are positive for anti-HBe antibodies and negative for HBeAg. This phenomenon has been explained by epidemiological features, as HDV has mainly been studied in Mediterranean basin, where most hepatitis B patients are HBeAg-negative [17,18]. However, our patients group (including HDV+ and HDV- patients) had the same geographic origin (data not shown). One other possibility for this finding is suppressive effect of HDV on HBV replication.

No difference in HBV-DNA levels or HBV-DNA qualitative assay was found between HDV+ and HDV- patients. All these patients, apart from one, were under ART including drugs against HBV/HIV and the majority of patients had very low levels or undetectable HBV-DNA. Therefore, it was not possible to evaluate a suppressive effect of HDV on HBV replication. We also found a higher frequency of HCV infection assessed by anti-HCV positivity in HDV+ patients with respect to HDV- patients, this finding likely reflects the different efficiency of HCV transmission comparing IVDU vs. sexual exposure, which was consistent with previous studies [10,11,19,20].

The complex interplay between multiple hepatitis infections has been described in previous reports which generally showed HDV to be the dominant virus leading to the suppression of HBV as well as of HCV [21-23]. Although we did not evaluate in all patients the presence of HDV-RNA, we showed that the majority of anti-HDV+ patients had a positive serology for HCV without evidence of HCV replicating virus. This finding suggests a strong inhibitory effect of HDV on HCV; however, the mechanism behind this inhibitory effect remains unknown. By multivariate analysis, the major determinants of HDV infection were IVDU and the years of ART. To the best of our knowledge, this is the first report showing an association between years of ART and HDV infection in HIV-1 subjects, suggesting that ART including or not drugs active on HIV/HBV is not protective against HDV infection. Longer duration of ART and concomitantly longer duration of HIV infection in HDV+ patients respect to HDV- patients may also reflect a longer period of exposure to the virus.

Conclusion

Our study was limited by the nature of cross-sectional investigation and small sample size. So, it is possible that a number of HBsAg+ patients were not tested for anti-HDV or their data were not reported in the database. In addition, HDV-RNA assay was not performed in all anti-HDV positive patients. Therefore, we considered as HDV infected those patients with anti-HDV positivity. In summary, we compared clinical and virologic characteristics of HBsAg+ patients with or without HDV, some of whom were also HCV coinfected. We confirmed the severity of liver disease by a non-invasive method for assessing liver fibrosis and added information on demographic, immunological and virologic features of HIV-1/HDV+ patients, that could be taken in consideration for the management of this difficult to treat group.

Conflict of Interest

Morsica G, Peano L, Bagaglio S, Poli A, Hasson H, Messina M, Uberti-Foppa C have declared that no competing interests exist.

References


