Changes to hepatocyte ploidy and binuclearity profiles during human chronic viral hepatitis

H Toyoda, O Bregerie, A Vallet, B Nalpas, G Pivert, C Brechot, C Desdouets

Background and aim: The importance of the hepatocyte ploidisation pattern to the control of cell proliferation and differentiation has been well established. However, there are no data that have characterised hepatocyte ploidy at various stages of chronic liver inflammation and fibrosis in vivo.

Methods: We therefore investigated hepatocyte ploidy/binuclearity patterns in 57 patients with chronic hepatitis, using a recently developed methodology which allows simultaneous hepatocyte ploidy and binuclearity analyses on the same liver section.

Results: The percentage of mononuclear diploid hepatocytes was significantly reduced in patients with high hepatitis activity and marked fibrosis (low activity: 75.1 (18.8%) v high activity: 61.8 (21.6%), p = 0.0111, and low fibrosis: 77.3 (13.8%) v high fibrosis: 57.4 (23.3%), p = 0.0002). Accordingly, the percentage of mononuclear polyploid hepatocytes increased in patients with high hepatitis activity and marked fibrosis (low activity: 11.9 (15.5%) v high activity: 22.2 (20.1%), p = 0.0166, and low fibrosis: 9.4 (10.7%) v high fibrosis: 26.4 (21.6%), p = 0.0001). In addition, the fraction of binuclear hepatocytes was significantly higher in patients with hepatitis B virus (HBV) than in those with hepatitis C virus (HCV) infections (HBV: 18.2 (7.6%) v HCV: 12.0 (4.8%), p = 0.0020). Under multivariate analysis, HBV infection was an independent factor accounting for the larger binuclear hepatocyte fraction (p = 0.0294).

Conclusion: Our results revealed an increase in the polyploid hepatocyte fraction which correlates with the severity of chronic hepatitis; moreover, we demonstrated that HBV and HCV related chronic hepatitis exhibited distinctive hepatocyte ploidy patterns, thus allowing the suggestion that these two viral infections may modulate liver ploidy through different mechanisms.
between March 1996 and March 2001. Twenty four patients were chronically infected with hepatitis B virus (HBV) while the remaining 33 patients were infected with hepatitis C virus (HCV). HBV patients had detectable hepatitis B surface antigen. At the time of liver biopsy, 11 patients had negative HBV DNA and 13 had positive serum HBV DNA (eight were infected by pre-C mutant virus and five by wild-type virus) using a hybridisation test (cut off value 700 000 copies/ml). There was no evidence of occult HBV infection in HCV patients.

All patients included in this study had no evidence of alcohol related liver disease. Based on histological findings, patients were grouped as a function of the degree of activity of hepatitis and degree of fibrosis, according to the METAVIR scoring system.21 The study was approved by the institutional review board for Hopital Necker Enfants Malades (Paris, France) and was carried out in accordance with the Helsinki declaration. Written informed consent was obtained from all patients before liver biopsy and prior to the study.

Immunofluorescence microscopy: definition of liver DNA ploidy
Liver tissues fixed in 10% phosphate buffered formalin were embedded in paraffin. Tissue sections (3 μm thick) were obtained using a conventional microtome. Sections were deparaffinised in xylene and placed in 100% ethanol. Sections were rehydrated in a descending gradient of ethanol-water and then boiled for 3 × 5 minutes in 10 mmol/l citrate buffer, pH 6.2, in a microwave oven. All subsequent antibody incubations were carried out in an antibody diluent (Dako, Trappes, France). Incubation with the primary antibody (anti-cytokeratin 1-50; KL1, Immunotech, SA, Marseille, France) was carried out for one hour at room temperature, followed by three washes with phosphate buffered saline (PBS). A secondary reagent was used to detect the primary antibody: biotinylated swine anti-goat, mouse, rabbit immunoglobulin solution (1:200; Dako) (30 minutes’ incubation) followed by FITC conjugated streptavidin (1:200; Dako) (30 minutes’ incubation). Hoechst 33342 (1 μg/ml) was included in the last incubation to stain DNA (20 minutes’ incubation). Following three final washes with PBS, the slides were mounted in 100% glycerol.

Tissue sections were then examined under a Zeiss (Axiovert 35; Carl Zeiss, Gottingen, Germany) inverted microscope equipped for epi-illuminations. Zeiss Plan Neofluar objectives 40 × (NA = 0.75) were chosen, enabling the collection of light from the entire thickness of the nucleus, these conditions being essential for adequate determination of DNA content.22 Images were captured using a cooled charged coupled device camera (KAF 1400-G2, class 2; Photometrics, Tucson, Arizona, USA) on 4056 grey levels. Automatic quantitative image analysis was performed in 12 bits using the IPLab Spectrum version 3.1 software. For measurement of nuclear DNA content on the image, we carefully selected only nuclei with a clear edge in order to avoid the effect of diffraction on the evaluation. Nuclei were termed mono- or binucleated hepatocytes by comparing fluorescent and membrane labelling images. Other liver cell types (defined by their morphological characteristics) overlapping nuclei or debris were eliminated interactively by inspection on the screen, using zoom and contrast enhancement where necessary. Parameters such as integrated fluorescent intensity or area of nucleus were stored in computer files for analysis using the Kaleidagraph and Imastat software (Imastat as developed by P Debe) 20 22 A minimum of 300 hepatocytes was studied, observed on 8–12 separate fields (analyse of at least two sections). Fields were randomly selected but always contained the same number of perportal, centrolobular, and mediolobular area from one patient to another. In order to eliminate fluorescence intensity value due to nuclear truncature, a control slide consisting on a pure 2n hepatocyte population was include in each experiments. This slide allowed us to determine the fluorescence intensity values for 2n DNA content; lower values (corresponding to nuclear truncature) were not taken into account. To ensure that the same cells were not counted twice, slides were read by two observers in a systematic manner, moving from the right to the left of the slide and then on successive descending lines. All analyses were performed blind.

Statistics
Results are expressed as mean (SD). Differences in proportions were tested by the χ² test. Mean quantitative values were compared using the non-parametric Mann-Whitney U test. Correlations were evaluated using Spearman’s test. Multivariate analysis was performed using logistic regression; the variables analysed were age at biopsy (<40 years v >40 years), sex, infective virus (HBV v HCV), activity of liver

![Image](http://gut.bmj.com/)
inflammation (A0/A1 v A3), and degree of liver fibrosis (F0–2 v F3/F4). Data analyses were performed with the SAS statistical package version 6 (SAS Institute, Cary, North Carolina, USA, 1990). All p values were two tailed and a level of <0.05 was accepted as statistically significant.

### RESULTS

#### Ploidy/binuclearity analyses by in vivo fluorescence imaging

Using a recently developed approach,20 we analysed the modification of ploidy during chronic hepatitis. By simultaneous

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**Table 1**  Patient characteristics

<table>
<thead>
<tr>
<th></th>
<th>Total</th>
<th>Low activity (A0, A1)</th>
<th>High activity (A3)</th>
<th>Low activity (A0, A1)</th>
<th>High activity (A3)</th>
</tr>
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<tr>
<td>No of patients</td>
<td>57</td>
<td>15</td>
<td>15</td>
<td>15</td>
<td>16</td>
</tr>
<tr>
<td>Sex (M/F)</td>
<td>41/16</td>
<td>7/8</td>
<td>10/5</td>
<td>10/1</td>
<td>14/2</td>
</tr>
<tr>
<td>Age (y) (mean (SD))</td>
<td>39.1 (5.0)</td>
<td>37.5 (4.6)</td>
<td>39.8 (5.5)</td>
<td>41.8 (4.6)</td>
<td>39.1 (5.1)</td>
</tr>
<tr>
<td>Hepatitis virus (HBV/HCV)</td>
<td>23/33</td>
<td>5/10</td>
<td>6/9</td>
<td>5/6</td>
<td>8/8</td>
</tr>
</tbody>
</table>

Activity and fibrosis were scored according to the METAVIR scoring system.21

HBV, hepatitis B virus; HCV, hepatitis C virus.

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**Figure 2**  Ploidy/binuclearity distributions, according to the progression of chronic hepatitis. (A, B) Fraction of mononuclear diploid hepatocyte; (C, D) fraction of mononuclear polyploid hepatocytes; and (E, F) fraction of binuclear hepatocytes. (A, C, E) Comparison between low and high activity of inflammation; and (B, D, F) comparison between low and high degrees of fibrosis. Box plots for the percentage of hepatocytes in each population (mononuclear diploid hepatocytes, mononuclear polyploid hepatocytes, and binuclear hepatocytes) out of the total number of hepatocytes analysed. The ends of each box represent the quartiles, and the median of distribution is indicated by a line within the box; n corresponds to the number of patients in each group. All p values were calculated using the Mann-Whitney U test.
We selected 57 patients with chronic hepatitis caused by HBV or HCV infection (24 with HBV and 33 with HCV). Based on their histological findings, we grouped these patients in relation to the degree of activity of hepatitis and the degree of fibrosis, according to the METAVIR scoring system (table 1).\textsuperscript{21} It is important to note that hepatocyte ploidy/binuclearity is reported to change according to age in normal human liver.\textsuperscript{7} Therefore, we selected age matched patients (39.1 (5.0) years) in order to reduce the influence of this parameter. Firstly, we analysed hepatocyte ploidy/binuclearity in age matched normal human liver samples (n = 3). The fractions of mononuclear diploid, mononuclear polyploid, and binuclear hepatocytes were 76.1 (2.7)%, 9.6 (2.2)%, and 14.3 (3.6)%, respectively. Only a few studies have analysed normal human livers but our values showed a proportion of the different fractions of cells which were “intermediate” between those found in two independent studies.\textsuperscript{5,7}

**Ploidy/binuclearity distribution according to the severity scores of the disease**

Given the very low fraction of mononuclear 8n and binuclear 2×4n in our studied cell population, the results were expressed globally as mononuclear polyploid and binuclear hepatocytes. Thus, proportions of mononuclear diploid hepatocytes, mononuclear polyploid hepatocytes, and binuclear hepatocytes were compared according to activity and fibrosis scores, respectively (fig 2). The percentage of mononuclear diploid hepatocytes was significantly reduced in both patients with high hepatitis activity and marked fibrosis (low activity: 75.1 (18.8)% vs high activity: 61.8 (21.6)%, p = 0.0111 (fig 2A); and low fibrosis: 77.3 (13.8)% vs high fibrosis: 57.4 (23.3)%, p = 0.0002 (fig 2B)). Accordingly, the percentage of mononuclear polyploid hepatocytes increased in both patients with high hepatitis activity and marked fibrosis (low activity: 11.9 (15.5)% vs high activity: 22.1 (20.1)%, p = 0.0166 (fig 2C); and low fibrosis: 9.4 (10.7)% vs high fibrosis: 26.4 (21.6)%, p = 0.0001 (fig 2D)). In contrast, we did not find any difference in binuclear hepatocyte percentages between either patients with low and high activity, or between patients with low and high fibrosis (fig 2E, F). Multivariate analysis showed that fibrosis was an independent factor associated with increased mononuclear polyploid (p = 0.035) and decreased mononuclear diploid (p = 0.04) cell fractions.

These results demonstrated a shift in the distribution of hepatocyte ploidy towards an increase in mononuclear.
polyplody hepatocytes associated with activity and, more strongly, with fibrosis (that is, severity of chronic hepatitis).

**Distribution of the fraction of mononuclear cells in the population of polyplody hepatocytes (FMP) according to progression of hepatitis**

In order to investigate changes in the balance between mononuclear (4n, 8n) and binuclear (2×2n, 2×4n) hepatocytes within the polyplody hepatocyte population, the fraction of mononuclear hepatocytes in the latter (FMP) was calculated for each patient. FMP levels increased in patients with high hepatitis activity (low activity: 0.36 (0.24) vs high activity: 0.47 (0.22); p = 0.0807) and in those with marked fibrosis (low fibrosis: 0.32 (0.20) vs high fibrosis: 0.54 (0.21); p = 0.0005). When the analysis was based on a combination of activity and fibrosis, we observed a gradual increase in FMP levels, together with an increase in the activity and fibrosis scores (fig 3).

**Comparison of ploidy/binularity profiles during HBV and HCV infections**

The distribution of hepatocyte ploidy/binularity was compared between patients with chronic hepatitis B and C to test for a direct effect of HBV and HCV on the distribution of hepatocyte ploidy/binularity. There were no significant differences in the distribution of mononuclear diploid and mononuclear polyplody hepatocyte fractions between patients with HBV and HCV infections although there was a trend towards a lower mononuclear diploid hepatocyte fraction in HBV infected patients (fig 4A, 4B). In contrast, the fraction of binuclear hepatocytes was significantly higher in patients with HBV than in those with HCV (HBV: 18.2 (7.6)% vs HCV: 12.8 (4.8)%; p = 0.0020) (fig 4C). In addition, multivariate analysis showed that HBV infection was an independent factor accounting for the higher percentage of binuclear hepatocytes (p = 0.0294, table 2). Our study therefore demonstrated a difference in the pattern of hepatocyte binucleation between patients with HBV and HCV related chronic hepatitis.

**DISCUSSION**

In this study, we assessed the evolution of liver cell ploidy status during progression of chronic hepatitis. We developed a methodology that allowed direct and simultaneous evaluation of cellular DNA content in liver cells identified in tissue sections. Using this approach, we demonstrated that chronic inflammation and fibrosis are associated with a marked increase in polyplody hepatocytes. In addition, our results showed distinct ploidy profiles in liver cells from patients with HBV and HCV related chronic hepatitis, suggesting different molecular pathways for the two viral infections.

In previous studies, human liver cell nucleus DNA content was evaluated in vitro by flow cytometry or Feulgen spectrophotometry. In some studies, DNA quantification was combined with a morphological evaluation of hepatocyte binularity, performed on serial sections from the same sample. In the present study, we validated a technique which allowed us to simultaneously identify on one tissue section mononuclear and binuclear liver cells and determine their DNA content in vivo.

Although several studies in humans have investigated patients with HCC only a few made a detailed analysis of liver sections from subjects with chronic hepatitis. In HCC, mononuclear diploid hepatocytes were mostly evidenced in precancerous nodules, early stage HCC, and well differentiated HCC. An increase in the mononuclear diploid hepatocyte fraction was also noted in the liver of patients with chronic hepatitis or cirrhosis. However, these studies did not compare patients with differing severities of inflammation and fibrosis, the two principal parameters indicating worsening of chronic hepatitis and progression to cirrhosis and HCC. In fact, we observed a decrease in mononuclear diploid hepatocyte fractions associated with an increase in mononuclear polyplody hepatocyte fractions in patients with high hepatitis activity and marked liver fibrosis. The contrast between our results and those of previous studies is likely to be due to the comparison we made between patients at different stages of liver disease. Furthermore, we selected age matched patients (57 patients, aged 39 (5) years (range 32–49)); this important issue was not taken into account for the 10 patients analysed by Melchiorri and colleagues. In addition, for the 18 patients analysed by Anti and colleagues, there was a marked distribution for patient age (58 (15) years (range 23–81)). Also, one study reported a similar distribution of ploidy/binularity in non-tumoural cirrhotic liver samples adjacent to HCC.

Another important finding in our study was the increased proportion of mononuclear versus binuclear fractions of polyplody hepatocytes (as estimated by FMP determination) in patients with high fibrosis and activity scores. High FMP levels have been reported in HCC in comparison with chronic hepatitis liver samples, and the suggestion was made that increased FMP levels may be of predictive value regarding the development of HCC. The mechanisms of liver cell polymorphosis are still largely unknown. The role of binuclear cell formation as an important step towards liver cell polymorphosis has been suggested. In fact, we have recently provided direct evidence for a pivotal role of such binuclear cells in the formation of 4n mononuclear cells and suggested the importance of centrosome homeostasis in this process. Our results may therefore be consistent with stimulation of binuclear cell DNA replication in the context of chronic inflammation and fibrosis. In addition, they do not exclude...
the possibility of a direct shift from mononuclear 2n to mononuclear 4n cells, in addition to intermediate binuclear cells.

Overall, our results in human chronic hepatitis liver samples did not demonstrate a shift of hepatocyte ploidy/binuclearity towards a predominance of mononuclear diploid hepatocytes, as was previously reported in human and animal experimentally induced HCC samples. One may hypothesise that the decrease in mononuclear diploid, and increase in polyploid, hepatocytes observed in our patients may reflect compensatory proliferation of hepatocytes in response to continuous loss of hepatocytes caused by chronic hepatitis, as has been observed after partial hepectomy.9 It is generally accepted that continuous liver injury, unlike other factors such as age, sex, activity of inflammation, and degree of fibrosis by multivariate analysis. In contrast, there was no significant difference in overall ploidy status between HBV and HCV infected subjects. The difference in the balance between mononuclear polyploid and binuclear hepatocytes indicated distinct patterns of hepatocyte polyploidisation/binucleation in HBV and HCV viral infections. HBV and HCV both trigger chronic liver inflammation and cirrhosis, a major risk factor for HCC. However, there is now strong evidence that these two viruses also act in liver carcinogenesis through direct and distinct molecular pathways (see reviews by Brechot and colleagues93 and Hildt and Hofschneider94). Our in vivo findings may pave the way towards mechanistic investigations; in particular, it is important to test whether HBV and HCV proteins may trigger some of the molecular signals which control the mononuclear/binuclear balance.

ACKNOWLEDGEMENTS

We would like to thank Jacques Emmanuel Guidotti, France Demaurex, and Dina Kremsdorf for helpful discussions. We would also like to thank Pascal Deby for the imaging facility at INRA UCR06/EA2703 Muséum National d’Histoire Naturelle.

Authors’ affiliations

H Toyoda, O Bregerie, B Nalpas, C Brechot, C Desdouets, Inserm U370-Pasteur Institute, CHU Necker, Paris, France
A Vallet, Liver Unit, Hospital Necker, Paris, France
G Pivot, Pathology Department, Hospital Necker, Paris, France

Conflict of interest: None declared.

REFERENCES