Quantitative hepatitis B core antibody level may help predict treatment response in chronic hepatitis B patients

We read with interest the recent article by Dandri emphasising the use of novel quantitative biomarkers as tools for predicting treatment response and disease progression in chronic hepatitis B infection (CHB).¹ The authors carried out a comprehensive review of the clinical implications of quantitative measurement of viral biomarkers both in blood and in liver, such as: serum HBsAg, serum HBeAg and intrahepatic cccDNA, based on current knowledge.¹ In addition to these, we would like to propose a new potential prognostic biomarker in CHB: the quantitative hepatitis B core antibody (anti-HBc) level which may help predict treatment response in CHB patients.

Anti-HBc is one of most classical serological markers for HBV infection.² However, the clinical significance of quantitative anti-HBc level is still largely unknown. By using a newly developed double-sandwich anti-HBc immunoassay validated by the WHO anti-HBc standards,^{3 4} we investigated its value in CHB patients.

In a cross-sectional cohort, we analysed the anti-HBc levels of 488 CHB patients and 350 healthy individuals with past HBV infection. The geometric mean level of anti-HBc in CHB patients was >1000-fold higher than that in individuals with past HBV infection (p<0.001, figure 1A). Among CHB patients, the anti-HBc levels in those who had elevated ALT levels (n=180) was significantly higher than those who had normal ALT levels (n=308) (p<0.001, figure 1A). Because elevated ALT levels in CHB is assumed to be due to a T-cell-



Figure 1 The quantitative anti-HBc levels in patients experiencing HBV infection. (A) Distribution of the serum anti-HBc levels during different phases of HBV infection. (B) The probability of seroconversion based on the baseline anti-HBc level grouping (high or low) according to the ROC-determined cut-off in patients receiving adefovir dipivoxil (Cohort A, 29 000 IU/mI) and peginterferon (Cohort B, 9000 IU/mI). PBI, past HBV infection; ULN, the upper normal limit.

mediated hepatocytolysis,⁵ the parallel increasing anti-HBc levels may reflect the host-adaptive anti-HBV immune activity. Moreover, the quantitative anti-HBc, thus, might also predict the response of patients receiving anti-HBV therapies.

To test this hypothesis, we further retrospectively investigated the usefulness of the baseline anti-HBc levels in predicting posttreatment response in two cohorts (A and B). Cohort A consisted of 49 HBeAgpositive patients treated with adefovir dipivoxil at 10 mg/day for 96 weeks and followed by a 12-week observation between years 2004-2007. Cohort B included 48 HBeAg-positive patients receiving peginterferon α -2a (180 µg/week) for 24 weeks, and followed by a 24-week observation in years 2005-2009 (Cohort B). As shown in table 1, patients with and without HBeAg seroconversion (SR) at the end of follow-up in the two cohorts had comparable (p>0.05) baseline values for age, gender, the levels of ALT (or stratification by 5×ULN), HBV DNA,

HBsAg and anti-HBc-IgM. However, the SR (+) patients had a significantly higher baseline anti-HBc levels than SR(-) patients, either in cohort A (p=0.005) or B (p=0.011). The predictive value (indicated by area under the curve for SR of the baseline anti-HBc level was 0.810 (95% CI 0.675 to 0.948, p=0.004) and 0.710 (95% CI 0.564 to 0.855, p=0.013) in cohorts A and B. respectively. Figure 1B showed the SR rate among patient groups classified by baseline anti-HBc level using the optimal cut-off indicated by the inflection point of the receiver operating characteristic in both cohorts. Patients with high baseline anti-HBc levels (≥29 000 IU/ml for cohort A or ≥9000 IU/ml for cohort B) had a higher SR rate than those with lower baseline anti-HBc levels (RR=7.22, 95% CI 1.69 to 30.9, p=0.006 in Cohort A; RR=2.10, 95% CI 1.06 to 4.17, p=0.021 in Cohort B). These results suggested that the pretreatment anti-HBc level may be an additional predictor for post-treatment SR both

 Table 1
 Analyses of baseline characteristics predicting HBeAg seroconversion (SR) in patients receiving adefovir dipivoxil (cohort A) and peginterferon (cohort B) treatments

Cohort A			Cohort B		
SR(+)	SR()	p Value	SR(+)	SR()	p Value
9	40	-	23	25	-
35±4	36±6	0.87	34±11	36±10	0.54
7/2	37/3	0.45	17/6	18/7	0.88
2/7	9/31	0.99	15/8	14/11	0.52
3/6	9/31	0.77	8/15	8/17	0.84
170±88	137 ± 79	0.28	198±129	213±149	0.71
7.03 ± 1.40	7.65 ± 1.13	0.16	7.64 ± 0.92	7.04 ± 1.61	0.12
4.32 ± 0.16	4.39 ± 0.65	0.78	4.01 ± 0.42	3.92 ± 1.07	0.70
3.13 ± 1.39	2.51 ± 2.49	0.47	3.38 ± 2.85	2.72 ± 3.43	0.47
$4.58\!\pm\!0.28$	4.15 ± 0.42	0.005	$4.32{\pm}0.66$	$3.81\!\pm\!0.68$	0.011
	$\begin{array}{c} \hline Cohort A \\ \hline SR(+) \\ 9 \\ 35 \pm 4 \\ 7/2 \\ 2/7 \\ 3/6 \\ 170 \pm 88 \\ 7.03 \pm 1.40 \\ 4.32 \pm 0.16 \\ 3.13 \pm 1.39 \\ 4.58 \pm 0.28 \end{array}$	$\begin{tabular}{ c c c c c } \hline Cohort A & \\ \hline SR(+) & SR(-) & \\ \hline 9 & 40 & \\ 35\pm4 & 36\pm6 & \\ 7/2 & 37/3 & \\ 2/7 & 9/31 & \\ 3/6 & 9/31 & \\ 170\pm88 & 137\pm79 & \\ 7.03\pm1.40 & 7.65\pm1.13 & \\ 4.32\pm0.16 & 4.39\pm0.65 & \\ 3.13\pm1.39 & 2.51\pm2.49 & \\ 4.58\pm0.28 & 4.15\pm0.42 & \\ \hline \end{tabular}$	$\begin{tabular}{ c c c c c c c } \hline Cohort A & & & & \\ \hline SR(+) & SR(-) & p \ Value & \\ \hline 9 & 40 & - & \\ 35\pm4 & 36\pm6 & 0.87 & \\ 7/2 & 37/3 & 0.45 & \\ 2/7 & 9/31 & 0.99 & \\ 3/6 & 9/31 & 0.77 & \\ 170\pm88 & 137\pm79 & 0.28 & \\ 7.03\pm1.40 & 7.65\pm1.13 & 0.16 & \\ 4.32\pm0.16 & 4.39\pm0.65 & 0.78 & \\ 3.13\pm1.39 & 2.51\pm2.49 & 0.47 & \\ 4.58\pm0.28 & 4.15\pm0.42 & 0.005 & \\ \hline \end{tabular}$	$\begin{array}{c c c c c c c c c c c c c c c c c c c $	$\begin{array}{c c c c c c c c c c c c c c c c c c c $

Data of Age, ALT activity, HBV DNA, HBsAg titier, IgM anti-HBc and anti-HBc level are expressed at Mean \pm SD; SR, HBeAg seroconversion.

in interferon or nucleos(t)ide analogue therapy.

In summary, baseline anti-HBc levels may serve as a useful marker indicating an ongoing host-immune activity against HBV, and the new findings may have clinical implications warranting further investigation.

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