Comparison of the new Alinity m HBV with the CAP/CTM HBV for quantification of HBV in serum and plasma

Background

HBV viral load monitoring is vital for guiding treatment decisions, and monitoring of treatment efficacy. Abbott Molecular recently released the Alinity m HBV assay to be run on the Alinity m System, a fully automated, continuous and random access analyser using ReadiFlex[™] technology.

Material and Methods

We investigated the performance of the Alinity m HBV assay in comparison to the cobas[®] AmpliPrep/cobas[®] Taqman HBV assay, version 2 (CAP/CTM HBV).

Eighty-eight serum and plasma samples with sufficient remaining sample volume after testing with the CAP/CTM HBV, were de-identified and tested with the Alinity m HBV assay. All quantifiable results were available in, or log transformed into, log₁₀ IU/mL for statistical analysis. Deming regression and Bland-Altman analysis were performed to assess correlation and agreement between the quantifiable results obtained with both assays.

Results

Fifty samples (22 plasma, 28 serum) had quantifiable results with both assays. A strong correlation (r = 0.981) was observed between CAP/CTM HBV and Alinity m HBV with Deming regression analysis (Figure 1). Bland-Altman analysis indicated that the mean difference between paired results (Alinity m minus CAP/CTM) was 0.14 log₁₀ IU/mL with a standard deviation (SD) of 0.277 $\log_{10} IU/mL$ (Figure 2).

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Alinity ₅

Figure 2. Bland-Altman analysis Alinity m HBV and CAP/CTM HBV (n = 50). Mean difference: 0.14 \log_{10} IU/mL (95% CI 0.06 – 0.22 log₁₀ IU/mL); SD: 0.277 log₁₀ IU/mL.

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Only four samples (4/50; 8.0%) had a difference greater than 0.5 log₁₀ IU/mL (difference ranged between 0.51 and 0.71 \log_{10} IU/mL).

Alinity m	CAP/CTM HBV				
HBV	> ULOQ	Quantifiable	< LLOQ	TND	Total
> ULOQ	1				1
Quantifiable	2	50	3	1	56
< LLOQ			5	4	9
TND			2	20	22
Total	3	50	10	25	88
Alinity m	HBV CAP/C	CTM HBV	Alinity (log., l	n HBV CAP/	CTM HBV

10510 10/11L) 8.32 8.33 **Alinity m HBV** $(\log_{10} IU/mL)$ 1.18

One sample tested above the upper limit of quantification (ULOQ) with both assays, and two samples tested above ULOQ with the CAP/CTM HBV but were quantifiable with the Alinity m HBV. Twenty samples were undetectable with both assays, and a further 5 samples were detected, but unquantifiable with both assays. Four samples were quantified with the Alinity m HBV (range $1.04 - 1.51 \log_{10} IU/mL$), but were either undetectable (n = 1) or detected but unquantifiable (n = 3) with the CAP/CTM HBV.

Conclusions

The Alinity m HBV assay compared well with the CAP/CTM HBV. More samples were quantifiable with the Alinity m HBV than the CAP/CTM HBV due to its broader quantification range.

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CAP/CTM HBV				
(log ₁₀ IU/mL)				
> 8.23				
> 8.23	J			
CAP/CTM HBV				
(log ₁₀ lU/mL)				
TND	J			

Alinity m HBV	CAP/CTM HBV	
(log ₁₀ IU/mL)	(log ₁₀ IU/mL)	
1.04	< 1.30	
1.13	< 1.30	
1.51	< 1.30	