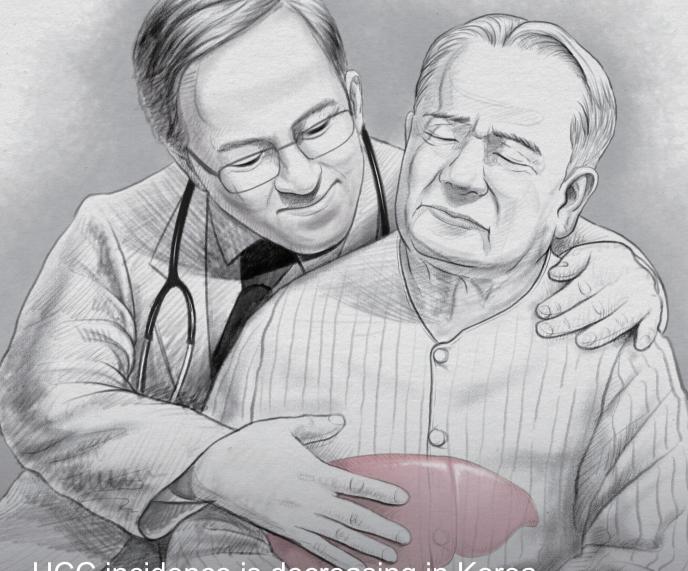
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HCC incidence is decreasing in Korea but increasing in elderly

Early changes in biomarkers predict HBsAg response Baveno-VII predicts decompensation in cACLD





Correspondence

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Correspondence on Editorial regarding "HBV pgRNA and HBcrAg reductions at week 4 predict favourable HBsAg response upon long-term nucleos(t)ide analogue in CHB"

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Keywords: Biomarkers; Treatment outcome

Dear Editor,

We sincerely appreciate the editorial piece from Liang et al. reviewing our recent paper on the role of early on-treatment decline in viral biomarkers in predicting favourable hepatitis surface antigen (HBsAg) response in chronic hepatitis B (CHB) infection, published in Clinical and Molecular Hepatology.² We agree with Liang and co-authors on the potential use of hepatitis B core-related antigen (HBcrAg) and hepatitis B virus (HBV) pre-genomic RNA (pgRNA) in multiple facets of management in the clinical context of CHB infection. Our study provided serum-liver correlations in the magnitude of decline in viral biomarkers upon nucleos(t) die analogue (NA) treatment-those with ≥1 log decline in covalently closed circular DNA (cccDNA) at week 48 had more significant reductions in serum pgRNA and HBcrAg at multiple timepoints of assessment. This further strengthens the proposition for these serum viral biomarkers to be used as surrogates for cccDNA activity.

The findings of our study suggest that subjects without

early biomarker response (defined as week 4 pgRNA decline ≥5.32 log copies/mL for hepatitis B envelope antigen (HBeAg)-positive subjects, or week 4 HBcrAg decline ≥2.05 log U/mL for HBeAg-negative subjects) had a low likelihood of achieving favourably low levels of quantitative HBsAg (gH-BsAq) (<100 IU/mL) or HBsAq seroclearance, and they should be prioritized for clinical trials while maintaining the NA therapy. As most current trials only consider gHBsAg and/or HBV DNA when screening patients for enrolment eligibility, HBcrAg and pgRNA would provide additional layer of information to identify patients who are most in need for new treatment approaches.3 HBsAg seroclearance plus HBV DNA undetectability >6 months after treatment cessation is the primary endpoint for phase III trials in the functional cure program of CHB. Notably, the benchmark of ≥30% patients achieving this endpoint⁴ has not been met by any of the currently developing novel compounds, despite initial promising results in qHBsAg knockdown by RNA interference-based therapy.^{5,6} This has engendered discussions about the practicability of such stringent treatment endpoint.⁷ Taking a step

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back, a 'looser' endpoint of achieving serum qHBsAg <10 IU/mL or <100 IU/mL (HBsAg cut-off levels still subjected to debate) by novel compounds might be more feasible, as such endpoint implies that a patient with CHB had a lower risk of off-therapy virological relapse and can potentially employ the 'stop-to-cure' approach to induce functional cure.⁸

The potential of serum HBcrAg and HBV RNA should not be limited to the context of novel compound development, but may also be applicable to consideration of NA withdrawal in those fulfilling criteria. The timing of biomarker assessment relative to NA therapy is an interesting point to consider. Our study looked at the early (as early as 4 weeks) on-treatment viral biomarker profiles instead of end-of-treatment (EOT) levels. The role of EOT pgRNA and/or HBcrAg in off therapy virological control have been investigated in multiple trials. Instead of having to wait for reaching EOT (≥3 years, which is the minimum consolidation period for NA in HBeAgnegative patients), early on-treatment profile of these biomarkers would provide valuable insights to identify patients potentially suitable for this treatment approach.

In summary, our study demonstrated that the degree of cccDNA silencing is the main determining factor for favourable HBsAg response, and can be reflected by early on-treatment changes in HBcrAg and HBV RNA. Patients without early biomarker response while on NA, as an additional consideration on top of qHBsAg levels, should be prioritized to participate in clinical trials in order to achieve functional cure.

Authors' contribution

LYM: literature review and original drafting; WKS and MFY: critical revision of article.

Conflicts of Interest -

LYM serves as advisor for Gilead Sciences. WKS received speaker's fees from AstraZeneca and Mylan, is an advisory board member of CSL Behring, is an advisory board member and received speaker's fees from AbbVie, and is an advisory board member, received speaker's fees and researching funding from Gilead Sciences. MFY serves as advisor/

consultant for AbbVie, Assembly Biosciences, Aligos Therapeutics, Arbutus Biopharma, Bristol Myer Squibb, Clear B Therapeutics, Dicerna Pharmaceuticals, Finch Therapeutics, GlaxoSmithKline, Gilead Sciences, Immunocore, Janssen, Merck Sharp and Dohme, Hoffmann-La Roche and Springbank Pharmaceuticals, Vir Biotechnology and receives grant/research support from Assembly Biosciences, Aligos Therapeutics, Arrowhead Pharmaceuticals, Bristol Myer Squibb, Fujirebio Incorporation, Gilead Sciences, Immunocore, Merck Sharp and Dohme, Hoffmann-La Roche, Springbank Pharmaceuticals and Sysmex Corporation.

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Abbreviations:

cccDNA, covalently closed circular DNA; CHB, chronic hepatitis B; EOT, end-of-treatment; HBcrAg, hepatitis B core-related antigen; HBeAg, hepatitis B envelope antigen; HBsAg, hepatitis surface antigen; HBv, hepatitis B virus; NA, nucleos(t)die analogue; pgRNA, pre-genomic RNA; qHBsAg, quantitative HBsAg

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