

BACKGROUND

Functional cure of chronic hepatitis B requires HBsAg loss and seroconversion to anti-HBs antibody. ARC-520 RNAi drug therapy targets cccDNA derived mRNA in the liver, including the HBsAg transcript, to promote host immune recovery and HBsAg seroclearance. Few biomarkers can reliably predict this outcome. Mapping the HBsAg profile based on epitope availability or occupancy has identified a predictive HBsAg Clearance Profile (CP) associated with HBsAg clearance in antiviral therapy cohorts. The developing anti-HBs response, possibly due to immune recovery on therapy, can be detected by analysis of antigen/antibody complexes.

OBJECTIVES

To evaluate the association between HBsAg response on ARC-520 RNAi therapy and the development of an HBsAg CP predictive of seroclearance, and concomitant development of a co-existing anti-HBs response suggestive of immune recovery.

MATERIALS & METHODS

Study Cohort: Consisted of 40 ARC-520 study HBeAg-negative (n=32) and HBeAgpositive (n=8) patients (under code: 30 ARC-520; 10 placebo), from pre-treatment to day 85¹. All were entecavir suppressed prior to (mean 5 years) and during ARC-520 therapy.

Diagnostic Serology: Study samples were batch analysed for quantitative HBsAg (IU/mL) and (where applicable) for HBeAg (PE IU/mL) to end-point using the Roche Cobas or Diasorin Liason platforms respectively.

HBsAg Clearance Profile (CP): Analysis of HBsAg CPs was performed using a 19plex HBsAg epitope mapping assay ^{2,3}, and results were related to HBsAg response ontreatment.

HBsAg clearance and presumably the selective pressure of an effective anti-HBs response **Complexed Anti-HBs Development:** We have developed an EIA for detection of anti-HBs therapy reduced HBsAg load in a dose-dependent manner. complexed with HBsAg, which is not able to be detected using current diagnostic assays. Briefly, HBsAg is captured by 2plex magnetic bead set conjugated with broadly specific Development of an HBsAg CP was predictive of HBsAg decline due to ARC-520 therapy, anti-HBs mAbs, and concomitant patient-derived anti-HBs identified indirectly via the value 0.038) to weeks 2 and 3 (p-values 0.019 and 0.003 respectively), and preceded or human IgG Fc domain detection. Detection of complexed anti-HBs, possibly reflective of immune recovery, coincided wit

Predicting HBsAg clearance responses during ARC-520 RNA interference (RNAi) therapy based on HBsAg epitope profile analysis

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RESULTS



HBsAg CP Development on ARC-520 Therapy

- There was a significant association between the development of an HBsAg CP and ARC-520 therapy (p-value 0.018), specifically identified at:
- Week 1 (p-value 0.038),
- Enhanced at weeks 2 and 3 (p-values 0.019 & 0.003, respectively)
- Late response association at week 6 (p-value 0.007)
- HBsAg CP preceded or coincided with an HBsAg response on ARC-520 therapy

HBsAg CP detection on ARC-520 (all timepoints)

HBsAg CP (Baseline vs on-therapy)



Cohorts 1-5	Baseline	Week1	Week2	Week3	Week4	Week6	Week8	Week12
RC-520 (n=30)	13	11	12	16	15	14	5 (5/24 tested)	5
placebo (n=10)	5	0	0	0	2	0	1 (1/8 tested)	1
p-value	0.730	0.038	0.019	0.003	0.145	0.007	1.000	1.000

Complexed Anti-HBs Development on ARC-520 Therapy

- The development of detectable anti-HBs complexed with HBsAg coincided with both:
- HBsAg response on ARC-520 therapy (trend only, did not reach significance)
- HBsAg CP detection
- Development of complexed anti-HBs may represent:
- Recovery of the immune response
- Reduction of 'free' HBsAg to increase the ratio of complexed anti-HBs/HBsAg

	REFERENCES
are key factors to achieve functional CHB cure. ARC-520	1. Yuen, M-F, <i>et al</i> . (2015) <i>Hepatol</i> 62 (Suppl): 1379A.
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