



AASLD Data Presentation

NASDAQ: ABUS

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Forward-Looking Statements

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AB-729-001 Phase 1a/1b Clinical Trial

Part 1 & 2:

Single-ascending dose

Robust HBsAg and HBV DNA declines in HBV DNA+ patients with AB-729 monotherapy (90mg single-dose)

Part 3: Multiple Ascending Dose in cHBV Patients (n=7/cohort)

E: 60mg Q4W
HBV DNA-

F: 60mg Q8W
HBV DNA-

G: 90mg Q8W + TDF
HBV DNA+

I: 90mg Q8W
HBV DNA-

J: 90mg Q12W
HBV DNA-

K: 90mg Q8W HBV DNA-,
HBeAg+ only

Baseline Demographics and Clinical Characteristics

Baseline Measure [#]	HBV DNA-					HBV DNA+
	Cohort E [‡] (n=7)	Cohort F (n=7)	Cohort I (n=6) [^]	Cohort J (n=7)	Cohort K [*] (n=7)	Cohort G (n=7)
Age in years, mean (range)	45.1 (33 – 63)	44.0 (31 – 59)	45.7 (38 – 54)	44.3 (35 – 61)	41.4 (21 – 57)	43.9 (34 – 50)
Male gender, n (%)	4 (57)	4 (57)	4 (67)	5 (71)	4 (57)	3 (43)
BMI, mean (SD)	27.7 (5.0)	23.7 (2.2)	25.5 (3.1)	28.7 (4.8)	25.0 (4.7)	23.8 (4.0)
Race, n (%)						
Asian	1 (14)	5 (71)	5 (83)	4 (57)	6 (86)	6 (86)
Black	0	1 (14)	0	0	0	0
White	6 (86)	1 (14)	1 (17)	3 (43)	0	1 (14)
Pacific Islander	0	0	0	0	1 (14)	0
ALT (U/L), mean (SD)	22.4 (10.5)	23.4 (15.2)	26.0 (10.2)	20.1 (7.2)	25.1 (8.9)	32.7 (15.8)
HBV eAg-, n (%) [◇]	7 (100)	6 (71) [◇]	5 (83)	4 (57)	0	7 (100)
HBsAg (IU/mL), mean (range)	5,372 (584 – 11,761)	5,354 (667 – 18,605)	4,691 (338 – 19,017)	6,911 (309 – 25,345)	2,221 (545 – 5,273)	1,818 (277 – 4,723)

[#] Genotype not determined

[‡] Patients switched to AB-729 60 mg Q12W for the extension phase

[^] n=6 due to 1 patient meeting exclusion criteria on D1 and a replacement patient receiving an incorrect dose on D1; both entered follow up and were excluded from analysis

[◇] One patient counted as HBeAg- was identified as “HBeAg borderline” (baseline HBeAg = 0.18 IU/mL, LLOQ = 0.11 IU/mL)

^{*} Cohort K Mean (SD) Baseline HBeAg = 22.7 (37.5) IU/mL

Robust HBsAg Declines Irrespective of Dose, Dosing Schedule, HBeAg or HBV DNA Status

Mean (SE) Baseline and $\Delta \log_{10}$ HBsAg by Visit

Nominal Visit	HBV DNA-					HBV DNA+
	Cohort E (n=7)	Cohort F (n=7)	Cohort I (n=6)	Cohort J ² (n=7)	Cohort K (n=7)	Cohort G (n=7)
Baseline (IU/mL)	3.51 (0.20)	3.53 (0.17)	3.36 (0.23)	3.37 (0.28)	3.23 (0.14)	3.14 (0.14)
Week 12	-1.10 (0.15)	-1.02 (0.11)	-1.30 (0.19)	-1.06 (0.31)	-1.63 (0.39)	-1.56 (0.32)
Week 24	-1.84 (0.16)	-1.57 (0.09)	-1.80 (0.23)	-1.56 (0.25)	-1.99 (0.35)	-1.82 (0.29)
Week 36	-1.84 (0.19)	-1.78 (0.10)	-2.06 (0.28)	-1.70 (0.39)	-2.50* (0.39)	-2.08 (0.32)
Week 48	-1.89 (0.18)	-1.90 (0.14)	1.91 (0.32)	-1.80* (0.41)	-2.57# (0.61)	-2.15 (0.34)
Week 12 Post Last Dose						
Week 12 Post Last Dose	-1.81 (0.17)	-1.74 (0.16)	-1.77 (0.31)	-1.80* (0.41)	-2.45# (0.66)	-1.97 (0.28)
Week 24 Post Last Dose						
Week 24 Post Last Dose	-1.54 (0.19)	-1.48 (0.24)	-1.67 (0.40)	-1.52 (0.40)	-2.31# (0.78)	-1.59 (0.31)

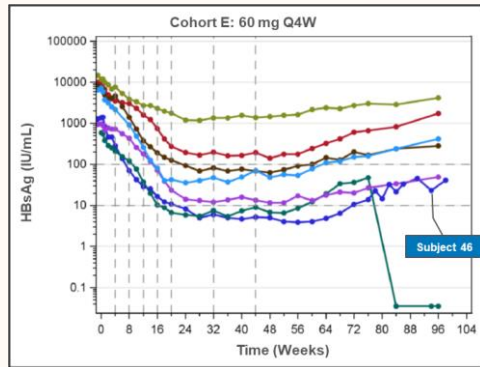
- Mean declines in HBsAg on treatment and post treatment continue to be comparable across cohorts
- Results to date from a dedicated HBeAg+ cohort (Cohort K) further support preliminary observations suggesting that baseline HBeAg status has no effect on response

Data shown as mean (SE) \log_{10} IU/mL; HBsAg LLOQ = 0.07 IU/mL, <LLOQ defined as 0.035 IU/mL Last AB-729 dose in Cohort K was at Week 40 *N=6; #N=5, 2 subjects did not receive Week 40 dose and were excluded from future timepoints

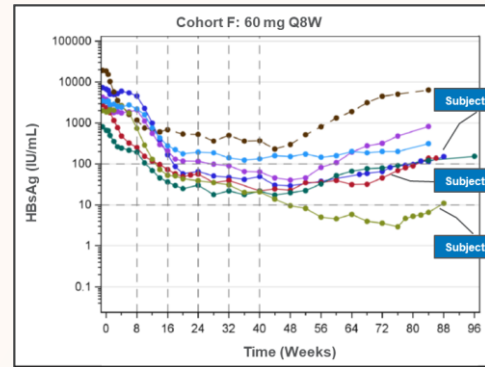
Robust HBsAg Declines Persist After Stopping AB-729

Change in HBsAg vs time

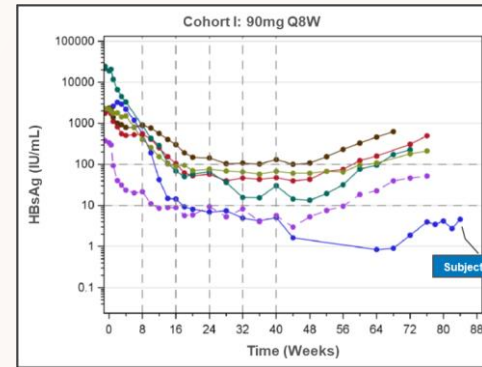
Cohort E



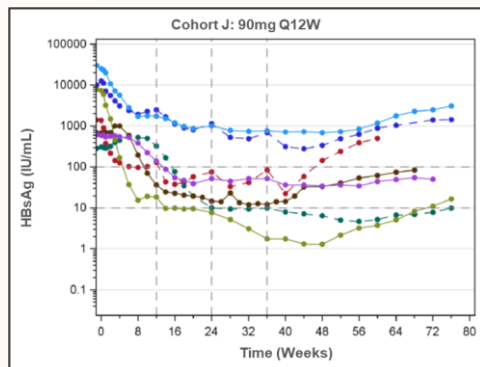
Cohort F



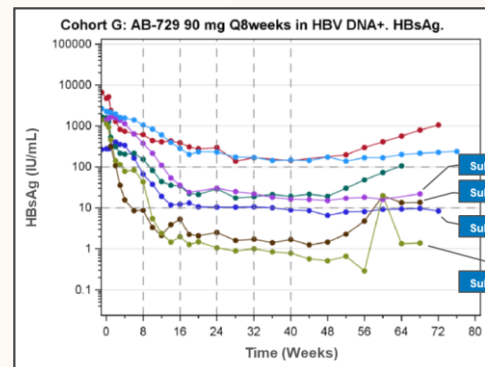
Cohort I



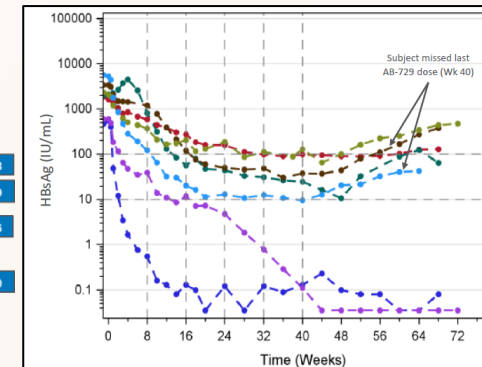
Cohort J



Cohort G

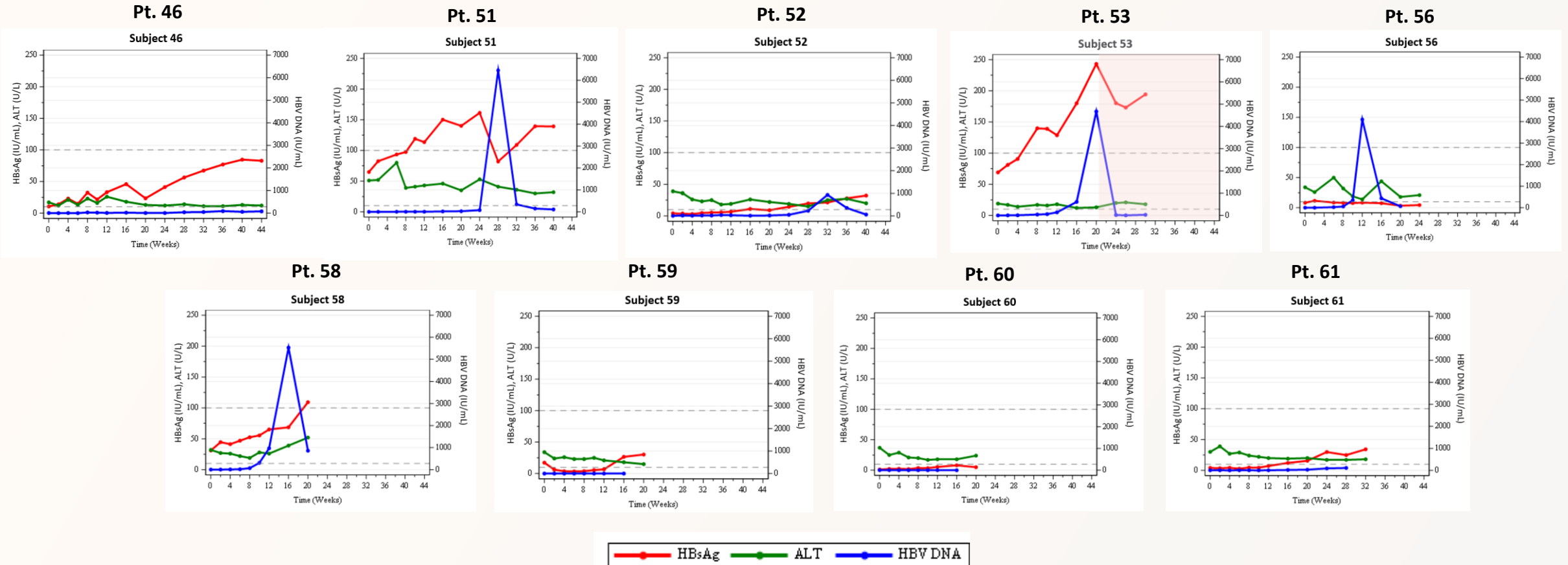


Cohort K



- 33 of 41 patients had HBsAg < 100 IU/mL at some point during the trial
- 1 patient in Cohort E (baseline HBsAg = 583.5 IU/mL) who qualified but declined to participate in NA discontinuation seroconverted at Week 84 (HBsAg < LLOQ and HBsAb = 189 IU/mL at last visit); liver enzymes remained within normal limits.
- 2 patients in Cohort K reached HBsAg < LLOQ on multiple visits with detectable HBsAb levels

HBV Control Maintained While Off-Treatment



- No patients have met virologic or clinical relapse criteria or restarted NA therapy to date
- HBV DNA has transiently increased in some patients and subsequently decreased with no intervention

AB-729-001 Safety Summary

- AB-729 generally safe and well-tolerated in clinical trial after single and repeat doses
- No treatment-related SAEs or discontinuations due to AEs
- No treatment-related Grade 3 or 4 AEs*
- No treatment-related Grade 3 or 4 laboratory abnormalities*
 - Grade 1 and Grade 2 ALT elevations have improved or stabilized with continued treatment
- Injection site TEAEs were mostly mild (erythema, pain, bruising, pruritis)
- No clinically meaningful changes in ECGs or vital signs

*1 patient (Cohort A) with rapid decline in HBsAg of ~ 2.0 log₁₀ IU/mL had an unrelated Gr 2 AE of food poisoning resulting in unrelated transient Grade 3 AEs of ALT/AST elevation (without bilirubin changes)

AB-729 Key Attributes

Unique, proprietary GalNAc delivery technology

Single trigger agent targeting all HBV transcripts

Robust reduction in HBsAg with 48 weeks of treatment

1.8 to 2.1 \log_{10} HBsAg decline in cHBV patients irrespective of dose, dosing interval or baseline characteristics

HBsAg/HBV DNA sustained when off all treatment at least 12-44 wks

No evidence of virologic or clinical relapse in 9/9 patients that discontinued AB-729 and NA-therapy

Immune activity

Patients treated with AB-729 experienced an increase in HBV-specific T-cell activation and decrease in exhausted T-cells.

Generally safe and well-tolerated in clinical trials to date

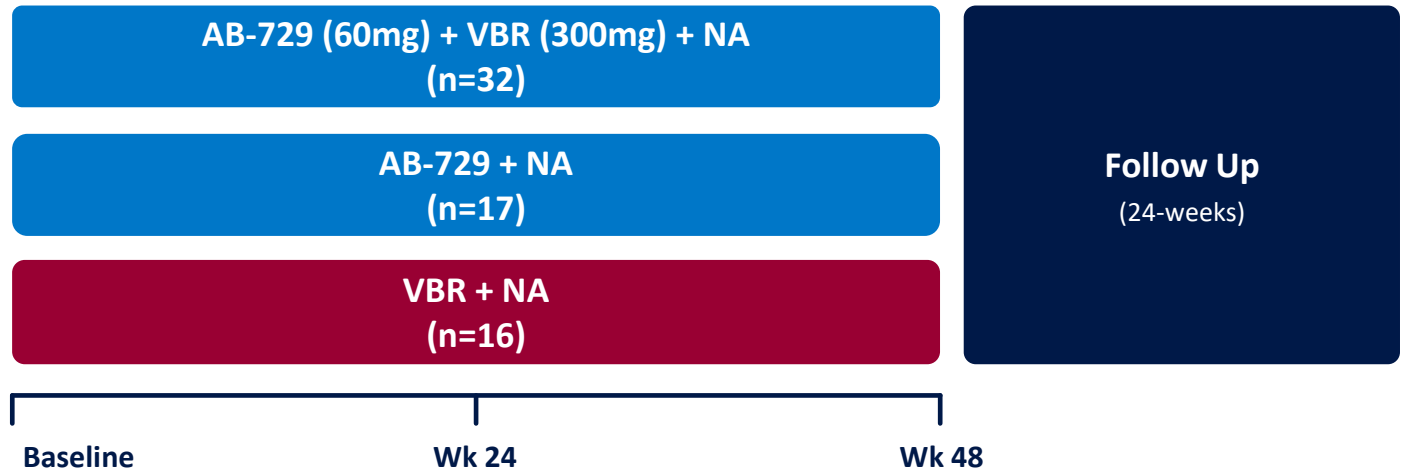
AB-729

Clinical Collaboration



**Provides accelerated
AB-729 combination**

POC with Assembly's capsid inhibitor and a NA



Primary objective: evaluate safety and tolerability of vebicorvir (VBR) in combination with AB-729 in patients with cHBV receiving NA therapy

n= 65 virologically-suppressed patients with cHBV infection

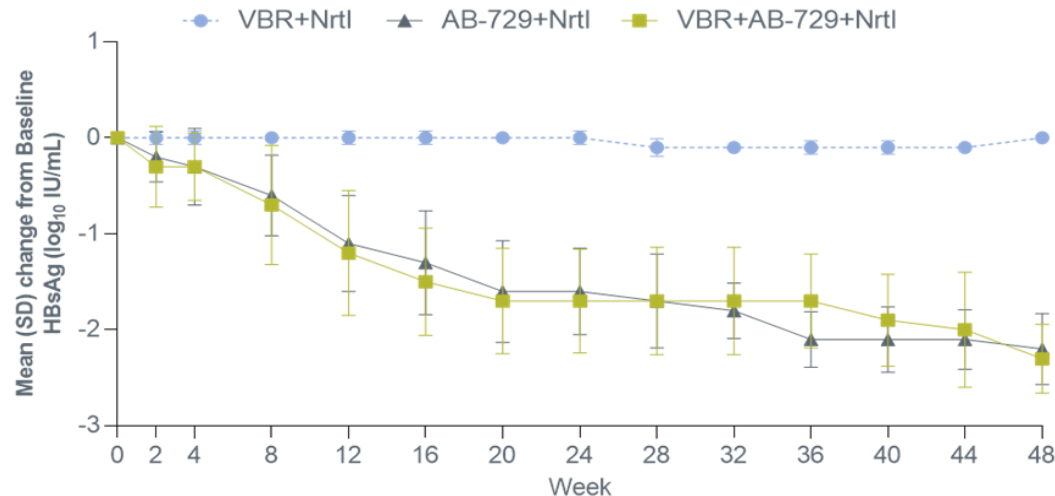
Preliminary results:

Adding VBR to AB-729+NA:

- Does not result in greater on-treatment improvements in HBV biomarkers as compared to AB-729+NA alone.
- Does not have a negative impact on reducing sAg.

Preliminary Phase 2a Triple Combination Data

Mean Changes in Virologic Parameters On-Treatment



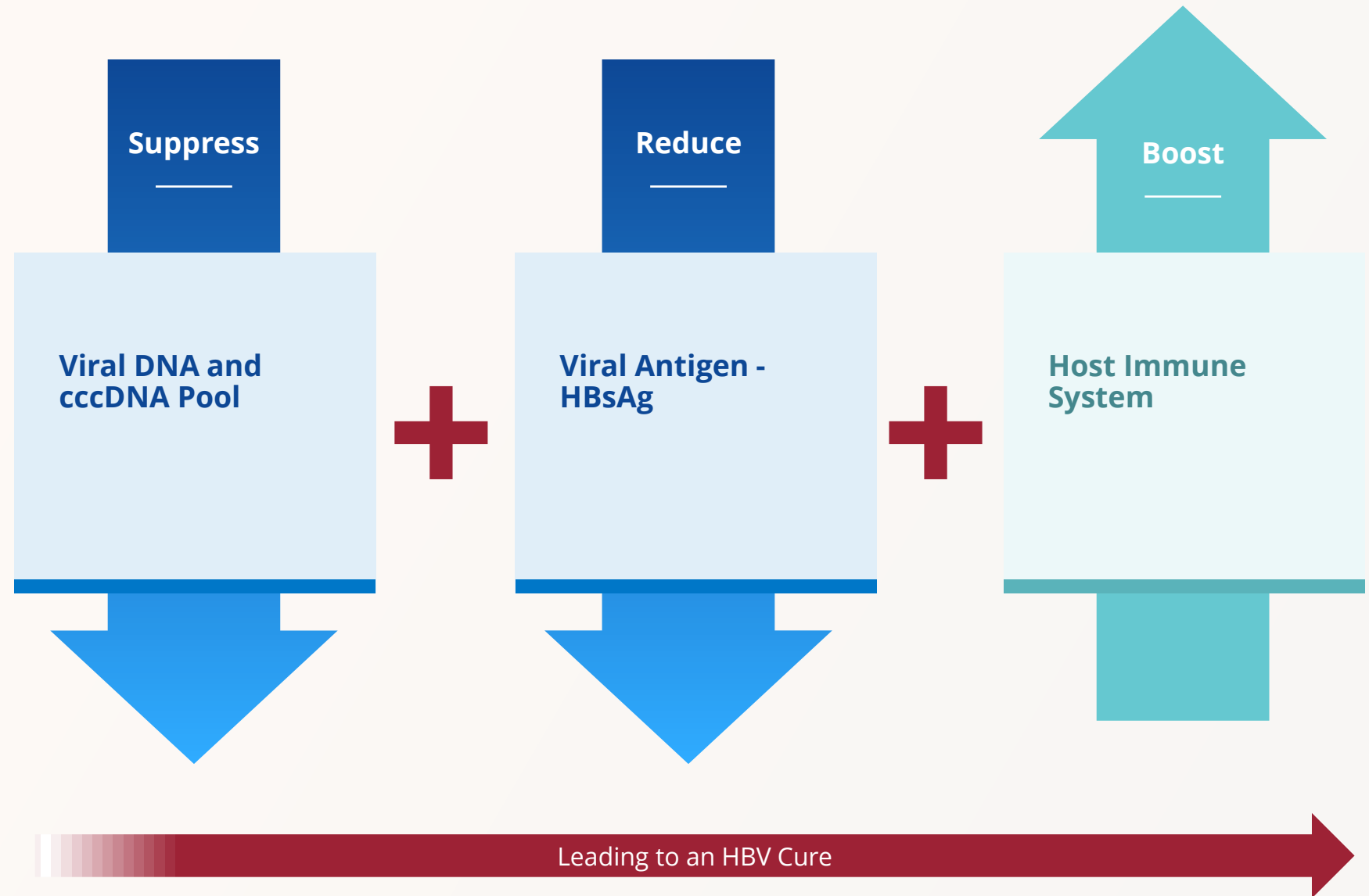
VBR+Nrtl (n):	16	15	16	14	15	13	15	14	11	10	7	4	3	2
AB-729+Nrtl (n):	17	17	15	17	17	17	16	13	10	9	6	5	3	2
VBR+AB-729+Nrtl (n):	32	32	30	29	29	28	27	24	23	19	14	7	6	4

HBsAg, hepatitis B surface antigen; NrtI, nucleos(t)ide reverse transcriptase inhibitor; SD, standard deviation; VBR, vebicorvir.

- The proportion of patients who achieved HBsAg levels <100 IU/mL and <10 IU/mL was similar in patients who received VBR+AB-729+NrtI and AB-729+NrtI
- No patients experienced HBsAg loss or seroconversion
- At the time of this analysis 0/1, 2/2 (100%), and 3/4 (75.0%) patients who received VBR+NrtI, AB-729+NrtI, and VBR+AB-729+NrtI respectively, with Week 48 data met the criteria to stop all treatment

3-Pronged Approach to Therapeutic Success

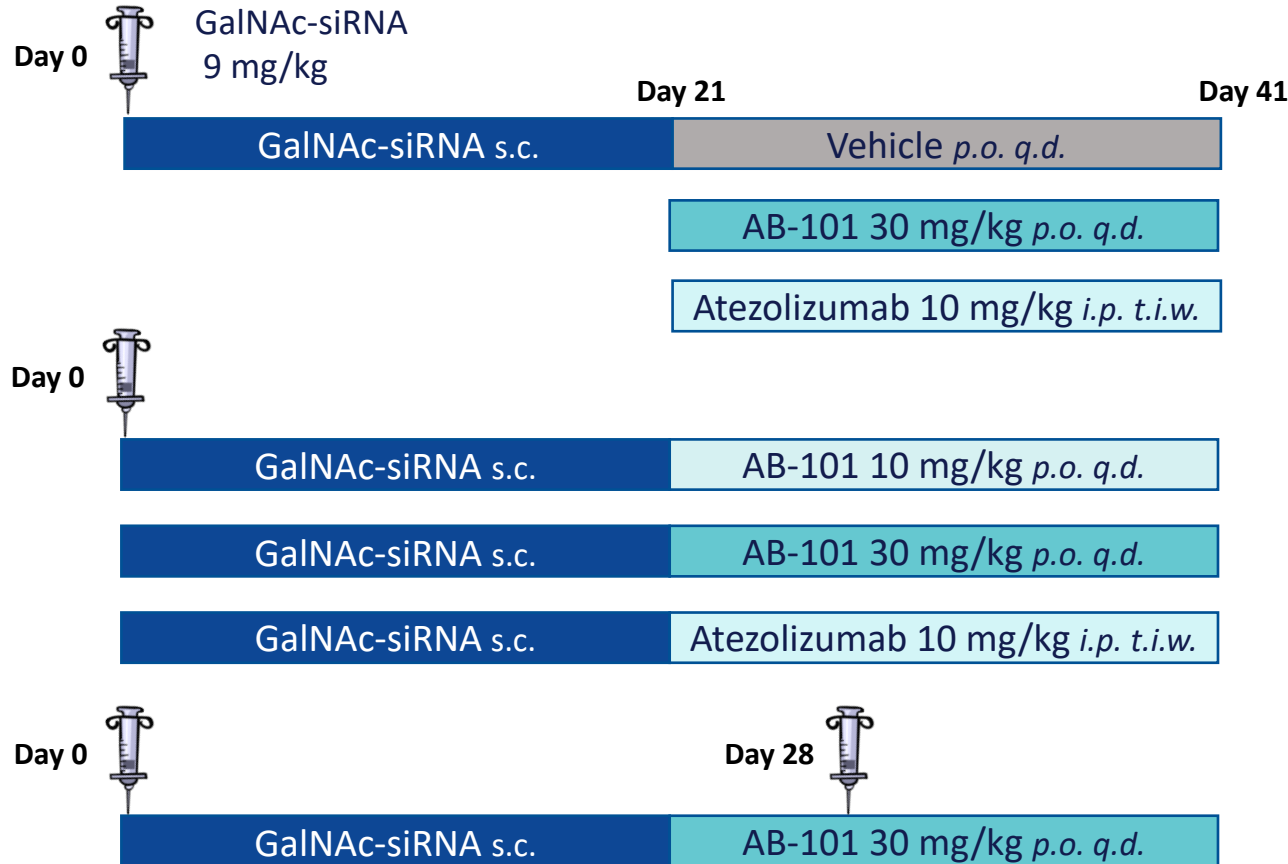
- ➔ **Suppress** HBV DNA
- ➔ **Reduce** viral antigens
- ➕ **Boost** host immune response



Therapeutic success will **require a combination of agents** with complementary MOAs.

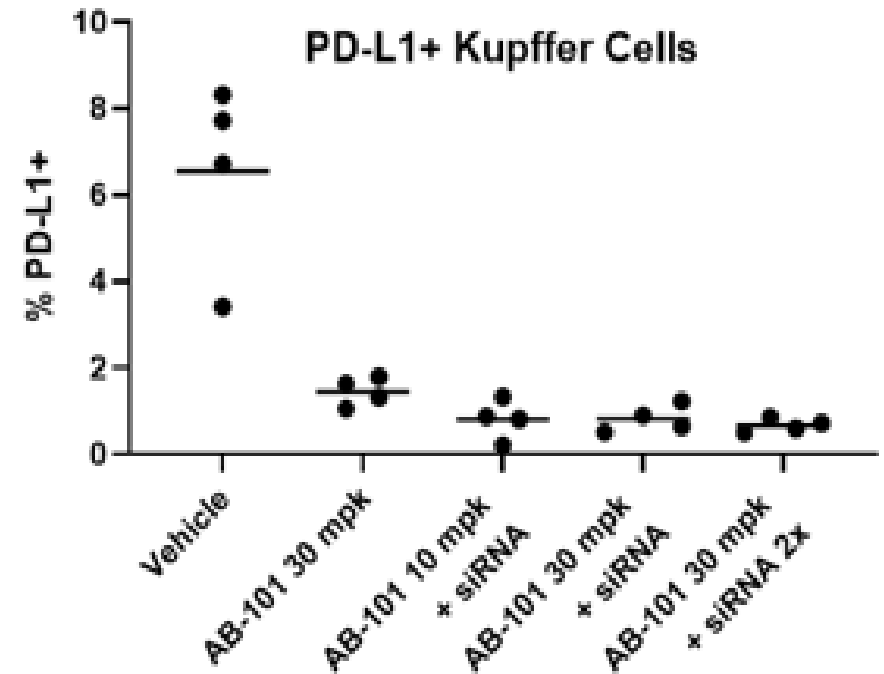
AB-101: Demonstrates Target Engagement in CD8+ Kupffer Cells in a cHBV Mouse Model

AAV-HBV hPD-L1/hPD-1 Mouse, 28 days post-AAV



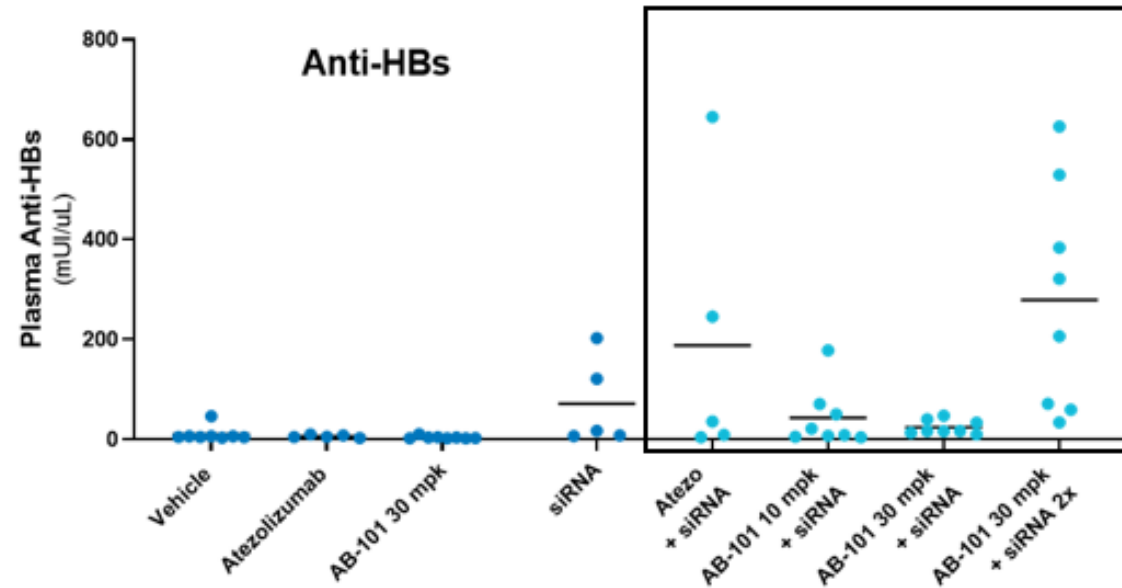
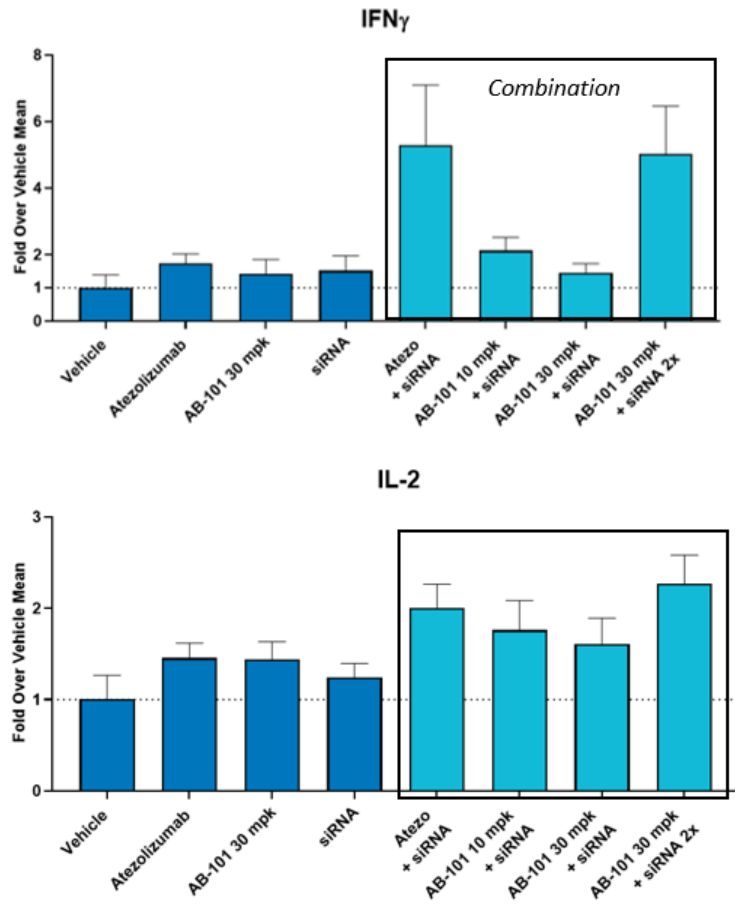
Vehicle, Naïve and positive immune response control (pHBV1.3 HDI C57BL/6) groups

AB-101 Treatment Reduces PD-L1 in Liver



AB-101+ siRNA Combination: Increases HBV-Specific T Cell Activity and HBsAb in Liver

Liver HBV-Specific T Cells



AB-101: Summary

- Oral small-molecule PD-L1 inhibitors have been identified which function through a novel internalization mechanism distinct from antibody approaches
- Combination treatment with AB-101 and HBV-targeting siRNA resulted in the activation of HBV-specific T cell and humoral responses in an AAV-HBV mouse model
- This favorable preclinical profile suggests this combination treatment strategy may provide additional benefit in increasing HBV immune responses, a key driver of CHB functional cure

Thank You

