

Long-term HBsAg suppression maintained after cessation of AB-729 treatment and comparable on-treatment response observed in HBeAg+ subjects

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SAT443

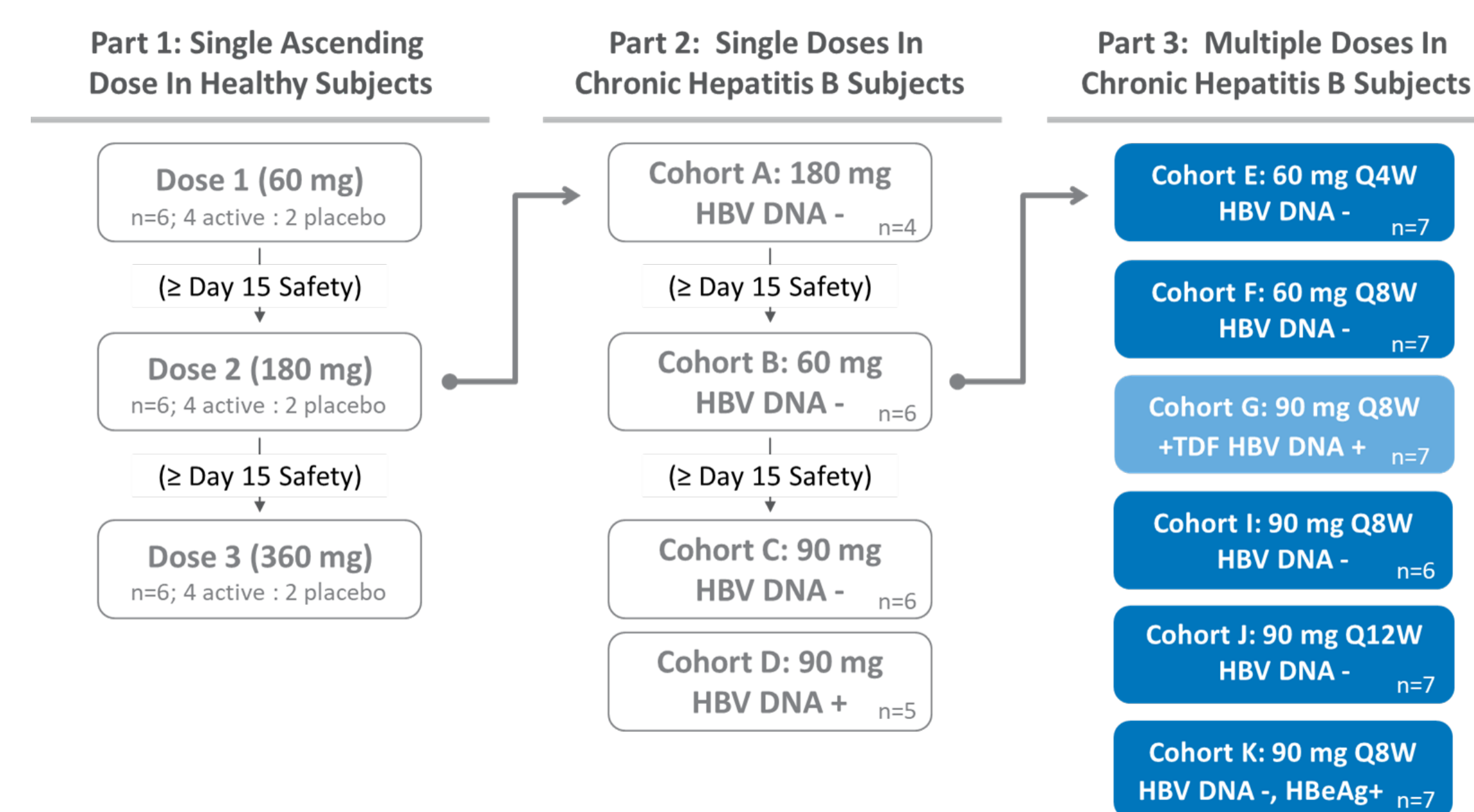


BACKGROUND

- Current therapies for chronic hepatitis B (CHB) slow or prevent the development of HBV-related liver complications, but do not typically lead to a cure.^{1,2,3} Thus, there is an unmet medical need for new finite HBV therapies that have the potential to provide a functional cure for CHB.
- AB-729 is a subcutaneously administered N-Acetylgalactosamine (GalNAc)-conjugated single trigger pan-genotypic RNA interference therapeutic that blocks all HBV RNA transcripts, including HBx, resulting in suppression of viral replication and all viral antigens. AB-729 targets the liver via proprietary technology based on GalNAc-ligand interaction with the asialoglycoprotein receptor (ASGPR). AB-729 is in Phase 2 clinical development for the treatment of CHB in combination with other agents.
- AB-729-001 is a 3-part study examining the safety and pharmacodynamics (PD) of single and repeat doses of AB-729 in healthy subjects and CHB subjects (both untreated and virologically-suppressed on nucleos(t)ide analogue [NA] therapy), and preliminary data have been reported previously.^{4,5,6,7}
- Here we report additional on-treatment and follow up data in CHB subjects following the last dose of AB-729, including the first reported data from a dedicated HBeAg+ cohort.
- Subjects meeting protocol-defined response criteria assessed at least 24 weeks after the last dose of AB-729 were given the option to discontinue NA therapy. More details for these subjects are presented in poster SAT448.

MATERIALS AND METHODS

Figure 1: AB-729-001 Study Design



- Cohorts E, F, I, J and K enrolled HBeAg positive and negative, HBV DNA- subjects on stable NA therapy. Cohort K enrolled HBeAg positive subjects only.
- Cohort G enrolled HBeAg positive and negative, HBV DNA+ subjects who began treatment with TDF concurrently with AB-729.
- All Part 3 repeat dose cohorts were initially designed for 24 weeks of treatment.
- Eligible subjects (>0.5 log₁₀ HBsAg reduction at Week 20) had the option to continue AB-729 through Week 48; all 41 subjects were eligible; 40 subjects agreed.
- Cohort E switched from AB-729 60mg Q4W to 60mg Q12W for the extension phase while the remaining cohorts maintained their initial regimen.
- Subjects are followed for at least 48 weeks after completion of AB-729 treatment.
- Assay Methods:
 - HBV DNA quantified by Abbott Realtime HBV viral load assay, LLOQ = 10 IU/mL; <LLOQ = 5 IU/mL
 - HBsAg quantified by Roche Elecsys HBsAg II quant II, LLOQ = 0.07 IU/mL; <LLOQ = 0.035 IU/mL
 - HBeAg quantified by Fujirebio Lumipulse G HBcrAg, LLOQ = 3.0 log₁₀ U/mL; <LLOQ = 2.9 log₁₀ U/mL
 - HBeAg quantified by Abbott Architect HBeAg, LLOQ = 0.11 IU/mL; <LLOQ = 0.055 IU/mL
 - ALT ULN = 48 U/L male, 43 U/L female

RESULTS

Table 1: Baseline Characteristics

Baseline Measure ^a	HBV DNA-						HBV DNA+
	Cohort E ^b (N=7)	Cohort F (N=7)	Cohort I (N=6) ^c	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 negative, n (%)	7 (100)	6 (71) ^o	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)	

^aGenotype not determined; ^bSubjects switched to AB-729 60 mg Q12W for the extension phase; ^cN = 6 due to one subject meeting exclusion criteria on Day 1 and a replacement subject receiving an incorrect dose on Day 1; both entered follow up and were excluded from the analysis; ^dOne subject counted as HBeAg negative was identified as "HBeAg borderline" (baseline HBeAg = 0.18 IU/mL, LLOQ = 0.11 IU/mL); ^eCohort K Mean (SD) Baseline HBeAg = 22.7 (37.5) IU/mL

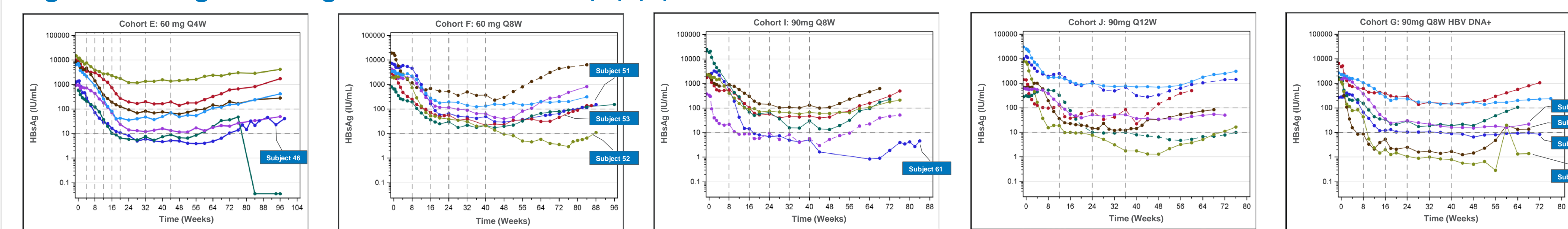
Table 2: Mean (SE) Baseline and Δ log₁₀ HBsAg by Visit

Nominal Visit	HBV DNA-						HBV DNA+
	Cohort E (N=7)	Cohort F (N=7)	Cohort I (N=6)	Cohort J ^o (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 (0.25)	-1.56 (0.35)	-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)		-2.15 (0.34)	
Week 12 Post Last Dose	-1.81 (0.17)	-1.74 (0.16)	-1.77 (0.31)	-1.80* (0.41)		-1.97 (0.28)	
Week 24 Post Last Dose	-1.54 (0.19)	-1.48 (0.24)	-1.67 (0.40)	-1.52 (0.40)		-1.59 (0.31)	

Note: Last dose Cohort E, Week 44; Cohorts F, I, G, K Week 40; Cohort J: Week 36. Mean (SE) values presented only if N≥5; ^oone subject in Cohort J chose not to extend treatment after Week 24; *Week 48 and 12 weeks post last dose are at the same visit for Cohort J

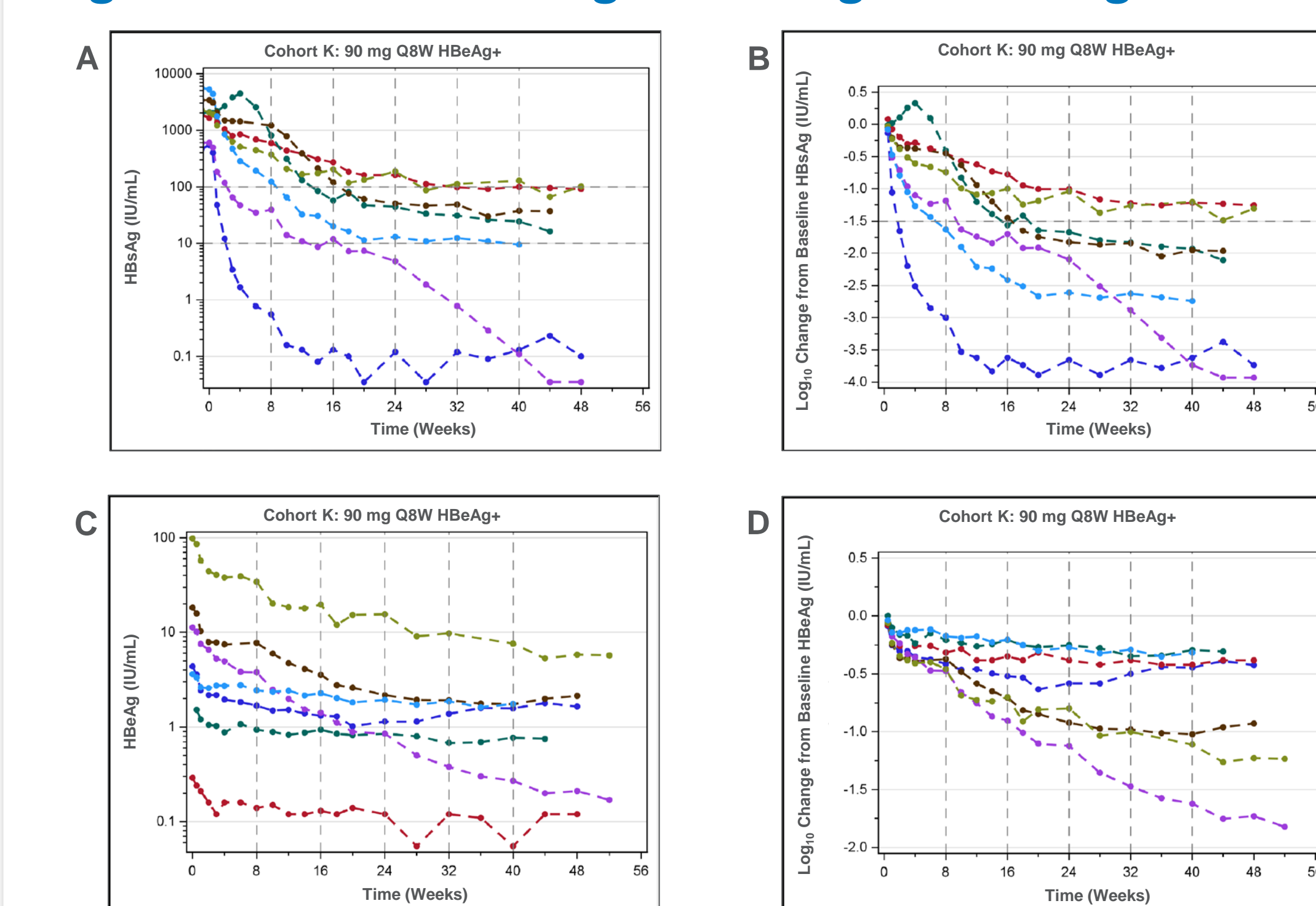
- Mean declines in HBsAg on treatment and post treatment continue to be comparable across cohorts
- Results to date from a dedicated HBeAg+ cohort (Cohort K) further support preliminary observations suggesting that baseline HBeAg status has no effect on response

Figure 2: Change in HBsAg vs time for Cohorts E, F, I, J, and G



- Robust declines in HBsAg were observed in most subjects and maintained well after the cessation of AB-729 treatment; mean log change from baseline 24 weeks post last dose is approximately -1.5 log₁₀ across cohorts
- Twenty-six of 34 subjects in these 5 cohorts had HBsAg < 100 IU/mL at some point during the study
- Eleven subjects in these cohorts were HBeAg-, HBV DNA < LLOQ, and HBsAg < 100 IU/mL with ALT < 2xULN at least 24 weeks post last dose of AB-729 and qualified to discontinue NA therapy; 9 subjects have consented
 - NA discontinuation subjects are identified in the figures above by a masked subject ID consistent with the AB-729-001 NA discontinuation poster SAT448
- One subject 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 mIU/mL at last visit); liver enzymes remained within normal limits.

Figure 3: Cohort K Change in HBsAg and HBeAg vs time



- HBsAg (panels A and B):
 - All subjects in Cohort K had HBsAg < 100 IU/mL during AB-729 treatment or follow up
 - To date, two subjects have reached HBsAg < LLOQ at one or more visits
- HBeAg (panels C and D):
 - The mean (SE) log₁₀ change from baseline in HBeAg at Week 48 was -0.94 (0.25) IU/mL
 - Log₁₀ change in HBeAg may have been limited by low baseline values (maximum HBeAg = 98.2 IU/mL)
 - One subject reached HBeAg < LLOQ and has remained near the LLOQ

Table 3: Adverse events and laboratory abnormalities

Subjects, n (%)	HBV DNA-						HBV DNA+	TOTAL [N=41]
	Cohort E [N=7]	Cohort F [N=7]	Cohort I [N=6]	Cohort J [N=7]	Cohort K [N=7]	Cohort G [N=7]		
Subjects with any TEAE	4 (57)	5 (71)	1 (17)	3 (43)	5 (71)	5 (71)	23 (56)	
Grade 1	3 (43)	4 (57)	0	2 (29)	4 (57)	4 (57)	17 (42)	
Grade 2	1 (14)	1 (14)	1 (17)	1 (14)	1 (14)	0	5 (12)	
Grade 3	0	0	0	0	0	1 (14)†	2 (5)	
SAEs (all unrelated)	0	0	0	1 (14)*	0	1 (14)‡	2 (5)	
Subjects with related TEAEs (all Grade 1)	2 (29)	4 (57)	1 (17)	2 (29)	5 (71)	2 (29)	16 (39)	
Most common related TEAEs (in ≥ 2 subjects):								
Injection site pain	0	2 (29)	0	1 (14)	4 (57)	1 (14)	9 (4)†	
Injection site erythema	2 (29)	1 (14)	0	0	1 (14)	0	5 (2)†	
Injection site bruising	2 (29)	0	1 (17)	0	0	0	3 (1)†	
Liver-related laboratory abnormalities:								
ALT elevation								
Grade 2	2 (29)	1 (14)	2 (33)	0	3 (43)	1 (14)	9 (22)	
Grade 3 or 4	0	0	0	0	0	0	0	
AST elevation								
Grade 2	1 (14)	0	0	0	0	1 (14)	2 (5)	
Grade 3 or 4	0	0	0	0	0	0	0	

- TEAE: treatment-emergent adverse event; SAE: serious adverse event; Grading criteria: Division of AIDS Table for Grading the Severity of Adult and Pediatric Adverse Events, V2.1 TEAE window was 12 weeks post-last dose of AB-729, data presented are cumulative from Screening/Study Day 1; worst grade of TEAE or lab abnormality reported
- *SAE was an unrelated Grade 3 diagnosis of cholangiocarcinoma >12 weeks post last dose of AB-729; †SAE was an unrelated Grade 3 thigh subcutaneous cyst abscess
- †n, % is number of events/out of 242 total AB-729 doses administered in Part 3
- 2 SAEs noted were both unrelated to AB-729 treatment:
 - Grade 3 cholangiocarcinoma diagnosed >4 months post-last dose of AB-729; subject withdrawn to undergo treatment
 - Grade 3 thigh subcutaneous cyst abscess required brief hospitalization for IV antibiotics; subject continued in the study
 - Mild to moderate ALT elevations observed in DNA- CHB subjects undergoing AB-729 repeat dosing may be associated with HBV-specific T-cell IFN-γ production (see poster SAT397).

Safety Summary

- There were no deaths or treatment discontinuations due to AEs
- Two SAEs were observed, both were unrelated to AB-729 and did not impact AB-729 treatment
- The most common TEAEs related to AB-729 were injection site-related
 - All were Grade 1 and did not appear to be dose or interval dependent
- All ALT and AST elevations were Grade 2 or lower; all were asymptomatic and not considered AEs by the Investigators
 - ALT/AST elevations improved or stabilized with continued dosing
 - No bilirubin or liver synthetic function changes were seen
- No clinically significant changes in other laboratory results, ECGs, or vital signs were seen

CONCLUSIONS

- AB-729 repeat dosing continues to be generally safe and well tolerated.
- AB-729 provided a robust HBsAg decline in a cohort of only HBeAg+ subjects that was comparable to cohorts composed of mostly HBeAg- subjects which further demonstrates an absence of effect of HBeAg status on AB-729 treatment response.
 - Two subjects in Cohort K reached HBsAg < LLOQ at one or more visits
- Further follow-up of a dedicated DNA+ cohort (Cohort G) continues to demonstrate HBsAg response comparable to DNA- subjects.
- Overall, 11 of 32 subjects who completed 48 weeks of treatment and who were HBeAg-, HBV DNA < LLOQ, HBsAg < 100 IU/mL with ALT < 2xULN at least 24 weeks post last dose of AB-729 and were given the opportunity to discontinue NA therapy [see poster SAT448].
- One subject in Cohort E who qualified for but chose not to discontinue NA became HBsAg < LLOQ and seroconverted at Week 40 post last dose of AB-729 with steadily increasing HBsAb (most recent = 189 mIU/mL).
- These data support the continued evaluation of AB-729 as the cornerstone of combination treatment to achieve functional cure of chronic HBV.

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