

Continued suppression of viral markers observed following discontinuation of nucleos(t)ide analogue therapy in chronic hepatitis B subjects with low hepatitis B surface antigen levels after 48 weeks of treatment with AB-729

M-F Yuen¹, SI Strasser², W Sukeepaisarnjaroen³, J Holmes⁴, V Sharma⁵, D Antoniello⁵, E Medvedeva⁵, EP Thi⁶, G Picchio⁵, T Eley⁵, KD Sims⁵

¹Queen Mary Hospital, The University of Hong Kong, Hong Kong, ²Royal Prince Alfred Hospital, Sydney, Australia, ³Srinagarind Hospital, Khon Kaen University, Khon Kaen, Thailand, ⁴St. Vincent's Hospital, Melbourne, Australia, ⁵Arbutus Biopharma Clinical Development, ⁶Arbutus Biopharma Research



SAT448



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 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}
- An amendment to AB-729-001 permitted the optional discontinuation of NA therapy in Part 3 subjects who completed 48 weeks of AB-729 treatment and who met protocol-defined NA stopping criteria assessed at least 24 weeks after the last dose of AB-729.
- Here we report preliminary safety and virology data from the subjects who have elected to participate in the NA discontinuation period to date.
- Follow up data for the remainder of the subjects in study AB-729-001 who continued on NA therapy for the duration of follow-up is presented in Poster SAT443, and additional immunology data for a subset of study subjects is presented in Posters SAT396 and SAT397.

MATERIALS AND METHODS

Figure 1: AB-729-001 Study Design (Part 3)

Part 3: Repeat Doses In Chronic Hepatitis B Subjects (open-label)

Cohort E: 60 mg Q4W HBV DNA - n=7

Cohort F: 60 mg Q8W HBV DNA - n=7

Cohort G: 90 mg Q8W + TDF: HBV DNA+ n=7

Cohort I: 90 mg Q8W HBV DNA - n=6

Cohort J: 90 mg Q12W HBV DNA - n=7

Cohort K: 90 mg Q8W HBV DNA-/HBeAg+ n=7

- Study AB-729-001 is ongoing; however AB-729 dosing is complete
- Cohorts E, F, I, and J 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 on Study Day 1
- The option to stop NA therapy was limited to those subjects that completed 48 weeks of AB-729 treatment (total number of AB-729 doses varied according to dosing schedule) via an optional 24 week treatment extension
- Eligibility was determined using the following criteria on or after 24 weeks post last dose of AB-729:
 - ALT < 2 × ULN, and
 - Undetectable (target not detected, TND) HBV DNA, and
 - HBeAg negative, and
 - At least one of the following:
 - HBsAg undetectable for at least 24 weeks after the last dose of AB-729
 - HBsAg < 100 IU/mL at two consecutive visits at least 24 weeks after the last dose of AB-729
 - HBsAb positive for at least 24 weeks after the last dose of AB-729

- After stopping NA, subjects were evaluated every 2 weeks for the first 12 weeks, then monthly

- Clinical laboratory testing and HBV parameters were collected at each visit

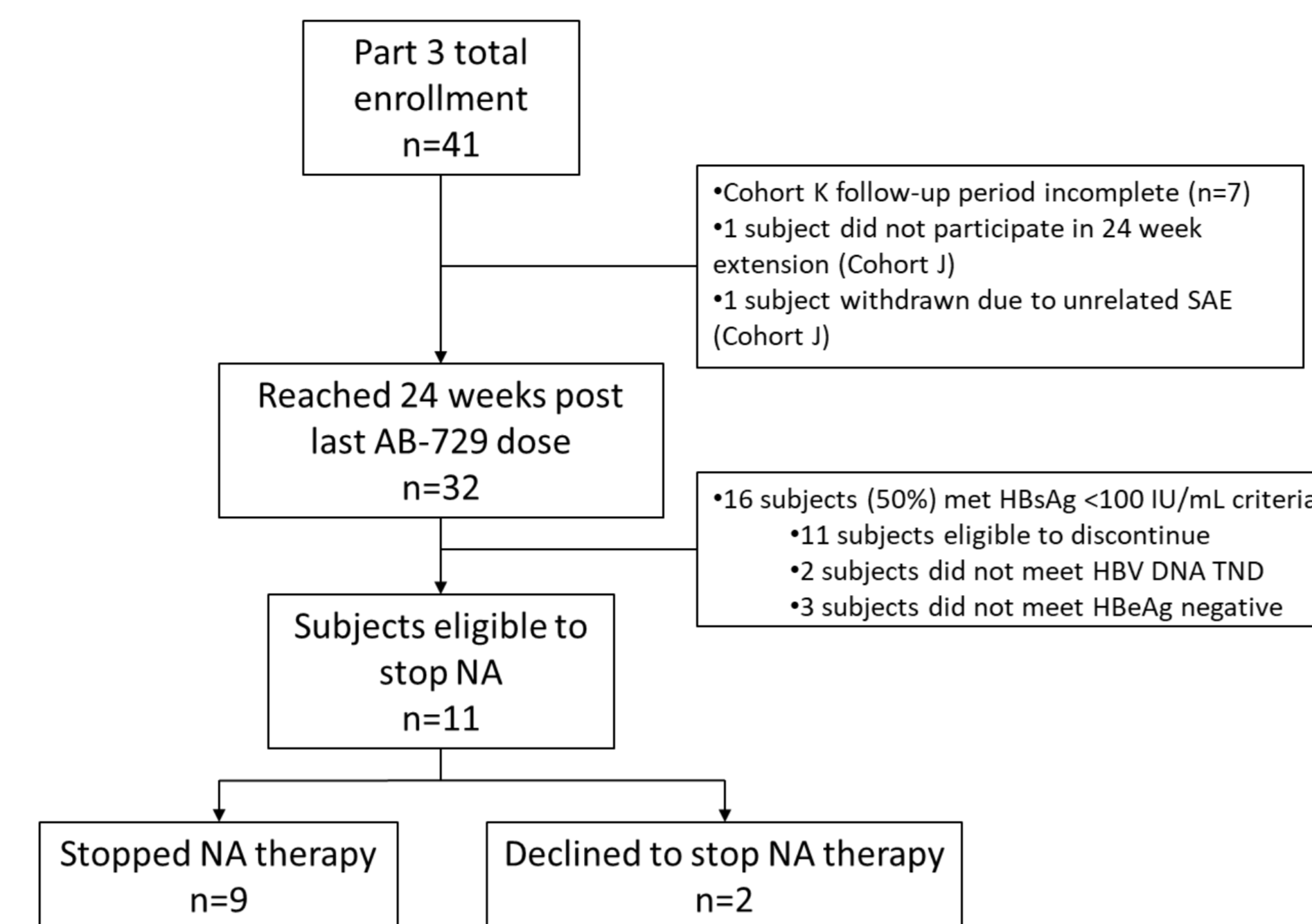
- NA therapy could be restarted if subjects met protocol-defined criteria

Study assay methods/cutoffs:

- HBV DNA was assessed with Abbott Realtime HBV viral load assay, LLOQ = 10 IU/mL
- HBsAg was assessed with Roche Elecsys HBsAg II quant II, LLOQ = 0.07 IU/mL
- HBcrAg was assessed with Fujirebio Lumipulse G HBcrAg, LLOQ = 3.0 log U/mL
- HBV RNA was assessed with Abbott RUO HBV RNA V1.0 or 2.0, LLOQ = 1.65 log₁₀ U/mL (V1.0) or 0.49 log₁₀ U/mL (V2.0)
- ALT upper limit of normal (ULN) = 48 U/L for males, 43 U/L for females

RESULTS

Figure 2: Subject Disposition



- Subjects with at least 4 weeks of follow-up data are presented (n=5); duration of subject follow-up ranged from 4 weeks to 24 weeks post-NA discontinuation

Table 1: Baseline Characteristics

Baseline Measure	Subject 46	Subject 51	Subject 52	Subject 53	Subject 61	Subject 56	Subject 58	Subject 59	Subject 60
Age (years)	35	49	36	61	56	52	50	36	46
Gender	Female	Male	Male	Female	Female	Female	Male	Male	Female
Race	Asian	Black	Asian	Asian	Asian	Asian	Asian	Asian	Asian
Study Cohort	E	F	F	F	I	G	G	G	G
NA therapy at study entry	ETV	ETV	TDF	TDF	ETV	none	none	none	none
Total duration of NA therapy	9 y, 7 m	6 y, 2 m	17 y	7 y, 5 m	6 y, 5 m	1 y, 6 m	1 y, 6 m	1 y, 6 m	1 y, 6 m

- All subjects who discontinued NAs were HBeAg negative at study entry

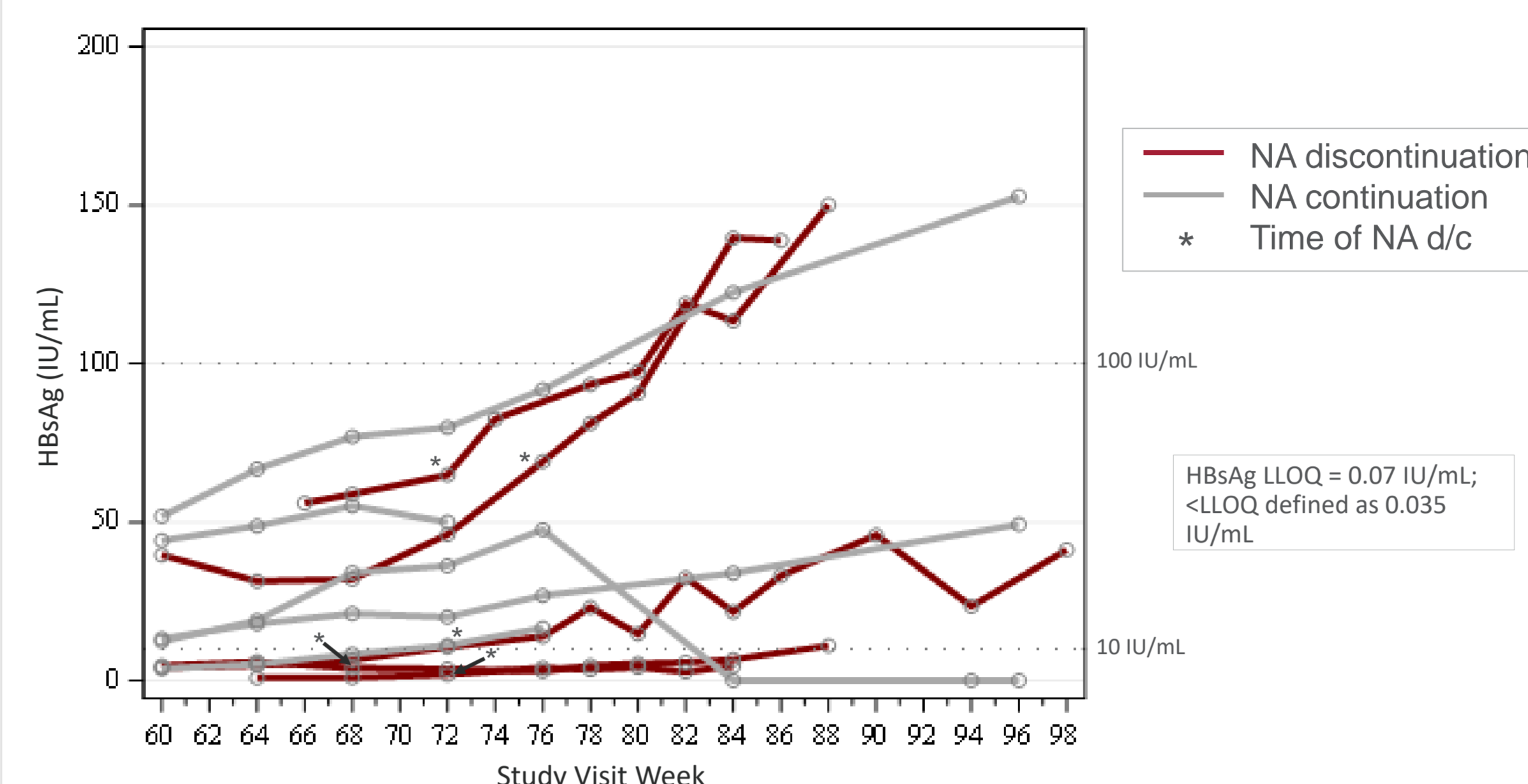
Table 2: HBV Markers

HBV Parameter	Subject 46	Subject 51	Subject 52	Subject 53	Subject 61	Subject 56	Subject 58	Subject 59	Subject 60
HBsAg (IU/mL)									
Study Day 1	1392	6765	1888	2368	2021	277	1397	1338	1128
Week 48/EOT	5.00	29.61	9.54	22.76	1.64	6.61	15.15	1.46	0.51
Last Visit prior to NA d/c	10.53	64.90	3.95	69.06	3.99	8.40	31.09	17.31	1.38
Last available post-NA d/c	41.22	150.1	10.97	138.9	4.58	N/A	N/A	N/A	N/A
HBcrAg (log U/mL)									
Study Day 1	3.8	<3.0	3.2	4.2	3.7	4.2	4.0	<3.0	3.1
Week 48/EOT	3.4	<3.0	3.0	4.4	3.4	3.6	4.0	<3.0	<3.0
Last Visit prior to NA d/c	3.4	<3.0	3.0	4.5	3.5	4.3	4.0	<3.0	3.0
Last available post-NA d/c	3.4	<3.0	3.1	4.5	3.6	N/A	N/A	N/A	N/A
HBV RNA (log₁₀ U/mL)									
Study Day 1	2.07	TND	<LLOQ	<LLOQ	N/A	3.34	2.76	1.15	1.74
Week 48/EOT	TND	TND	0.70	TND	TND	TND ^a	TND ^b	0.78 ^c	TND ^d
Last Visit prior to NA d/c*	1.29	1.07	1.20	TND	1.43	N/A	N/A	N/A	N/A
Last available post-NA d/c	1.16	1.31	1.36	1.08	1.09	N/A	N/A	N/A	N/A

Last available HBV RNA timepoint was Treatment Extension Week 44 or 40†; y = year; m = month; d/c = discontinuation; EOT = end of treatment

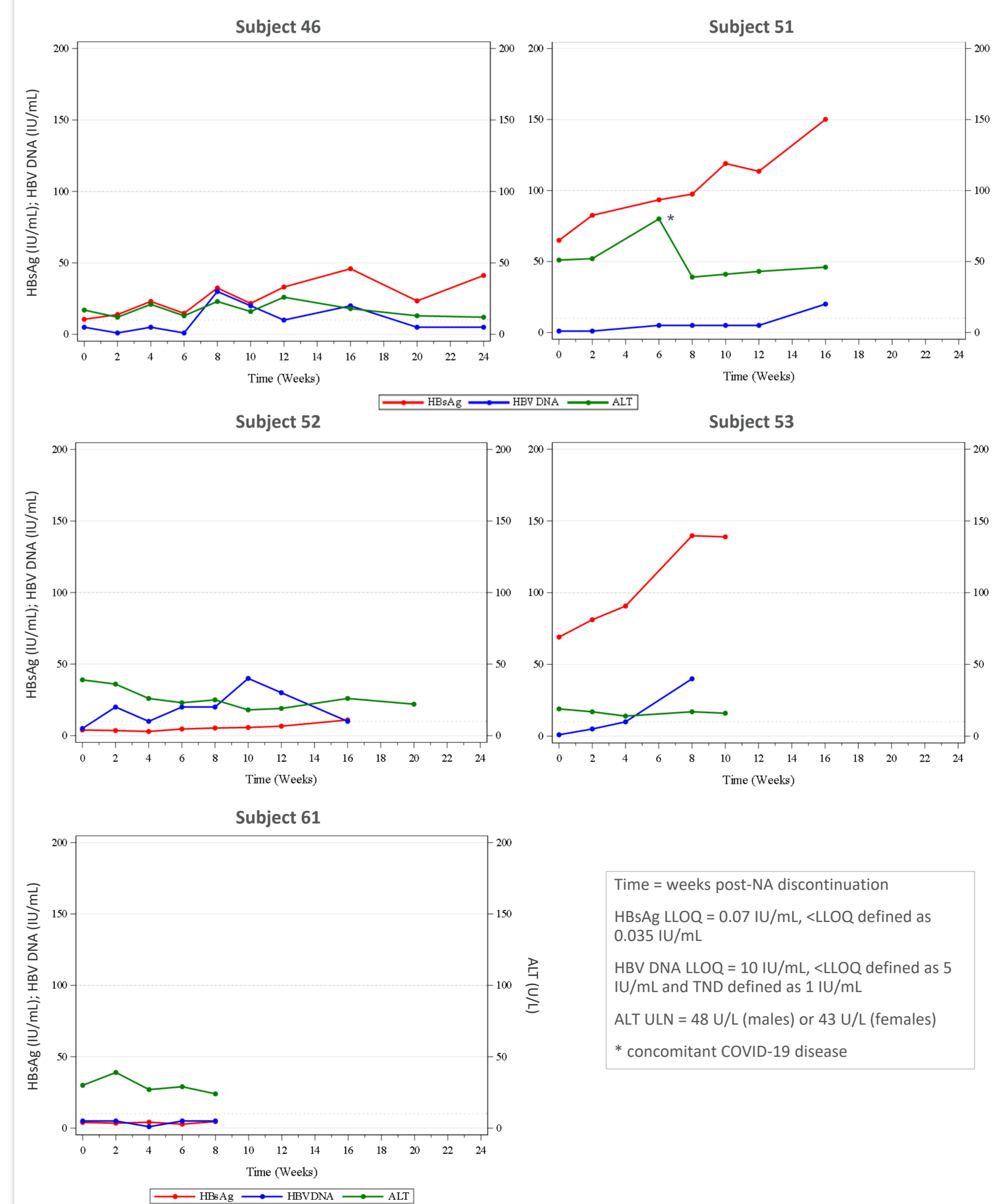
- No subjects have met criteria to restart NA therapy or had evidence of viral (confirmed HBV DNA > 2000 IU/mL) or clinical relapse (confirmed HBV DNA > 2000 IU/mL plus ALT ≥ 2 × ULN and 2 × baseline)
- Three adverse events have been reported in 2 subjects during the follow-up period, all related to COVID-19 disease
- One subject had ALT of 80 U/L at the Week 6 visit coincident with COVID-19 disease, HBV DNA was <LLOQ; ALT returned to normal at Week 8 and HBV DNA remained <LLOQ
- all other ALT values for other subjects have been < ULN

Figure 3: HBsAg levels ≥ 24 weeks post-last AB-729 dose in HBeAg-negative subjects who reached <100 IU/mL



- Data shown are for NA discontinuation subjects (red, n=5) and comparable HBeAg-negative subjects with HBsAg <100 IU/mL 24 weeks post-last dose of AB-729 that did not discontinue NA (gray, n=5)
- Most subjects maintained HBsAg <100 IU/mL for the available follow-up period
- Discontinuation of NAs did not appear to negatively impact HBsAg kinetics

Figure 4: Individual plots of HBV DNA, HBsAg and ALT post-NA discontinuation



- Data shown are individual subject plots of HBsAg (red) and HBV DNA (blue) vs ALT (green) over times since NA discontinuation
- No subjects have met viral or clinical relapse criteria or restarted NA
- HBV DNA has transiently increased in some subjects and subsequently decreased with no intervention

CONCLUSIONS

- AB-729 treatment for 48 weeks at varying doses and intervals led to continued HBsAg declines to <100 IU/mL in 50% of subjects 24 weeks after the last dose of AB-729
- Eleven of these 16 subjects met protocol-defined NA stopping criteria
- No evidence of viral or clinical relapse has been detected in the first 5 subjects to discontinue NA therapy with at least 8 - 24 weeks of follow up data available
- No subjects have restarted NA therapy
- HBsAg remains well below pre-study levels in all subjects
- Additional data is needed to determine the value of HBV RNA and HBcrAg in predicting outcomes once NA therapy is discontinued
- Discontinuation of NA therapy for up to 24 weeks has been safe and well-tolerated to date with no ALT flares observed
- Subjects will continue to be followed every 2-4 weeks for 1 year after stopping NA therapy, and longer term follow up is being amended into the protocol to monitor for sustained viral response and functional cure.

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Please see Posters SAT395, SAT396, SAT397, and SAT443 for additional data regarding AB-729 and Study AB-729-001.

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CONTACT INFORMATION AND DISCLOSURES

Karen Sims, MD/PhD
Vice President, Clinical Development
Arbutus Biopharma Inc., 701 Veterans Circle, Warminster, PA 18974
Email: ksims@arbutusbio.com

Authors affiliated with Arbutus Biopharma are employees and may own company stock.

WEBSITE:
www.arbutusbio.com