Effects of the siRNA JNJ-3989 and/or the Capsid Assembly Modulator JNJ-6379 on Viral Markers of Chronic Hepatitis B: Results From the REEF-1 Study

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Introduction



- JNJ-73763989 (JNJ-3989) is a liver-targeted short-interfering RNA (siRNA) designed to target all hepatitis B virus (HBV) RNAs for degradation, thereby reducing all HBV viral proteins and pregenomic RNA¹
- JNJ-56136379 (JNJ-6379) is a capsid assembly modulator that interferes with HBV replication by causing the formation of structually normal capsids that are devoid of HBV DNA and RNA (CAM-N)²
- The phase 2b REEF-1 study (ClinicalTrials.gov Identifier: NCT03982186) assessed the efficacy and safety of 48 weeks of JNJ-3989 and/or JNJ-6379 in combination with nucleos(t)ide analogues (NA) in patients with chronic hepatitis B (CHB)³
- JNJ-3989 treatment resulted in a dose-dependent reduction in hepatitis B surface antigen (HBsAg) through follow-up Week 24; the greatest decline was observed with the subcutaneous (SC) 200 mg dose received every 4 weeks (Q4W)
- There was no beneficial effect of coadministration with JNJ-6379 on HBsAg decline
- JNJ-3989 and/or JNJ-6379 were safe and well tolerated

Objective



o assess JNJ-3989- and/or JNJ-6379-induced changes to viral markers in CHB patients who were not currently treated (NCT) or virologically suppressed (VS) with NA treatment and who were hepatitis B e antigen (HBeAg)+ or HBeAg

Methods



- **Study Design and Participants**
- REEF-1 is a phase 2b, multicenter, double-blind, active-controlled, randomized study; results through follow-up Week 24 are reported here
- Eligible patients included those aged 18 to 65 years
- Patients were randomized to 6 treatment arms (**Figure 1**), all of which included NA, and received study treatment for 48 weeks
- Patients who met the criteria for stopping NA treatment at Week 44 (primary endpoint; **Figure 1**) terminated NA
- treatment at the Week 48 visit and began a 48-week NA-free follow-up phase • Patients could stop NA treatment and enter a 48-week NA-free follow-up phase at any time if NA stopping criteria were met
- Patients having met NA stopping criteria and having stopped NA treatment were monitored for HBV DNA and alanine aminotransferase (ALT); NA treatment restarted based on predefined NA retreatment criteria

Figure 1. Study design. Analysis up to follow-up Week 24 Inclusion criteria: -6379 250 mg PO QD (n = 48) • Active CHB (NCT or VS) HBsAg >100 IU/mL at screening NJ-3989 40 mg SC Q4W (n = 93) NA continued when NA stopping criteria⁺ not met Fibrosis stage FO-F2 x ULN, HBV DNA <LLOQ, HBeAg–, and HBsAg <10 IU/mL NJ-3989 100 mg SC Q4W (n = 93) Stratification: IA stopping criteria reassessed at every follow-up visit • HBeAg+ vs HBeAg-JNJ-3989 200 mg SC Q4W (n = 96) Treatment history (NCT vs VS) NJ-3989 100 mg SC Q4W + JNJ-6379 250 mg PO QD (n = 9 All patients received NA* during treatment F12 F24 F36 F48 **Primary endpoint:** Proportion of patients meeting NA stopping criteria

(ALT <3x ULN, HBV DNA <LLOQ, HBeAg–, and HBsAg <10 IU/mL) at Week 48 ETV, entecavir; F, follow-up; LLOQ, lower limit of quantitation; PBO, placebo; PO, oral; QD, daily TAF, tenofovir alafenamide; TDF, tenofovir disoproxil fumarate; ULN, upper limit of normal.

*NA = ETV/TDF/TAF.

Results

Participants

• A total of 470 patients with CHB were included, with a mean age of 43 years and a mean duration of CHB infection of 25.4 years; 66% were male and 40% were Asian (**Table 1**)

• 95% of patients enrolled completed the 48-week treatment phase, with no differences across treatment groups Table 1. Baseline Demographics and Clinical Characteristics by Treatment History and HBeAg Status at Screening

	N	СТ	V	′S		
Characteristic*	HBeAg+ (n = 75)	HBeAg- (n = 97)	HBeAg+ (n = 67)	HBeAg- (n = 231)		
Male, n (%)	45 (60.0)	61 (62.9)	49 (73.1)	155 (67.1)		
Asian, n (%)	42 (56.0)	16 (16.5)	48 (71.6)	84 (36.5)		
Age, years	37.2 (10.99)	40.6 (10.17)	41.7 (9.06)	46.2 (10.21)		
HBsAg, log ₁₀ IU/mL	4.48 (0.79)	3.95 (0.50)	3.61 (0.58)	3.46 (0.61)		
HBV DNA, log ₁₀ IU/mL	7.93 (1.11)	5.07 (1.37)	Not applicable ⁺	Not applicable ⁺		
ALT, U/L	107.1 (103.16)	92.5 (94.08)	24.4 (11.60)	23.7 (12.21)		
HBeAg, log ₁₀ IU/mL	2.31 (1.16)	_	0.27 (0.79)	_		
HBcrAg, n (%) <lloq‡< td=""><td>0</td><td>19 (19.8)</td><td>0</td><td>110 (48.2)</td></lloq‡<>	0	19 (19.8)	0	110 (48.2)		
HBV RNA, n (%) ≺LOD§	0	30 (31.9)	23 (34.8)	202 (89.0)		
Liver stiffness," kPa	6.30 (1.88)	5.90 (1.45)	4.84 (1.33)	4.92 (1.38)		

HBcrAg, hepatitis B core related antigen; LOD, limit of detection; SD, standard deviation. *Mean (SD) unless otherwise noted.

⁺98% of patients had HBV DNA <LLOQ (1.3 \log_{10} IU/mL = 20 IU/mL). *LLOQ = 3.0 log₁₀ lU/mL.

 $^{\text{s}}$ LOD = 2.49 log₁₀ copies/mL.

"Measured with FibroScan® Paris, France.

References

- 1. Gane E, et al. Presented at: European Association for the Study of the Liver (EASL)
- Digital International Liver Congress™; August 27-29, 2020; Virtual. Oral GS10.
- 2. Berke JM, et al. Antimicrob Agents Chemother. 2020;64(5):e02439-19.
- 3. Yuen MF, et al. Presented at: American Association for the Study of Liver Diseases (AASLD) The Liver Meeting; November 12-15, 2021; Virtual. Oral LO10.

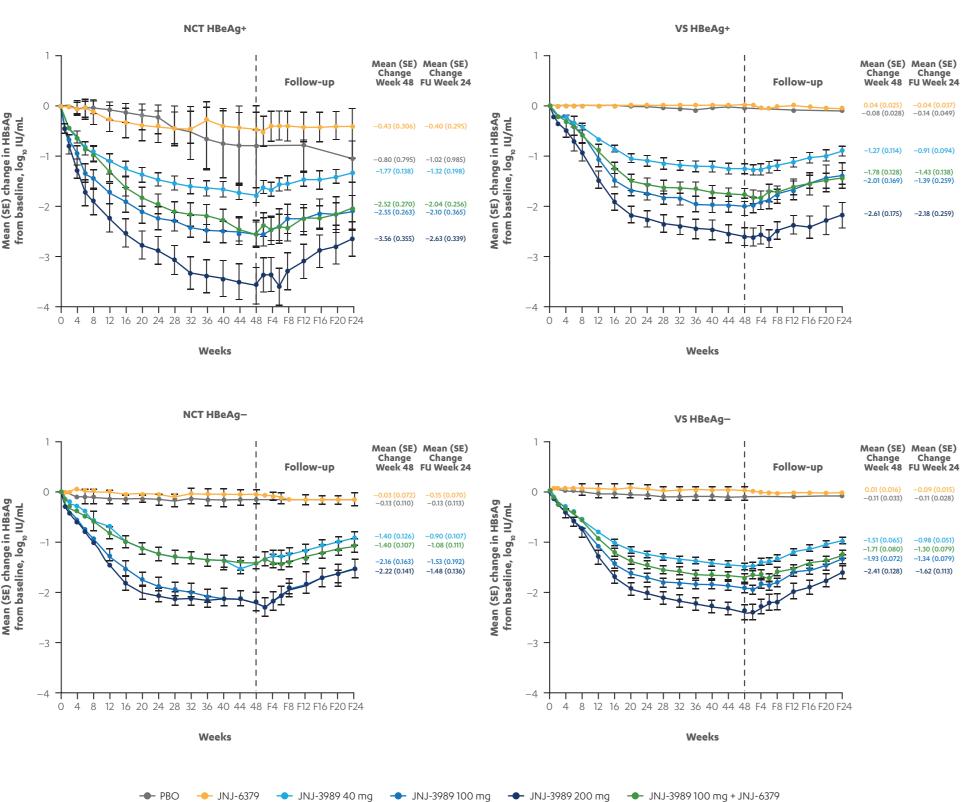
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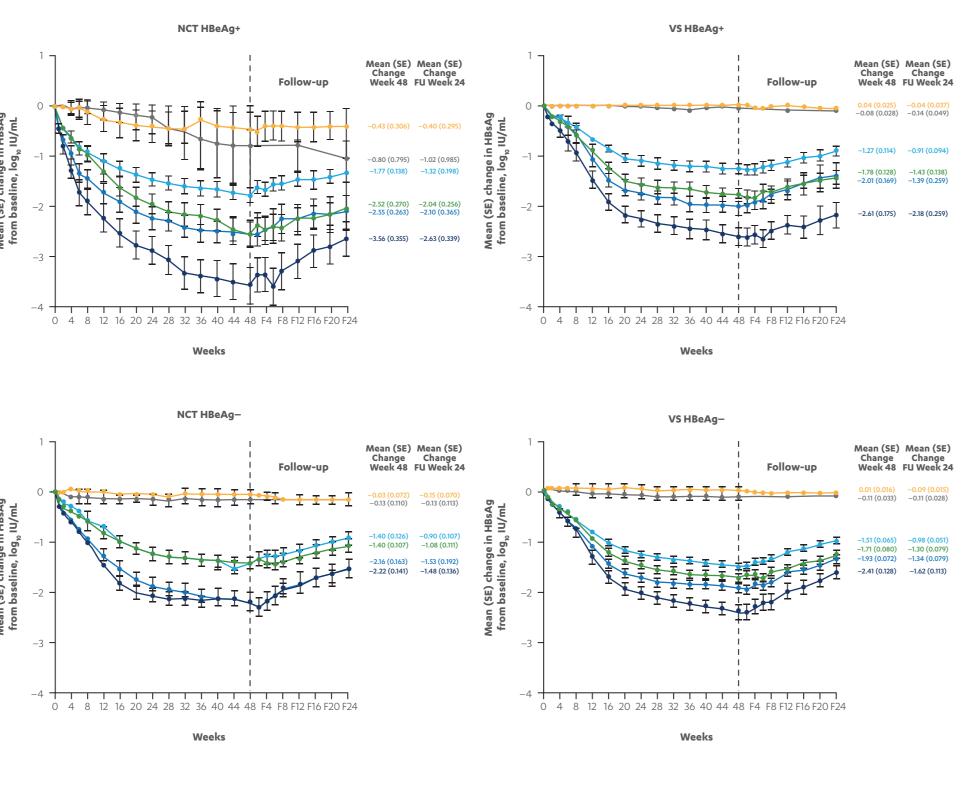
This study was supported by Janssen Research & Development, LLC. Medical writing support was provided by Kim Caldwell, PhD, of Cello Health Communications/MedErgy, and was funded by Janssen Global Services, LLC.

Presented at the European Association for the Study of the Liver (EASL) International Liver Congress™; June 22-26, 2022; London, UK & Online.

Primary Endpoint

- criteria through follow-up Week 24
- **Changes in HBsAg**

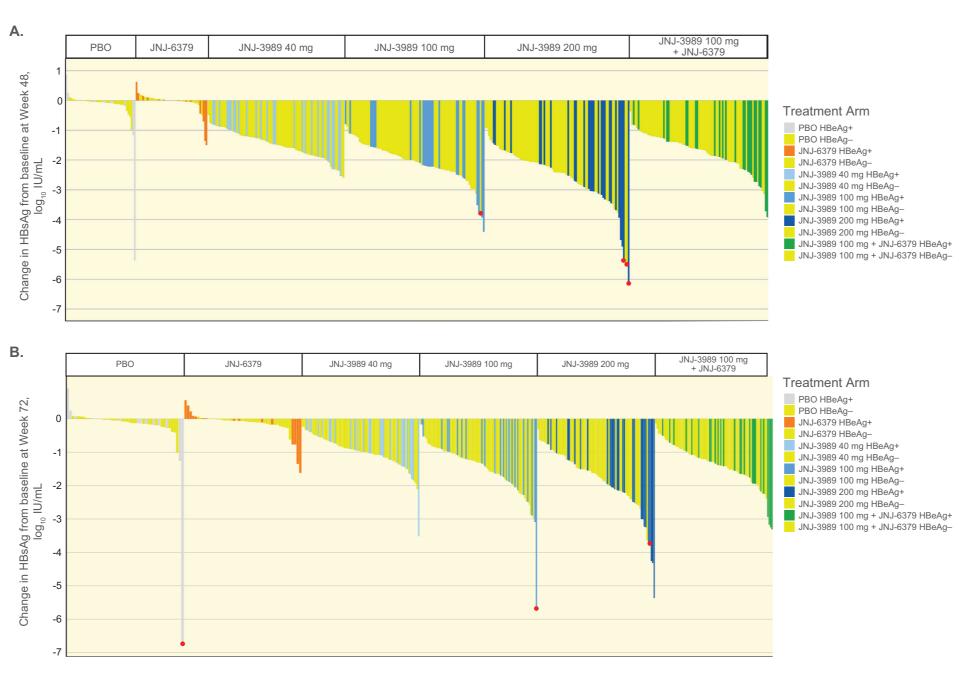






FU, follow-up; SE, standard error.

10.8%, respectively; **Figure 3B**)



Patients who achieved HBsAg loss are noted with red dots.

- Changes in HBeAg
- and Table 2)

• 19.1% (18/94) and 29.8% (28/94) of patients (all of whom were VS, except for 7) met NA stopping criteria with JNJ-3989 200 mg at Week 48 and until follow-up Week 24, respectively

- For HBeAg- patients, the main reason for not meeting NA stopping criteria was not achieving HBsAg <10 IU/mL – For HBeAg+ patients, the main reasons for not meeting stopping criteria were not achieving HBeAg seroclearance and/or HBsAg <10 IU/mL and, in those who were also NCT, not achieving HBV DNA <LLOQ

• 2 of 63 (3.2%) patients who met NA stopping criteria and stopped NA treatment subsequently met NA restarting

• Consistent with the overall population,³ JNJ-3989 reduced HBsAg in a dose-dependent manner in subgroups by treatment history and HBeAg status (**Figure 2**); the greatest mean declines were generally seen with the 200 mg dose • The largest reductions of HBsAg were observed in NCT HBeAg+ patients (**Figure 2**)

Figure 2. Mean change in HBsAg (±SE) from baseline by treatment history and HBeAg status.

• The JNJ-3989 200 mg arm had the highest proportion of patients who achieved HBsAg reduction ≥2 log₁₀ and ≥3 log₁₀ at Week 48 (73.6% and 27.5%, respectively; **Figure 3A**) and at the time of follow-up Week 24 (37.3% and

Figure 3. Individual changes in HBsAg from baseline at (A) Week 48 and at (B) follow-up Week 24 in individual patients by HBeAg status.

• Declines in HBeAg were observed across all treatment arms and sustained during follow-up, with reductions being JNJ-3989 dose dependent (Figure 4). Of note, the combination arm of JNJ-3989 100 mg + JNJ-6379 had the numerically greatest decline among NCT patients

• Reductions were most pronounced in NCT patients, potentially due to higher baseline HBeAg levels (Figure 4

• The number of patients in each treatment arm who achieved HBeAg <LLOQ was similar between NCT and VS patients, with the greatest proportion of patients reaching HBeAg <LLOQ in the JNJ-3989 100 mg (n = 3, 30.0%) and JNJ-3989 100 mg + JNJ-6379 (n = 4, 28.6%) treatment arms in VS patients (**Table 2**)

Disclosures

M-FY serves as advisor/consultant for AbbVie, AlloVir International, Arbutus Biopharma, Bristol Myers Squibb, ClearB Therapeutics, Dicerna Pharmaceuticals, Gilead Sciences, GlaxoSmithKline, Janssen, Merck Sharp & Dohme, Roche, and Spring Bank Pharmaceuticals; and receives grant/research support from Assembly Biosciences, Arrowhead Pharmaceuticals, Bristol Myers Squibb, Fujirebio Diagnostics Inc., Gilead Sciences, Merck Sharp & Dohme, Roche, Spring Bank Pharmaceuticals, and Sysmex Corp. TA serves as advisor/consultant for AbbVie, Antios Therapeutics, Enyo Pharma, Gilead Sciences, GlaxoSmithKline, Janssen, and Roche. IMJ serves as a consultant for Aligos, Arbutus, Janssen, and Gilead; has received research funds from Janssen and Assembly Biosciences; and serves on a data monitoring committee for GlaxoSmithKline. MB serves as a consultant and speaker for AbbVie, Gilead Sciences, Janssen, Esai, MSD, and Roche. HLAJ received grants from AbbVie, Arbutus, Gilead Sciences, Janssen, and Roche; and is a consultant for Arbutus, Arena, Enyo, Gilead Sciences, GlaxoSmithKline, Janssen, Merck, Roche, Vir Biotechnology Inc., and Viroclinics. TT has received lecturer fees from Bristol Myers Squibb, Gilead Sciences, and GlaxoSmithKline; research funding from Janssen: JLH serves as advisor/consultant for Aligos, Assembly, Gilead Sciences, Johnson & Johnson, and Roche; and receives grant/research support from Bristol Myer Squibb. TNK, TL, RK, CG-A, CM, JJ, TV, OL, US, and MB are employees of Janssen Pharmaceuticals and may be Johnson & Johnson stockholders.

Figure 4. Mean (±SE) change in HBeAg over time in HBeAg+ patients (A) NCT or (B) VS.

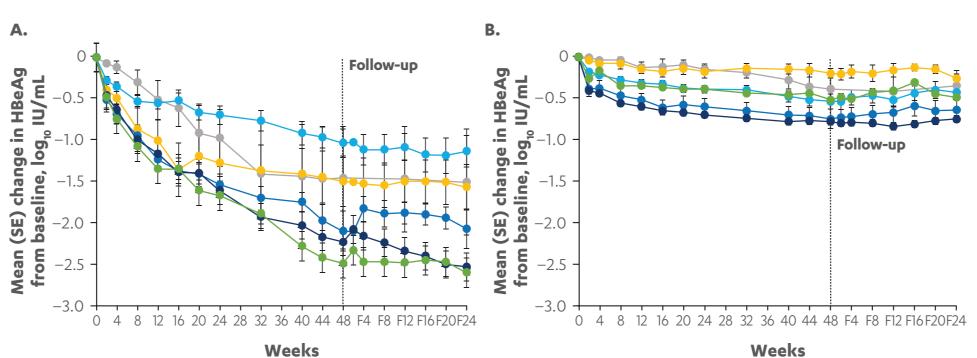


Table 2. Baseline and Change From Baseline HBeAg Values

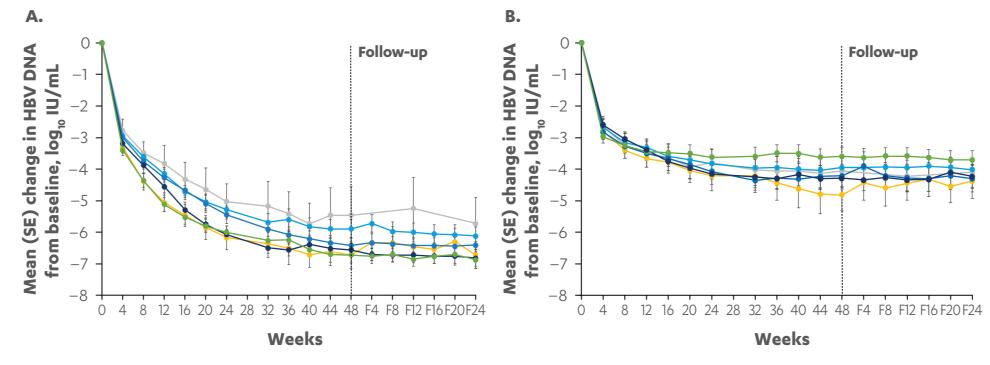
	NCT and HBeAg+									VS and HBeAg+								
	N	BL, log ₁₀ IU/mL	Change from BL at W24, log ₁₀ IU/mL	<lloq at W24, n/N (%)</lloq 	Change from BL at W48, log ₁₀ IU/mL	<lloq at W48, n/N (%)</lloq 	Change from BL at FU W24, log ₁₀ IU/mL	<lloq at<br="">FU W24, n/N (%)</lloq>	N	BL, log ₁₀ IU/mL	Change from BL at W24, log ₁₀ IU/mL	<lloq at W24, n/N (%)</lloq 	Change from BL at W48, log ₁₀ IU/mL	<lloq at W48, n/N (%)</lloq 	Change from BL at FU W24, log ₁₀ IU/mL	<lloq at<br="">FU W24, n/N (%)</lloq>		
PBO	7	1.92 (0.54)	-0.97 (0.35)	0/7	-1.45 (0.63)	1/7 (14.3)	-1.50 (0.64)	1/7 (14.3)	6	-0.02 (0.07)	-0.15 (0.07)	0/6	-0.39 (0.40)	1/6 (16.7)	-0.35 (0.36)	1/6 (16.7)		
JNJ-6379	8	1.60 (0.42)	-1.27 (0.31)	1/7 (14.3)	-1.49 (0.31)	1/7 (14.3)	-1.56 (0.28)	1/8 (12.5)	7	-0.11 (0.18)	-0.18 (0.08)	0/6	-0.20 (0.16)	0/7	-0.26 (0.26)	0/7		
JNJ-3989 40 mg	15	1.86 (0.38)	-0.69 (0.12)	1/15 (6.7)	-1.03 (0.17)	2/13 (15.4)	-1.13 (0.19)	2/15 (13.3)	15	0.67 (0.22)	-0.39 (0.05)	1/15 (6.7)	-0.54 (0.35)	1/14 (7.1)	-0.43 (0.36)	1/12 (8.3)		
JNJ-3989 100 mg	14	2.77 (0.22)	-1.53 (0.13)	1/14 (7.1)	-2.09 (0.25)	2/14 (14.3)	-2.06 (0.24)	2/13 (15.4)	11	0.29 (0.30)	-0.60 (0.10)	3/11 (27.3)	-0.75 (0.39)	3/10 (30.0)	-0.64 (0.40)	3/10 (30.0)		
JNJ-3989 200 mg	16	2.65 (0.25)	-1.60 (0.23)	0/15	-2.22 (0.30)	1/14 (7.1)	-2.52 (0.33)	2/14 (14.3)	14	0.40 (0.22)	-0.70 (0.12)	1/14 (7.1)	-0.78 (0.56)	1/14 (7.1)	-0.75 (0.53)	1/13 (7.7)		
JNJ-3989 + JNJ-6379	13	2.52 (0.25)	-1.66 (0.20)	0/15	-2.48 (0.25)	1/12 (8.3)	-2.59 (0.26)	1/12 (8.3)	14	0.02 (0.69)	-0.39 (0.12)	2/14 (14.3)	-0.52 (0.48)	4/14 (28.6)	-0.49 (0.50)	1/13 (7.7)		

Values are mean (SE) unless otherwise noted; $LLOQ = 0.11 IU/mL = -0.96 \log_{10} IU/mL$. **Changes in HBV DNA**

• A numerically greater decline in HBV DNA was seen with JNJ-3989 100 and 200 mg and JNJ-6379-containing arms compared to control in NCT HBeAg+ patients (**Figure 5A**)

• Assessment of mean change from baseline in HBV DNA in NCT HBeAg- patients was limited by a high proportion of patients reaching HBV DNA <LLOQ in all treatment arms beginning at early time points (**Figure 5B** and **Table 3**)

Figure 5. Mean (±SE) change in HBV DNA over time in patients NCT and (A) HBeAg+ or (B) HBeAg-.



→ PBO → JNJ-6379 → JNJ-3989 40 mg → JNJ-3989 100 mg → JNJ-3989 200 mg → JNJ-3989 100 mg + JNJ-6379
Table 3. Baseline and Change From Baseline HBV DNA Values

	NCT and HBeAg+									NCT and HBeAg-							
	N	BL, log ₁₀ IU/mL	Change from BL at W24, log ₁₀ IU/mL	<lloq at W24, n/N (%)</lloq 	Change from BL at W48, log ₁₀ IU/mL	<lloq at W48, n/N (%)</lloq 	Change from BL at FU W24, log ₁₀ IU/mL	<lloq at<br="">FU W24, n/N (%)</lloq>	N	BL, log ₁₀ IU/mL	Change from BL at W24, log ₁₀ IU/mL	<lloq at W24, n/N (%)</lloq 	Change from BL at W48, log ₁₀ IU/mL	<lloq at W48, n/N (%)</lloq 	Change from BL at FU W24, log ₁₀ IU/mL	≺LLOQ at FU W24, n/N (%)	
PBO	7	7.92 (0.49)	-5.03 (0.56)	2/7 (28.6)	-5.46 (0.90)	4/7 (57.1)	-5.72 (0.82)	5/7 (71.4)	9	5.20 (0.47)	-3.81 (0.43)	5/9 (55.6)	-4.07 (0.48)	9/9 (100)	-4.11 (0.47)	8/9 (88.9)	
JNJ-6379	8	8.03 (0.35)	-6.17 (0.38)	1/7 (14.3)	-6.72 (0.43)	5/7 (71.4)	-6.72 (0.42)	6/8 (75.0)	10	5.24 (1.74)	-4.22 (0.47)	7/9 (77.8)	-4.83 (0.50)	8/8 (100)	-4.39 (0.55)	9/9 (100)	
JNJ-3989 40 mg	15	7.57 (0.38)	-5.28 (0.25)	2/15 (13.3)	-5.89 (0.29)	8/14 (57.1)	-6.12 (0.29)	8/15 (53.3)	18	4.87 (0.30)	-3.84 (0.29)	15/16 (93.8)	-3.96 (0.36)	15/16 (93.8)	-4.02 (0.33)	16/16 (100.0)	
JNJ-3989 100 mg	14	7.77 (0.29)	-5.44 (0.28)	3/14 (21.4)	-6.41 (0.22)	8/14 (57.1)	-6.40 (0.25)	8/13 (61.5)	19	5.29 (0.33)	-4.08 (0.23)	15/18 (83.3)	-4.22 (0.36)	13/18 (72.2)	-4.31 (0.42)	13/16 (81.3)	
JNJ-3989 200 mg	16	8.29 (0.21)	-6.07 (0.16)	2/15 (13.3)	-6.56 (0.20)	3/14 (21.4)	-6.82 (0.26)	7/14 (50.0)	19	5.26 (0.35)	-4.16 (0.32)	14/18 (77.8)	-4.28 (0.35)	16/18 (88.9)	-4.24 (0.38)	15/17 (88.2)	
JNJ-3989 + JNJ-6379	13	8.04 (0.28)	-6.00 (0.19)	2/13 (15.4)	-6.72 (0.24)	6/12 (50.0)	-6.87 (0.27)	9/12 (75.0)	20	4.62 (0.24)	-3.63 (0.26)	18/19 (94.7)	-3.60 (0.29)	17/18 (94.4)	-3.71 (0.29)	18/18 (100)	

Values are mean (SE) unless otherwise noted.

Changes in HBcrAg

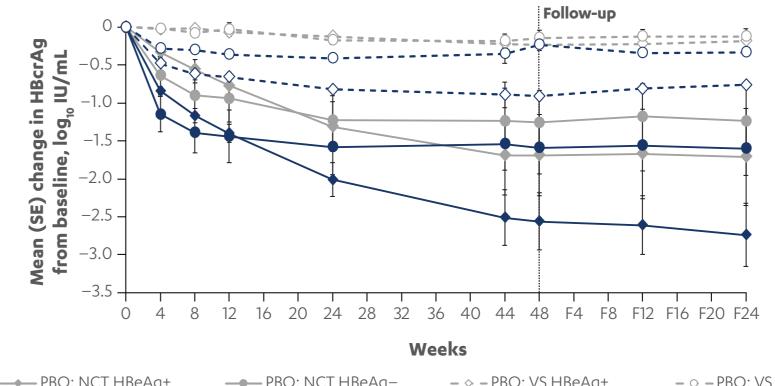
• JNJ-3989 200 mg resulted in pronounced reduction of HBcrAg in NCT patients, with up to 2.56 log₁₀ reduction at Week 48 in the HBeAg+ subgroup, while HBcrAg decline in VS patients was limited, potentially due to lower baseline levels (Figure 6 and Table 4)

• The proportion of patients reaching HBcrAg <LLOQ was greatest in the NCT and HBeAg – patients who received JNJ-3989 200 mg (Table 4)

*Presenting author.

---- PBO ---- JNJ-6379 ---- JNJ-3989 40 mg ---- JNJ-3989 100 mg ---- JNJ-3989 200 mg ---- JNJ-3989 100 mg + JNJ-6379

Figure 6. Mean (±SE) change in HBcrAg over time by treatment history and HBeAg status in patients with HBcrAg >LLOQ at baseline who received treatment with either PBO or JNJ-3989 200 mg.



-- PBO: NCT HBeAg- - - + PBO: VS HBeAg+ – o – PBO: VS HBeAg– → JNJ-3989: NCT HBeAg+ → JNJ-3989: NCT HBeAg- → JNJ-3989: VS HBeAg+ → JNJ-3989: VS HBeAg-

Table 4. Baseline and Change From Baseline Mean (SE) HBcrAg by Subgroup for NA and JNJ-3989 200 mg

 Treatment Arms in Patients With HBcrAg >LLOQ at Baseline

	N*	BL, log ₁₀ IU/mL	Change from BL at W24, log ₁₀ IU/mL	<lloq at="" w24,<br="">n/N (%)</lloq>	Change from BL at W48, log ₁₀ IU/mL	<lloq at="" w48,<br="">n/N (%)</lloq>	Change from BL at FU W24, log ₁₀ IU/mL	<lloq at<br="">FU W24, n/N (%)</lloq>
NCT HBeAg+								
PBO	7	7.76 (0.47)	-1.31 (0.33)	0/7	-1.69 (0.53)	0/7	-1.71 (0.59)	0/7
JNJ-3989 200 mg	16	8.43 (0.28)	-2.01 (0.18)	0/15	-2.56 (0.25)	0/14	-2.74 (0.30)	0/14
NCT HBeAg-								
PBO	8	4.8 (0.55)	-1.23 (0.32)	3/8 (37.5)	-1.26 (0.43)	3/8 (37.5)	-1.24 (0.45)	3/8 (37.5)
JNJ-3989 200 mg	16	4.71 (0.37)	-1.58 (0.37)	6/16 (37.5)	-1.59 (0.35)	7/16 (43.8)	-1.60 (0.36)	6/15 (40.0)
VS HBeAg+								
PBO	6	5.42 (0.14)	-0.12 (0.04)	0/6	-0.23 (0.08)	0/6	-0.18 (0.04)	0/6
JNJ-3989 200 mg	14	5.80 (0.22)	-0.82 (0.14)	0/14	-0.91 (0.17)	0/14	-0.76 (0.15)	0/14
VS HBeAg-								
PBO	11	3.73 (0.19)	-0.17 (0.06)	3/11 (27.3)	-0.14 (0.10)	1/11 (9.1)	-0.12 (0.10)	1/11 (9.1)
JNJ-3989 200 mg	22	3.92 (0.12)	-0.41 (0.07)	4/22 (18.2)	-0.23 (0.13)	2/22 (9.1)	-0.33 (0.09)	4/18 (22.2)

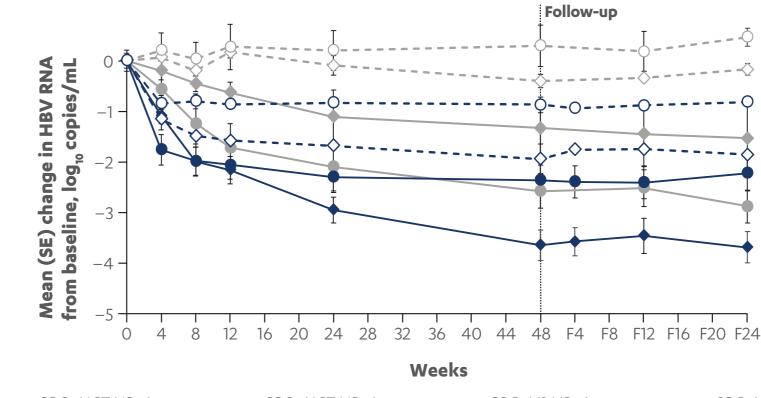
*Patients with detectable levels of HBcrAg at baseline Values are mean (SE) unless otherwise noted; $LLOQ = 3 \log_{10} IU/mL$.

Changes in HBV RNA

• JNJ-3989 200 mg resulted in pronounced reduction of HBV RNA across all patient populations, with up to 3.66 \log_{10} reduction in NCT HBeAg+ patients (**Figure 7** and **Table 5**)

- Reductions in HBV RNA were sustained through 24 weeks of follow-up
- Assessment of RNA reduction in VS patients was hampered by a high proportion of patients achieving HBV RNA <LOD during treatment

Figure 7. Mean (±SE) change in HBV RNA over time by treatment history and HBeAg status in patients with HBV RNA >LOD at baseline who received treatment with either PBO or JNJ-3989 200 mg.



→ PBO: NCT HBeAg – → PBO: VS HBeAg+ → JNJ-3989: NCT HBeAg+ → JNJ-3989: NCT HBeAg- - ◇ - JNJ-3989: VS HBeAg+ - ○ - JNJ-3989: VS HBeAg-

Table 5. Baseline and Change From Baseline Mean (SE) HBV RNA by Subgroup for PBO and JNJ-3989

 200 mg Troatmost Arms in Dationts With HPV/ DNA NOD at Pacelin

	N*	BL, log ₁₀ copies/mL	Change from BL at W24, log ₁₀ copies/mL	<lod at="" w24,<br="">n/N (%)</lod>	Change from BL at W48, log ₁₀ copies/mL	<lod at="" w48,<br="">n/N (%)</lod>	Change from BL at FU W24, log ₁₀ copies/mL	<lod at<br="">FU W24, n/N (%)</lod>
NCT HBeAg+								
PBO	6	6.90 (0.67)	-1.12 (0.53)	0/6	-1.34 (0.57)	0/6	-1.54 (0.62)	0/6
JNJ-3989 200 mg	16	7.33 (0.23)	-2.96 (0.26)	2/15 (13.3)	-3.66 (0.30)	4/14 (28.6)	-3.70 (0.31)	5/13 (38.5)
NCT HBeAg-								
PBO	5	5.02 (0.32)	-2.11 (0.47)	2/5 (40.0)	-2.59 (0.34)	3/5 (60.0)	-2.89 (0.33)	3/4 (75.0)
JNJ-3989 200 mg	13	4.64 (0.32)	-2.31 (0.30)	10/13 (76.9)	-2.37 (0.30)	11/13 (84.6)	-2.23 (0.36)	10/11 (90.9)
VS HBeAg+								
PBO	5	3.31 (0.47)	-0.10 (0.20)	1/5 (20.0)	-0.41 (0.14)	3/5 (60.0)	-0.18 (0.12)	1/5 (20.0)
JNJ-3989 200 mg	9	4.29 (0.44)	-1.69 (0.26)	5/9 (55.6)	-1.95 (0.35)	7/9 (77.8)	-1.87 (0.30)	6/8 (75.0)
VS HbeAg-								
PBO	3	3.25 (0.47)	0.20 (0.40)	1/3 (33.3)	0.29 (0.41)	1/3 (33.3)	0.46 (0.18)	0/2
JNJ-3989 200 mg	9	3.13 (0.14)	-0.84 (0.15)	8/9 (88.9)	-0.88 (0.17)	8/9 (88.9)	-0.75 (0.26)	5/7 (71.4)

Values are mean (SE) unless otherwise noted; N = patients with detectable HBV RNA at baseline; LOD = $2.49 \log_{10} \text{ copies/mL}$.



– \circ – PBO: VS HBeAg–

Key Findings

Similar to HBsAg, a JNJ-3989 dose-response relationship was observed for HBeAg, HBcrAg, HBV DNA, and HBV RNA The greatest reductions in viral markers were observed during the 48-week treatment phase and in patients who were NCT and HBeAg+ The reductions in HBeAg, HBcrAg, and HBV RNA generally remained stable or declined during the 24-week follow-up phase Safety results have been previously reported and showed that all regimens were generally safe and well tolerated³

Conclusions

JNJ-3989, with or without JNJ-6379, reduced all viral markers, with the strongest effect observed for HBsAg This study supports the continued development of JNJ-3989 in combination therapies that may provide functional cure to patients with CHB

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