Resistance and Pharmacokinetic/Pharmacodynamic Analyses of GS-1720, a Once-Weekly Oral Integrase Strand Transfer Inhibitor

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Conclusions

- No observed cases of emergent integrase (IN) resistance occurred after GS-1720 monotherapy treatment in this Phase 1b proof-of-concept study
- GS-1720 showed robust antiviral activity (≥1.5 log₁₀ copies/mL decline in HIV-1 RNA from baseline) at Day (D) 11 concentrations above inhibitory quotient (IQ) 2
- The pharmacokinetics (PK)/pharmacodynamics (PD) data and lack of observed resistance support further clinical development
- An oral combination regimen of once-weekly GS-1720 and GS-4182^a is being evaluated in Phase 2 studies among virologically suppressed and treatment-naïve people with HIV-1 (PWH)^b

Plain Language Summary

- GS-1720 is a medicine that is being studied to treat HIV, but it is not yet approved for people to take outside of a clinical trial
 - GS-1720 can be taken just once a week unlike many other HIV medicines that need to be taken every day
- We tested many doses of GS-1720 in people with HIV to see how well it works and to study if the HIV virus developed any changes that helped It resist the effects of GS-1720 (called treatment resistance)
- We found that GS-1720 worked well to treat HIV, and people who took GS-1720 did not develop resistance to this medicine
- We are planning more studies to test if GS-1720 combined with another drug called GS-4182 can be taken once a week to treat people with HIV

Background

- Adherence to HIV-1 treatment reduces the risk of virologic failure, yet can be challenging; an unmet need for long-acting antiretroviral therapies (ART) remains^{1,2}
- GS-1720 is an oral integrase strand transfer inhibitor (INSTI) with potent anti-HIV-1 activity and a median half-life of 9.3 days, supportive of weekly dosing³
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- In a Phase 1b study in PWH, GS-1720 resulted in a mean decline in HIV-1 RNA from baseline to D11 of >2 log₁₀ copies/mL in three of the four GS-1720 dose cohorts, with a 1.74 log₁₀ copies/mL decline observed in the lowest dose cohort⁴

Objective

To assess GS-1720 resistance and PK/PD from the current Phase 1b study

Methods

- This Phase 1b, open-label, multicohort substudy enrolled PWH^c
- Participants with detectable HIV-1 viral load were enrolled into four cohorts (n=7/cohort) and administered oral GS-1720 on D1 and D2, then switched on D11 to bictegravir/emtricitabine/tenofovir alafenamide (B/F/TAF) or an alternative standard of care (SOC) ART regimen (Figure 1)
- Participants were tested for genotypic and phenotypic IN resistance at baseline (screening visit) and D11 using the GeneSeq[®] Integrase^d and PhenoSense[®] Integrase^e assays^f
- Participants with suboptimal virologic response (SVR) after D11, defined as HIV-1 RNA ≥50 copies/mL and <1 log₁₀ HIV-1 RNA reduction from D11, qualified for further genotypic and phenotypic resistance testing
- Intensive PK sampling was collected on D1 and D2 up to 12 hours post-dose, followed by single anytime plasma PK sampling throughout the study
- GS-1720 plasma concentrations were quantified using a validated high-performance liquid chromatography-tandem mass spectrometry bioanalytical method

Figure 1. Study Design

Key eligibility criteria:	30 mg (n=7) B/F/TA	F or SOC
Aged 18–65 years	150 mg (n=	7) B/F/TA	F or SOC
• HIV-1 RNA 5000–≤400,000 cpm	450 mg (n=	7) B/F/TA	F or SOC
 CD4+ T-cells >200 cells/µL 	900 mg (n=	7) B/F/TA	F or SOC
 Treatment naïve OR treatment experienced, but naïve to INSTIs and off ART for ≥12 weeks 	D1 2 GS-1720 Oral	I 11 ▲ Primary Endpoint	60 ▲ Last Study
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ART, antiretroviral therapy; B/F/TAF, bictegravir/emtricitabine/tenofovir alafenamide; cpm, copies/mL; D, day; INSTI, integrase strand transfer inhibitor; SOC, standard of care

Administration

Results

Viral Load Decline

- One participant in the lowest dose cohort had virologic rebound during the monotherapy period
- Maximum HIV-1 RNA reduction was –1.11 log₁₀ copies/mL at D7, increasing to –0.26 log₁₀ copies/mL at D11, with resuppression to <50 copies/mL on B/F/TAF
- No INSTI resistance associated mutations (RAMs) or phenotypic changes to licensed INSTIs were detected at D11 By D60, 24/27 participants had HIV-1 RNA <50 copies/mL
- The three participants who did not reach HIV-1 RNA <50 copies/mL included one participant in the GS-1720 150 mg cohort and two participants in the GS-1720 450 mg cohort
- All participants had considerable viral load decrease from baseline

Resistance Analysis

- All participants were phenotypically susceptible to GS-1720 and INSTIs at baseline
- No primary INSTI RAMs were observed (Table 2)
- Secondary INSTI RAMs were detected, with no impact on phenotypic sensitivity (Table 3)
- No treatment-emergent resistance to the INSTI class was detected at D11 or in those with SVR (Tables 2 and 3)
 Phenotyping results for all participants at D11 demonstrated sustained susceptibility after monotherapy dosing relative to the wild-type for bictegravir, dolutegravir, elvitegravir, raltegravir, and GS-1720
- No treatment-emergent primary RAMs to the INSTI class were detected at D11 or follow-up (Table 2)
- No treatment-emergent secondary mutations to the INSTI class were detected at D11 or follow-up (Table 3)

Table 2. Participants With Primary INSTI RAMs

	Participants with Primary INSTI RAMs				
GS-1720 dose	Baseline D11ª		Follow-up in Participants with SVR ^b	Treatment-Emergent Resistance	
30 mg (n=7)	0	0	-	0	
150 mg (n=7)	0	0	0	0	
450 mg (n=7)	0	0	0	0	
900 mg (n=7)	0	0	-	0	

^aD11 samples were analysed for 27 participants. Genotypic data were obtained from 20/27 participants (74%) and phenotypic data were obtained from 12/27 participants (44%). ^bTwo participants in the 150 mg group and two participants in the 450 mg group qualified for further INSTI genotypic and phenotypic resistance testing. D, day; INSTI, integrase strand transfer inhibitor resistance; RAM, resistance-associated mutation; SVR, suboptimal virologic response.

Table 3. Participants With Secondary INSTI RAMs

	Participants with Secondary INSTI RAMs (n)				
GS-1720 dose	Baseline	D11ª	Follow-up in Participants with SVR ^b	Treatment-Emergent Resistance	
30 mg (n=7)	M50I (1) S119P/R/T (3) V151A/I/L (1) E157K/Q (1)	M50I (1) S119P/R/T (1) V151A/I/L (1) E157K/Q (1)	-	0	
150 mg (n=7)	0	0	0	0	
450 mg (n=7)	M50I (3) S119P/R/T (2)	M50I (3) S119P/R/T (2)	M50I (1) S119P/R/T (1)	0	
900 mg (n=7)	M50I (2) S119P/R/T(2)	M50I (1) S119P/R/T (1)	-	0	

^aD11 samples were analysed for 27 participants. Genotypic data were obtained from 20/27 participants (74%) and phenotypic data were obtained from 12/27 participants (44%). ^bTwo participants in the 150 mg group and two participants in the 450 mg group qualified for further INSTI genotypic and phenotypic resistance testing. D, day; INSTI, integrase strand transfer inhibitor resistance; RAM, resistance associated mutation; SVR, suboptimal virologic response.

PK/PD Analysis

- Mean GS-1720 concentrations at D11 and HIV-1 RNA reductions from baseline to D11 are displayed in Table 4
- At D11, participants with GS-1720 concentrations above two-fold the IQ (IQ2; 3.876 µg/mL) showed robust antiviral activity of ≥1.5 log₁₀ copies/mL reduction in HIV-1 RNA from baseline (Figure 2)

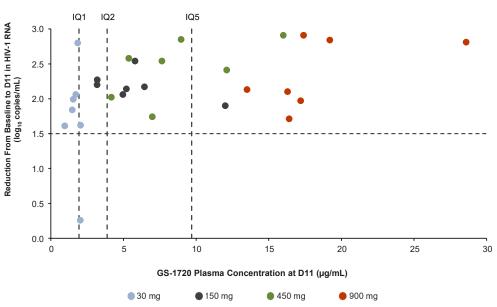
Table 4. Mean GS-1720 Concentrations and HIV-1 RNA Reductions From Baseline at D11

GS-1720 dose	Mean GS-1720 Concentrations at D11 (µg/mL)	Relative Mean IQ Values at D11 (µg/mL)	Mean Reduction From Baseline in HIV-1 RNA at D11 (log ₁₀ copies/mL)	
30 mg (n=7)	1.64	0.8	1.74	
150 mg (n=7)	=7) 5.87 3.0		2.18	
450 mg (n=7)	8.78	4.5	2.44	
900 mg (n=7)	18.4	9.5	2.37	

D, day; IQ, inhibitory quotient.

Visit

Figure 2. PK/PD Analysis of GS-1720 at D11



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Study Population

- Twenty-eight participants were enrolled
- Median (range) age was 33 (18–62) years, and 10.7% were female (Table 1)

Table 1. Baseline Characteristics

	30 mg (n=7)	150 mg (n=7)	450 mg (n=7)	900 mg (n=7)	Total participants (N=28)
Median (range) age, years	35 (18–55)	38 (27–61)	28 (25–62)	31 (24–43)	33 (18–62)
Female sex at birth, n (%)	1 (14.3)	1 (14.3)	1 (14.3)	0	3 (10.7)
Race, n (%) American Indian/Alaska Native Asian Black Native Hawaiian/Pacific Islander White Other	1 (14.3) 0 3 (42.9) 0 1 (14.3) 2 (28.6)	1 (14.3) 0 3 (42.9) 0 3 (42.9) 0	0 0 2 (28.6) 3 (42.9) 2 (28.6)	0 2 (28.6) 0 2 (28.6) 3 (42.9)	2 (7.1) 2 (7.1) 6 (21.4) 2 (7.1) 9 (32.1) 7 (25.0)
Ethnicity, n (%) Hispanic or Latinx	4 (57.1)	2 (28.6)	2 (28.6)	5 (71.4)	13 (46.4)
Median (Q1–Q3) HIV-1 RNA, log ₁₀ copies/mL	4.36 (4.08–5.19)	4.74 (4.55–4.98)	5.31 (5.12–5.42)	4.90 (4.51–5.29)	4.90 (4.48–5.30)
Median (Q1–Q3) CD4+ T-cells/µL	454 (334–505)	264 (194–389)	350 (336–430)	440 (276–475)	370 (275–450)
ART naïve, n (%)	6 (85.7)	4 (57.1)	6 (85.7)	6 (85.7)	22 (78.6)

ART, antiretroviral therapy; Q, quartile

Footnotes "Results from the GS-1182 Phase 1a study are presented in the HIV Glasgow 2024 poster #P036 (accessible via QR code)^{1,} *NCT06544733 includes virologically suppressed PWH; NCT06613685 includes treatment-naive PWH; *NCT0558507, *GeneSeq0 Integrase sequences the N gene to identify known RAM to the INSTI class. *PhenoSense® Integrase determines the phenotypic sensitivity to all currently approved antifetrovinal class (biologizer), doublegravir, advelogravir, advelogravity and GS+172, *Morgania Biosciences, South Sam Francisco, California, USA.

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Author Disclosures: Brie Falkard, Haeyoung Zhang, Mutaz Jaber, Eva Mortensen, Furong Wang, Christian Callebaut, and Dhananjay D. Marathe are all employees and shareholders of Glead Sciences, Inc.

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