

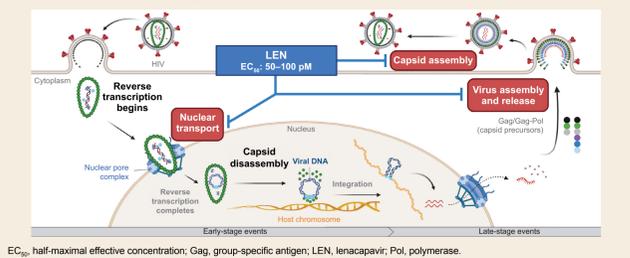


Resistance Analysis of Long-Acting Lenacapavir in Highly Treatment-Experienced People With HIV After 52 Weeks of Treatment

Nicolas Margot, Laurie VanderVeen, Vidula Naik, Hadas Dvory-Sobol, Martin S. Rhee, Christian Callebaut — Gilead Sciences, Inc., Foster City, California, USA

Introduction

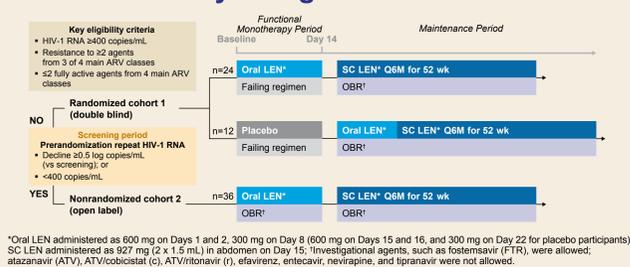
Lenacapavir Targets Multiple Stages of HIV Replication Cycle^{1,2}



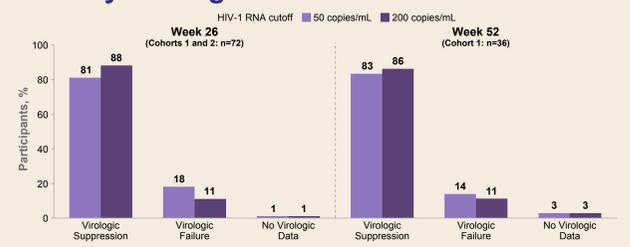
LEN: Long-Acting Inhibitor of HIV-1 Capsid

- Highly potent activity (EC₅₀: 50–100 pM), with low clearance and slow release kinetics³
 - Can be administered orally (daily or weekly) or subcutaneously (SC) every 6 months (Q6M)³⁻⁵
- Fully active against HIV with resistance to existing antiretroviral (ARV) classes (nucleoside reverse transcriptase inhibitors [NRTIs], non-NRTIs [NNRTIs], protease inhibitors [PIs], integrase strand transfer inhibitors [INSTIs], and entry inhibitors)^{1,6,7}
- In vitro selected resistance-associated mutations (RAMs; L56I, M66I, Q67H, K70N, N74D/S, and T107N) had low replication capacity, except Q67H^{1,8}
- In viremic heavily treatment-experienced people with HIV (PWH) with multidrug resistance (CAPELLA study; ClinicalTrials.gov NCT04150068⁹⁻¹¹), LEN in combination with an optimized background regimen (OBR) led to 83% (n=30/36) virologic suppression at Week 52¹²
- In treatment naïve PWH (CALIBRATE study; ClinicalTrials.gov NCT04143594)^{13,14}:
 - SC LEN, initially in combination with emtricitabine (FTC)/tenofovir alafenamide (F/TAF) and later with oral TAF or bictegravir (BIC), was well tolerated, and achieved and maintained high rates of virologic suppression through 1 year (90% and 85%, respectively)
 - Oral LEN in combination with F/TAF had similar efficacy (85%)

CAPELLA Study Design⁹⁻¹¹



Efficacy Through Weeks 26 and 52¹²



- Durable efficacy through Week 52
- Cohort 1 (randomized): mean increase in CD4 of 83 cells/μL at Week 52
- Cohort 2 (nonrandomized): mean increase in CD4 of 98 cells/μL at Week 26

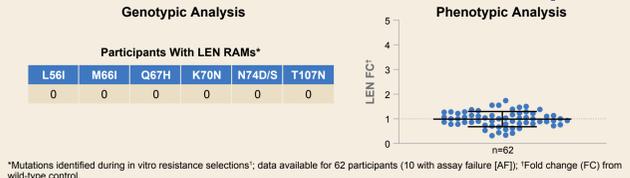
Baseline Class Resistance¹⁵

4 Main ARV Classes

Resistance Within Classes	Participants, n (%)			
	Cohort 1 n=36	Cohort 2 n=36	All n=72	
NRTI	17 (47)	16 (44)	33 (46)	
NNRTI	9 (25)	13 (36)	22 (31)	
PI	8 (22)	5 (14)	13 (18)	
INSTI	2 (6)	0	2 (3)	
*M184V/I alone was not sufficient to fulfill NRTI-resistance (R) criteria in study; 11 participant had INSTI-R and NNRTI-R, with only partial resistance to 1 NRTI (didanosine) in presence of M184V/I (FTC-R and lamivudine [3TC]-R not counting in total NRTI-R when M184V/I was present per protocol requirement).				

- Entry Criteria:** resistance to ≥2 ARVs in ≥3 of 4 main ARV classes
- Nearly half of all participants had resistance to all 4 classes

Absence of Baseline Resistance to Lenacapavir¹⁵



- Evaluated with PhenoSense[®] Gag-Pro (Monogram Biosciences, South San Francisco, California, USA)
 - No LEN-R mutations detected
 - Wild-type LEN phenotypic susceptibility: mean FC = 1.0 (range 0.3–1.7)

Objective

- To describe the resistance analyses conducted through Week 52 in CAPELLA

Methods

Virologic Failure (VF) Criteria

- Suboptimal virologic response: confirmed HIV-1 RNA ≥50 copies/mL and <1-log₁₀ ↓ from LEN start (assessed at Week 4)
- Virologic rebound: after suppression, confirmed HIV-1 RNA ≥50 copies/mL or >1-log₁₀ ↑ from nadir
- Viremia at last visit

On-Treatment Resistance Analyses: Participants With VF

- Initial or confirmatory VF visit analyzed for capsid protein (CA) resistance: PhenoSense Gag-Pro (genotypic and phenotypic assay) and alternate deep-sequencing assay (Seq-IT GmbH & Co. KG, Kaiserslautern, Germany) used for retest samples
- Confirmatory VF visit analyzed for protease (PR), reverse transcriptase (RT), and integrase (IN) resistance: PhenoSense GT, GenoSure[®] MG, PhenoSense Integrase, and GeneSeq[®] Integrase (Monogram)
- Drug plasma concentrations measured using liquid chromatography–tandem mass spectrometry

Results

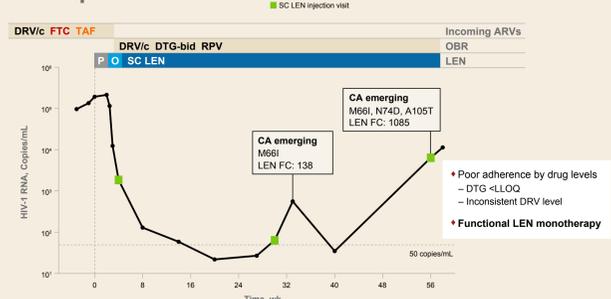
Postbaseline Resistance Analysis Week 52 Interim Analysis

Resistance Category, n (%)	Cohort 1: n=36	Cohort 2: n=36	All: N=72
Resistance analysis population	11 (31)	10 (28)	21 (29)
With data	11 (31)	9 (25)	20 (28)
Resuppressed <50 copies/mL	4 (11)	4 (11)	8 (11)
With CA-R emerging	4 (11)	4 (11)	8 (11)
M66I, n	4	2	6
Q67H/K70N, n	1	2	3
K70H/N74D, n	1	3	4
N74D, n	3	0	3
A105S/T, n	3	1	4
T107A/C/N, n*	1	3	4
No CA-R emergence	7 (19)	5 (14)	12 (17)

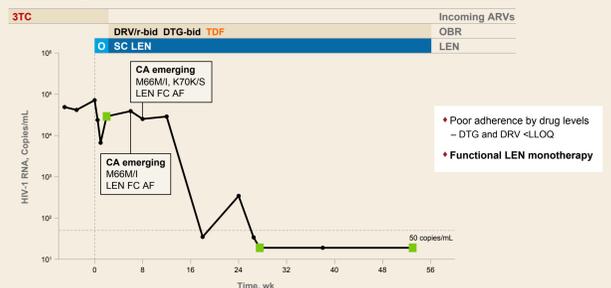
*1 participant had emergent T107A mutation in CA with no loss in LEN susceptibility before achieving HIV-1 RNA suppression; participant was not categorized as having CA-R emergence.

- 21 of 72 participants were analyzed for resistance
- 8 of 72 participants had CA-R emerging by Week 52, with no change since Week 26 interim analysis

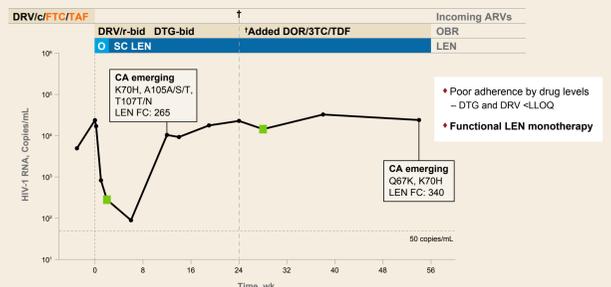
Viral Response and Resistance: OBR Adherence Issue* Participant 1



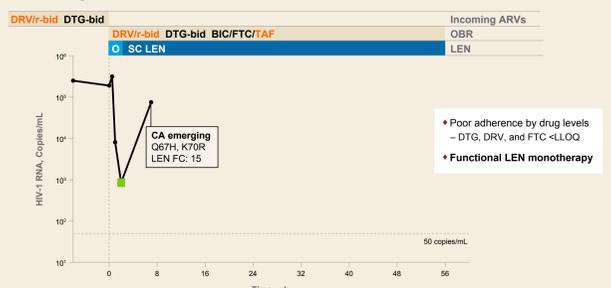
Participant 2



Participant 3

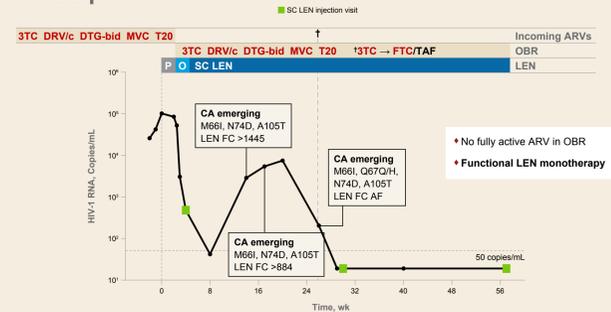


Participant 4

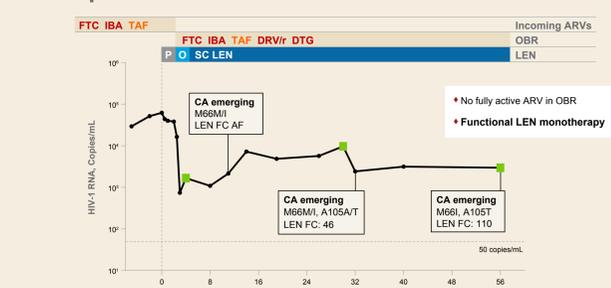


*Drugs in red are not active (OSS: 0); drugs in orange are partially active (OSS: 0.5); drugs in black are fully active (OSS: 1). DOR, doravirine; DRV, darunavir; DTG, dolutegravir; LLOQ, lower limit of quantification; O, oral; P, placebo; RPV, rilpivirine; TDF, tenofovir disoproxil fumarate.

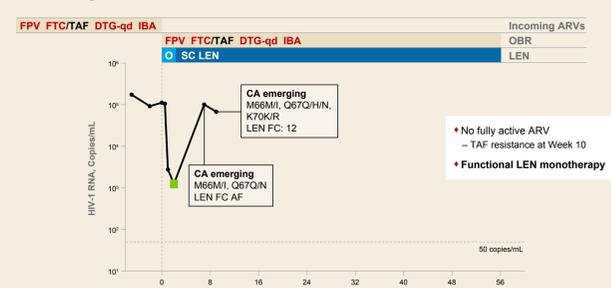
Viral Response and Resistance: No Fully Active ARV in OBR* Participant 5



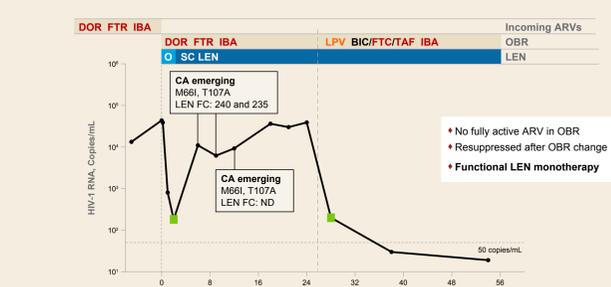
Participant 6



Participant 7



Participant 8



*Drugs in red are not active (OSS: 0); drugs in orange are partially active (OSS: 0.5); drugs in black are fully active (OSS: 1). FPV, fosamprenavir; IBA, balizumab; LPV, lopinavir; MVC, maraviroc; ND, not determined; T20, enfuvirtide.

Summary of Participants With CA Resistance

Participant	1st Visit With CA-R	CA RAMs	LEN FC	No. of Fully Active Drugs	Reason for Functional LEN Monotherapy
1	Week 26	M66I	138	3	
2	Week 4	M66M/I, K70K/S	ND	2	OBR adherence issue*
3	Week 10	K70H, A105A/S/T, T107T/N	265	2	
4	Week 4	Q67H, K70R	15	3	
5	Week 10	M66I, N74D, A105T	>1445	0	
6	Week 4	M66M/I	46	0	No fully active ARVs
7	Week 4	M66M/I, Q67Q/H/N, K70K/R	12	1 [†]	in OBR
8	Week 4	M66I, T107A	240	0	

*Adherence based on drug plasma concentrations; [†]TAF was fully active at baseline (PhenoSense GT), but was inactive at Week 10 (GenoSure).

Conclusions

- In heavily treatment-experienced PWH with baseline multidrug resistance:
 - LEN + OBR led to durable high rates of virologic suppression (83%) and increases in CD4 count through Week 52
 - LEN was well tolerated, with no adverse events leading to discontinuation
- Postbaseline emergence of LEN RAMs was observed in 8 of 72 participants
- All 8 cases with LEN RAM emergence were associated with functional LEN monotherapy at the time of resistance emergence
- Absence of postbaseline emergence of LEN RAMs was observed in 12 of 72 participants with viremia
- Minimal emergence of resistance to OBR drugs was observed, with no adverse virologic effects (participants resuppressed after OBR drug-resistance emergence)

References: 1. Link JO, et al. Nature 2020;584:614-8. 2. Zila V, et al. Cell 2021;184:1032-46. 3. Begley R, et al. AIDS 2020, abstr PEB0265. 4. Begley R, et al. CROI 2020, abstr 470. 5. Daar EM, et al. CROI 2020, poster 3691. 6. Margot N, et al. Antimicrob Agents Chemother 2021;65:e02057-20. 7. Margot N, et al. CROI 2022, abstr 508. 8. Margot N, et al. J Antimicrob Chemother 2022;77:989-995. 9. Molina J-M, et al. IAS 2021, abstr CALX01LB02. 10. Segal-Maurer S, et al. CROI 2021, abstr 127. 11. Segal-Maurer S, et al. N Engl J Med 2022;386:1793-803. 12. Ogibagu O, et al. CROI 2022, abstr 1047. 13. Gupta SK, et al. CROI 2022, abstr 138. 14. Gupta SK, et al. IAS 2021, abstr OALB0302. 15. Margot N, et al. EACS 2021, oral OS1/1. Acknowledgments: We are grateful to all the individuals who participated in this trial, and their partners and families. Participating study investigators and their study teams: Canada: J Brunetto, B Trotter, Dominican Republic: E Koenig, France: J-M Molina, S Ronot-Bregiere, Y Yazdanzhanah, Germany: H-J Stellbrink, Italy: A Antinori, A Castagna, F Castelli, Japan: T Shirasaka, Y Yokomaku, South Africa: M Passolunghi, Spain: J Mellado, Taiwan: C-C Hung, Thailand: A Avihingsanon, P Chetchotisakd, K Sirigum, W Ratanasumrit, USA: DS Berger, M Berne, C Brown, CM Creticos, GE Crofoot, E DeJesus, D Higgins, T Hodge, K Lichtenstein, J McCowan, O Ogbagu, O Ojayemi, GJ Richmond, MN Rampopal, PJ Ruane, W Sanchez, S Segal-Maurer, J Sims, GJ Sinclair, DA Wheeler, A Wozniak, K Workowski, C Zuzawski. We also thank Monogram Biosciences for resistance analyses and Seq-IT for sequence analyses. This study was funded by Gilead Sciences, Inc. Editing and production assistance were provided by BioScience Communications, New York, New York, USA, funded by Gilead.