



Terapia dei pazienti multifalliti

Stefano Rusconi

Sistema Socio Sanitario
 Regione
Lombardia
ASST Ovest Milanese

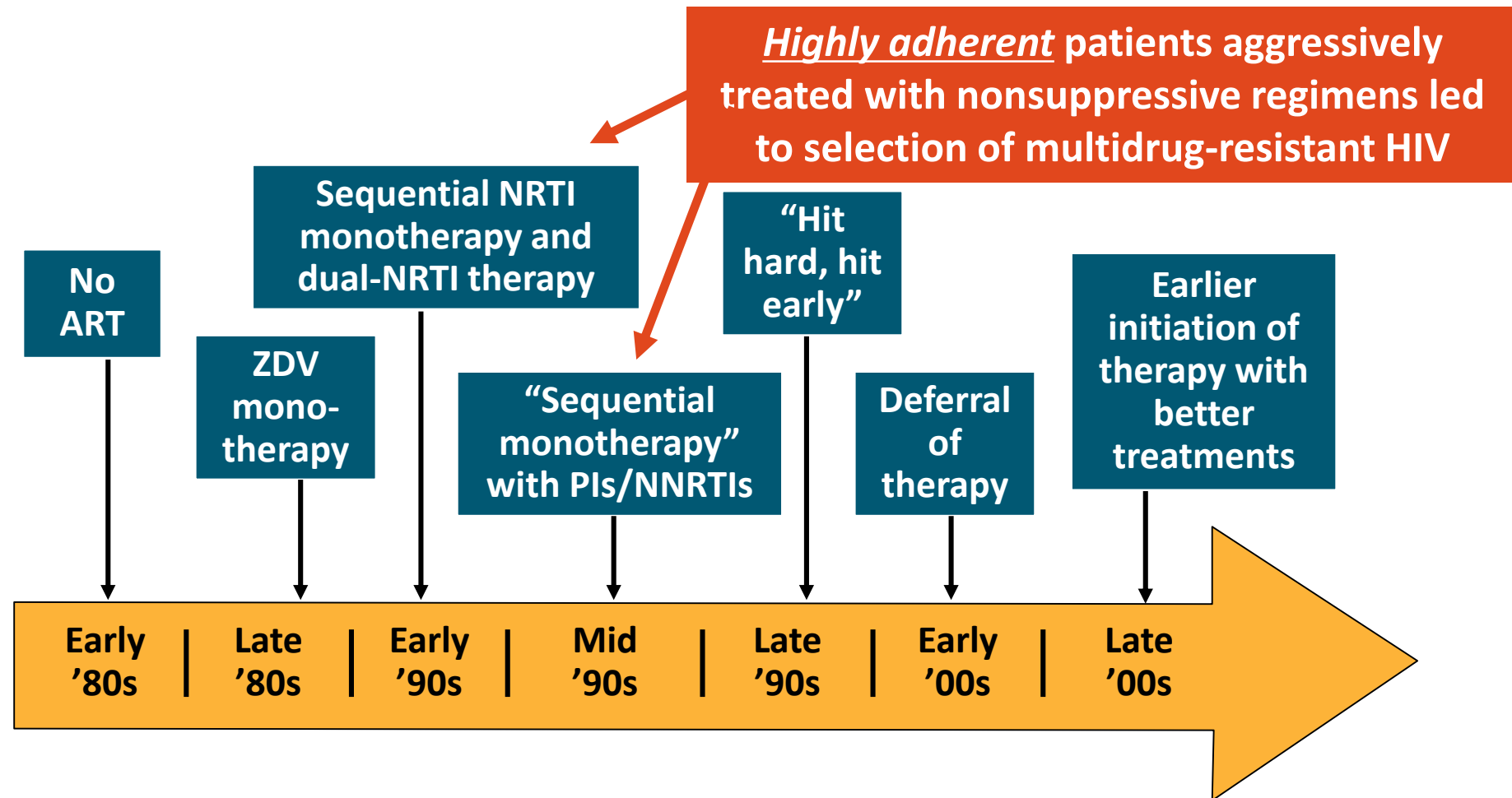


COIs

*GSK, Gilead, Janssen-Cilag, MSD,
ViiV Healthcare, Theratechnologies*

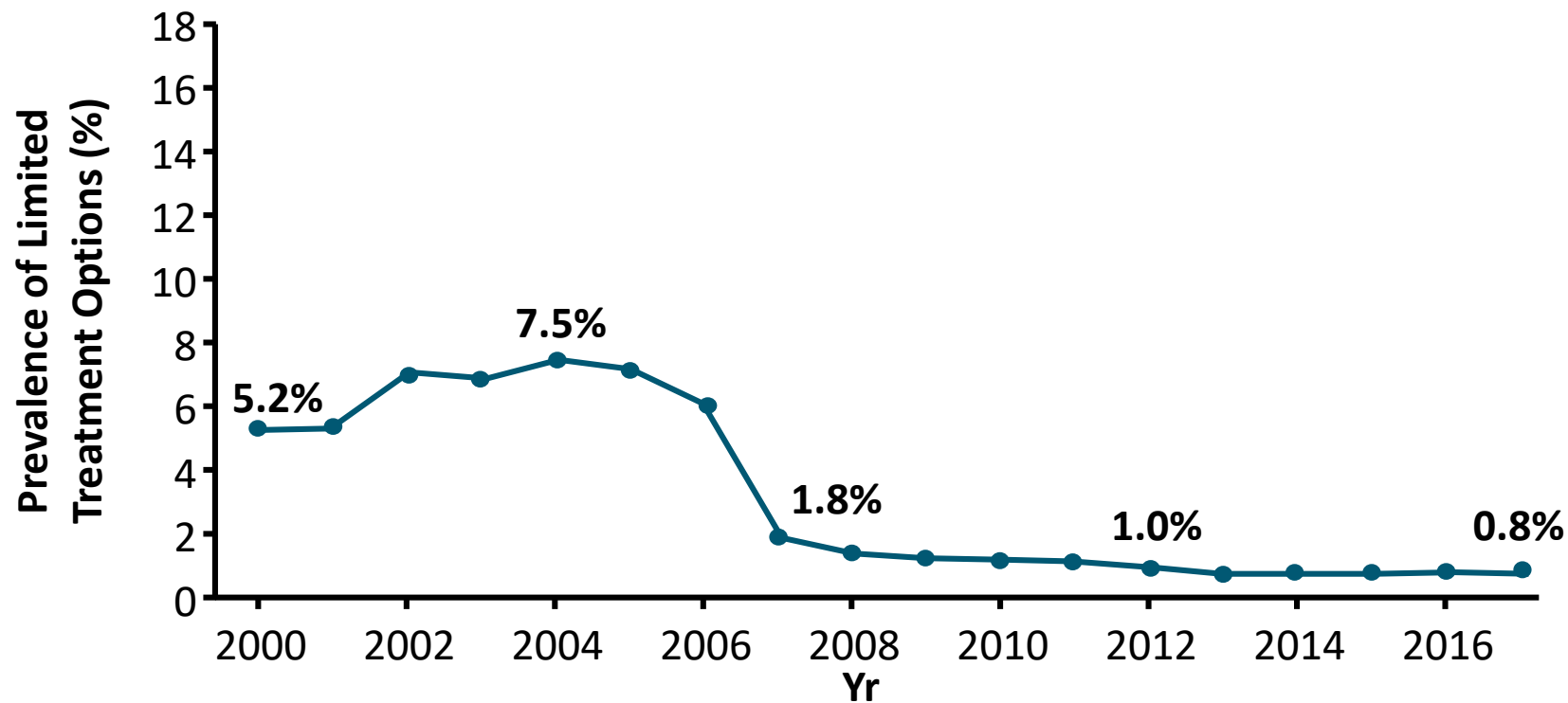


Who Are the People With Multidrug Resistant HIV?

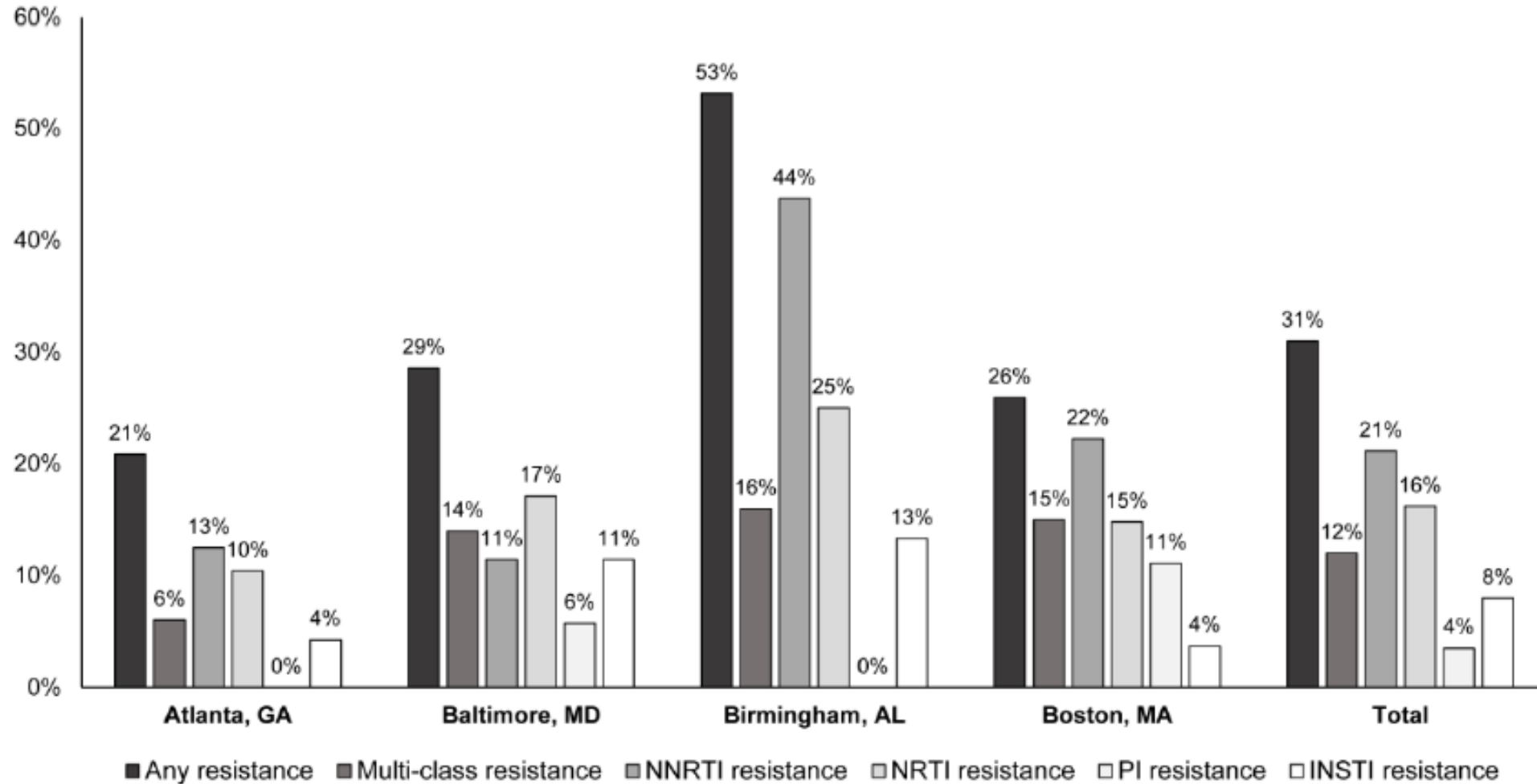


Prevalence of Heavily Treatment–Experienced Patients With Multiclass Resistance/Limited Treatment Options

- CNICS cohort of > 26,000 ART-experienced people with HIV receiving care in the US
- Limited treatment options defined as ≤ 2 available classes with ≤ 2 active drugs per class by resistance testing



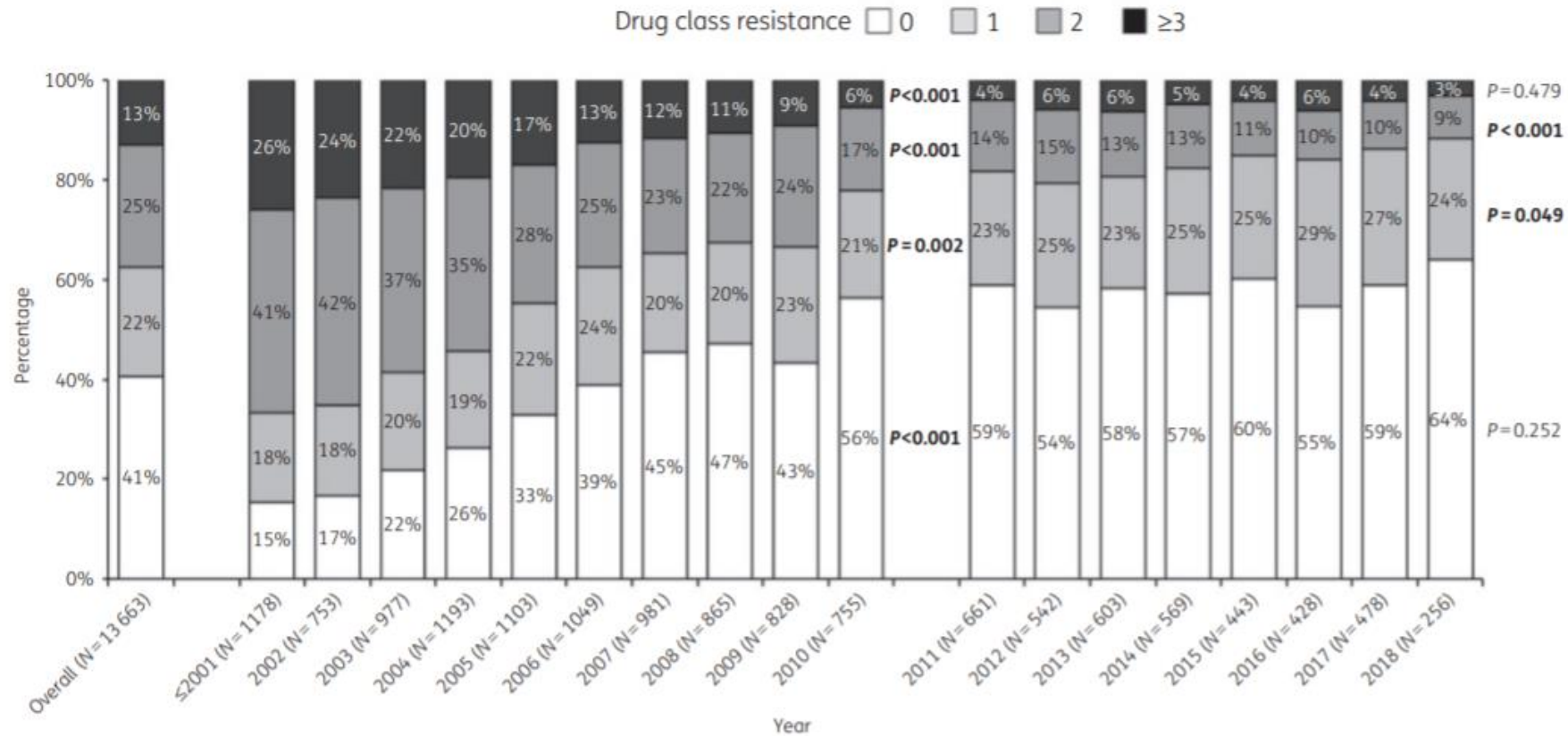
HIV drug resistance in a cohort of HIV-infected MSM in the United States



Jessica M. Fogel et al. AIDS 2020, 34:91–101

Heavily Treatment-Experienced (HTE) patients

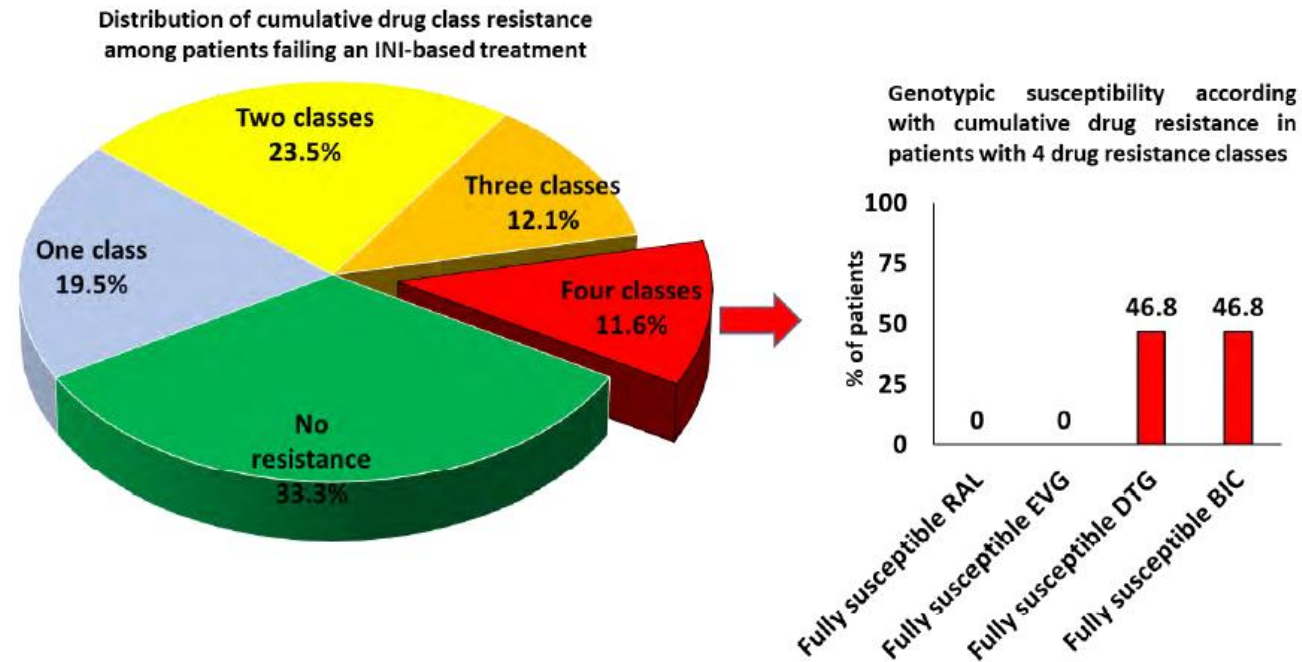
HIV MDR is still a relevant issue despite its dramatic drop over the years



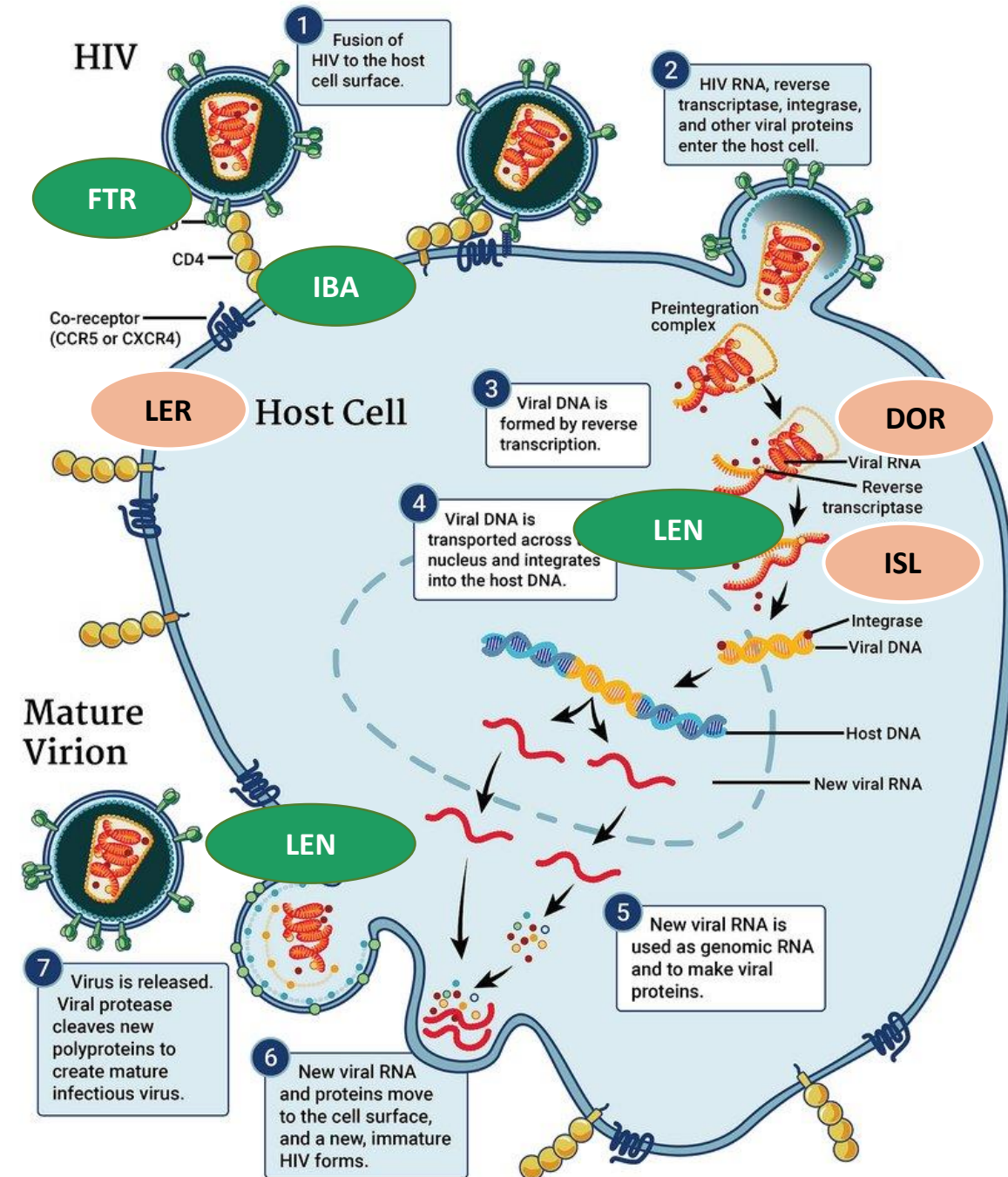
Heavily Treatment-Experienced (HTE) patients

INTEGRASE INHIBITOR RESISTANCE DYNAMICS FROM 2007 TO 2017 IN ITALIAN CLINICAL ISOLATES

Among 405 INI-failing patients, 47 (11.9%) patients showed cumulative four-drug class resistance. Of them, only 22 (46.8%) showed full GS to DTG or BIC.



HIV life cycle and upcoming drugs for MDR HIV



• Novel drugs

- NNRTI: Doravirine* (**DOR**)
- NRTI: Islatravir (**ISL**)
- Anti-CCR5: Leronlimab (**LER**)

• Novel classes

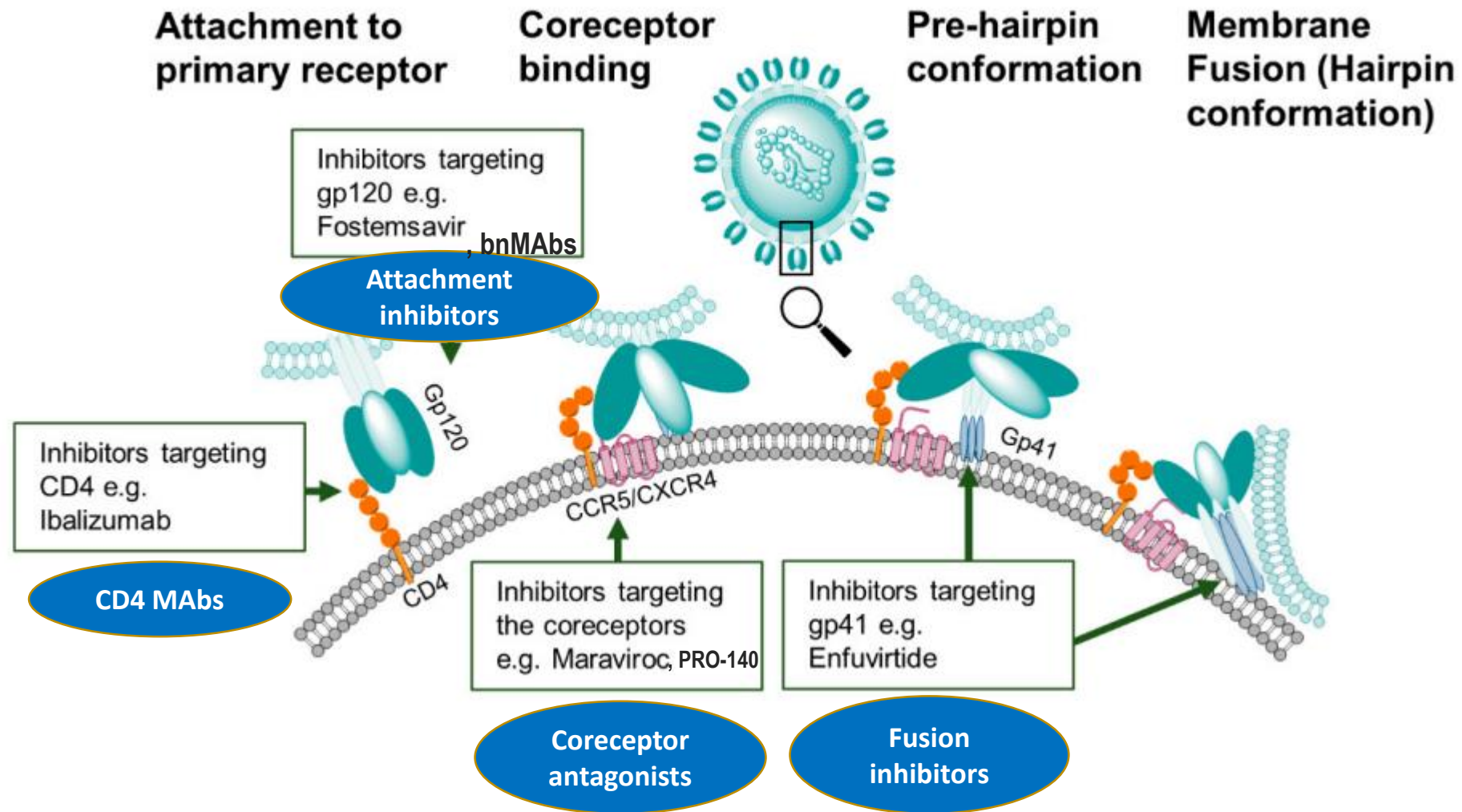
- Anti-CD4: Ibalizumab* (**IBA**)
- Anti-gp120: Fostemsavir* (**FTR**)
- Capsid inhibitors: Lenacapavir (**LEN**)

New options for MDR HIV

Genotypic interpretation

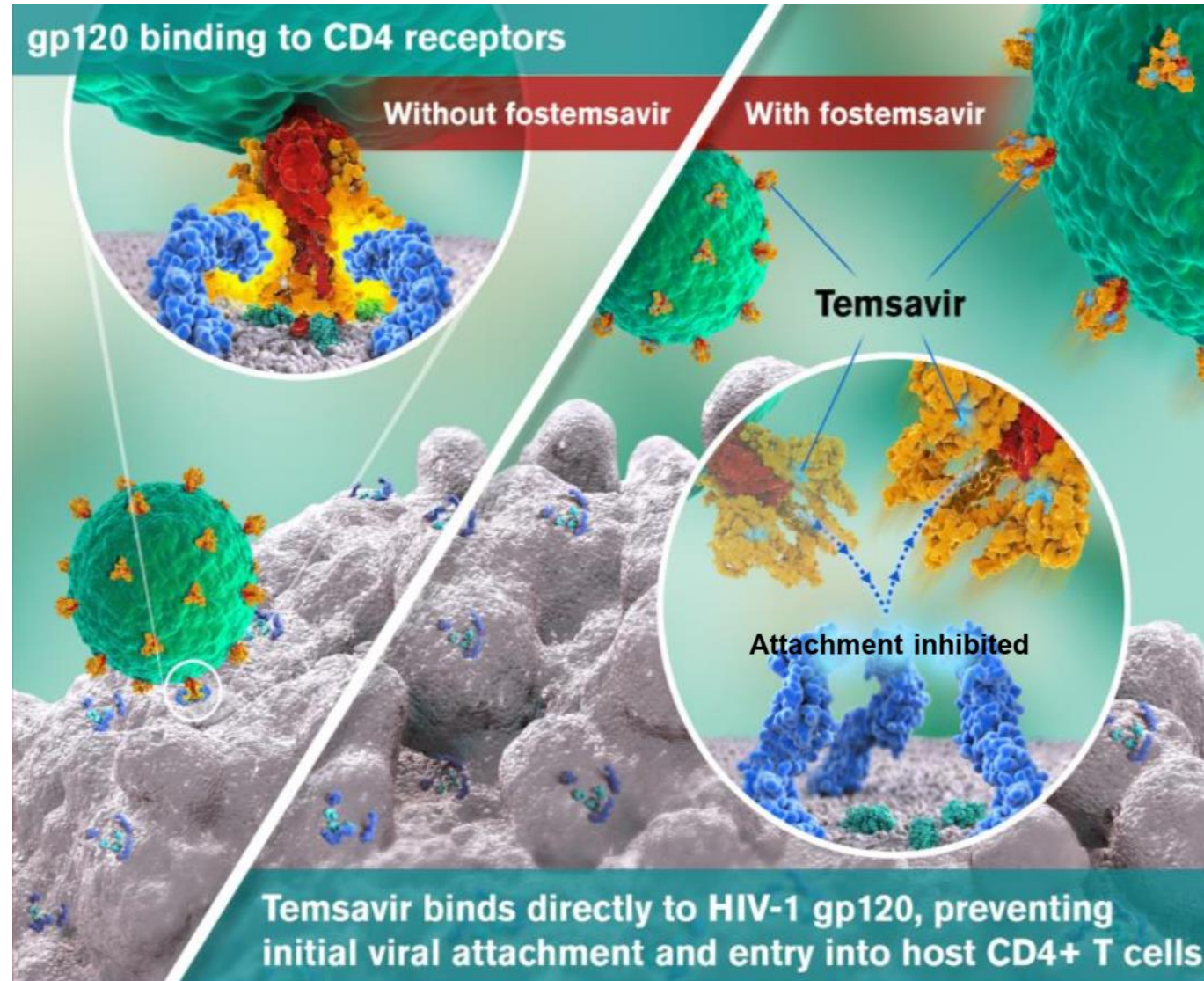
Drug	Target	Genotypic interpretation	Comments
<u>Doravirine</u> *	RT	Available	Role of minor RAMs possibly underestimated
Islatravir	RT	Not available	Signature RAMs and cross-resistance to be further defined
Leronlimab	CCR5 (host)	Available (geno2pheno)	Resistance through use of Leronlimab-bound CCR5, similar to MVC, not defined
<u>Ibalizumab</u> *	CD4 (host)	Not available	Most resistant strain occur through use of ibalizumab-bound CD4 following loss of N-glycosylation site(s) in gp120 V5 domain
<u>Fostemsavir</u> *	Gp120	Not available	Signature RAMs to be further defined, genotype and phenotype poorly predictive of treatment response
bnMAbs	Gp120	Not available	Resistance hardly predictable by genotype
Lenacapavir	Capsid	Not available	Signature RAMs to be further defined

HIV entry steps and entry inhibitors



New options for MDR HIV

Fostemsavir → targets HIV gp120



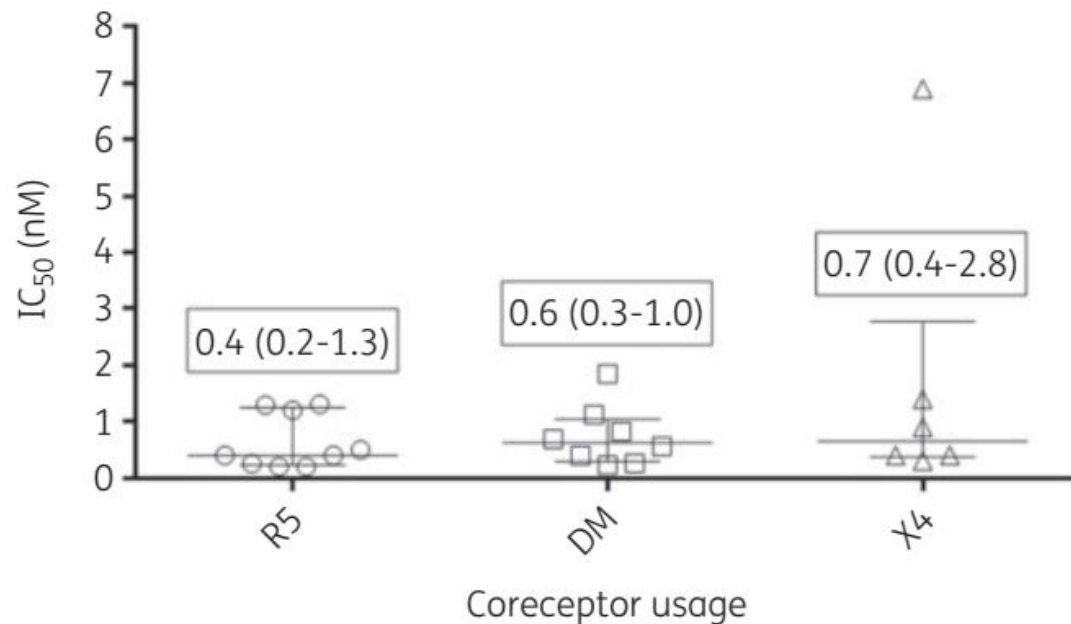
Susceptibility of global HIV-1 clinical isolates to fostemsavir using the PhenoSense Entry assay

Subtype	#	IC ₅₀ <1 nM (%)	IC ₅₀ >1 to <10 nM (%)	IC ₅₀ >10 to <100 nM (%)	IC ₅₀ >100 nM (%)
All	1337	719 (53.8)	352 (26.3)	146 (10.9)	120 (9.0)
B	881	544 (61.7)	196 (22.2)	85 (9.6)	56 (6.4)
C	156	78 (50.0)	56 (35.9)	15 (9.6)	7 (4.5)
A1	17	5 (29.4)	11 (64.7)	0 (0)	1 (5.9)
A	43	10 (23.3)	21 (48.8)	8 (18.6)	4 (9.3)
BF	19	5 (26.3)	8 (42.1)	2 (10.5)	4 (21.1)
F1	48	8 (16.7)	15 (31.3)	13 (27.1)	12 (25.0)
B, F1	29	4 (13.8)	9 (31.0)	5 (17.2)	11 (37.9)
CRF01_AE	5	0 (0)	0 (0)	0 (0)	5 (100)

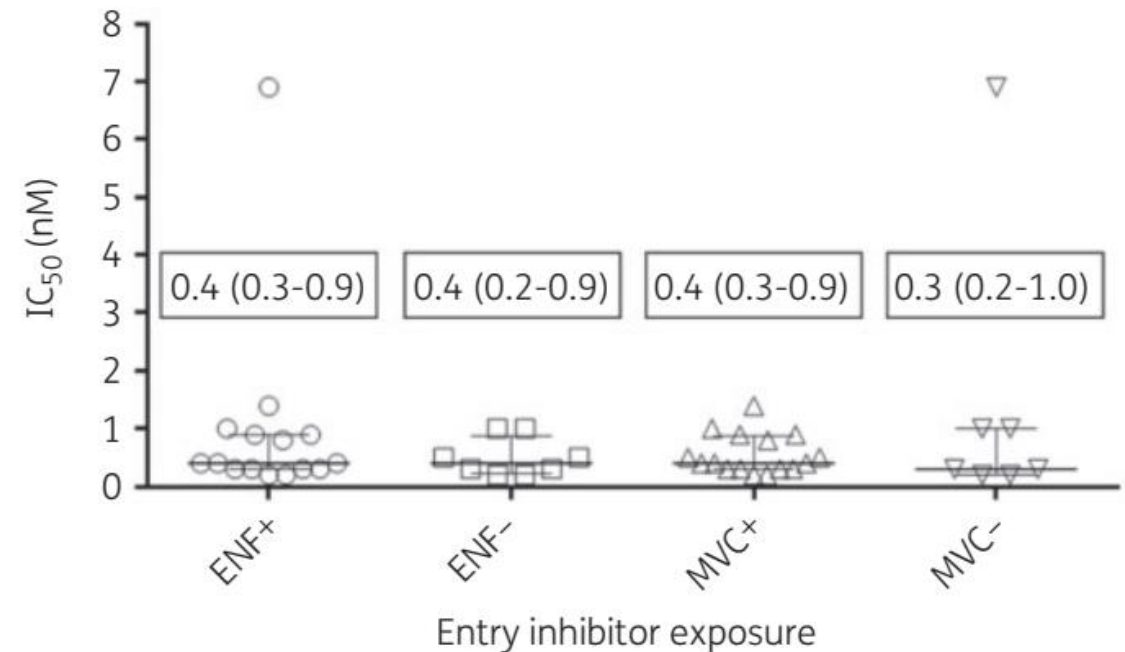
CRF01_AE is the least susceptible variant, however variable proportions of strains with reduced susceptibility may be present also within other subtypes or recombinants (mainly F-related)

Susceptibility to the novel attachment inhibitor Fostemsavir in patients harboring 4-class resistant HIV

Phenotypic analysis of 24 patients harboring MDR HIV from the PRESTIGIO registry



Susceptibility to Fostemsavir is comparable for viruses using different coreceptors



Susceptibility to Fostemsavir is not affected by past exposure to enfuvirtide or maraviroc

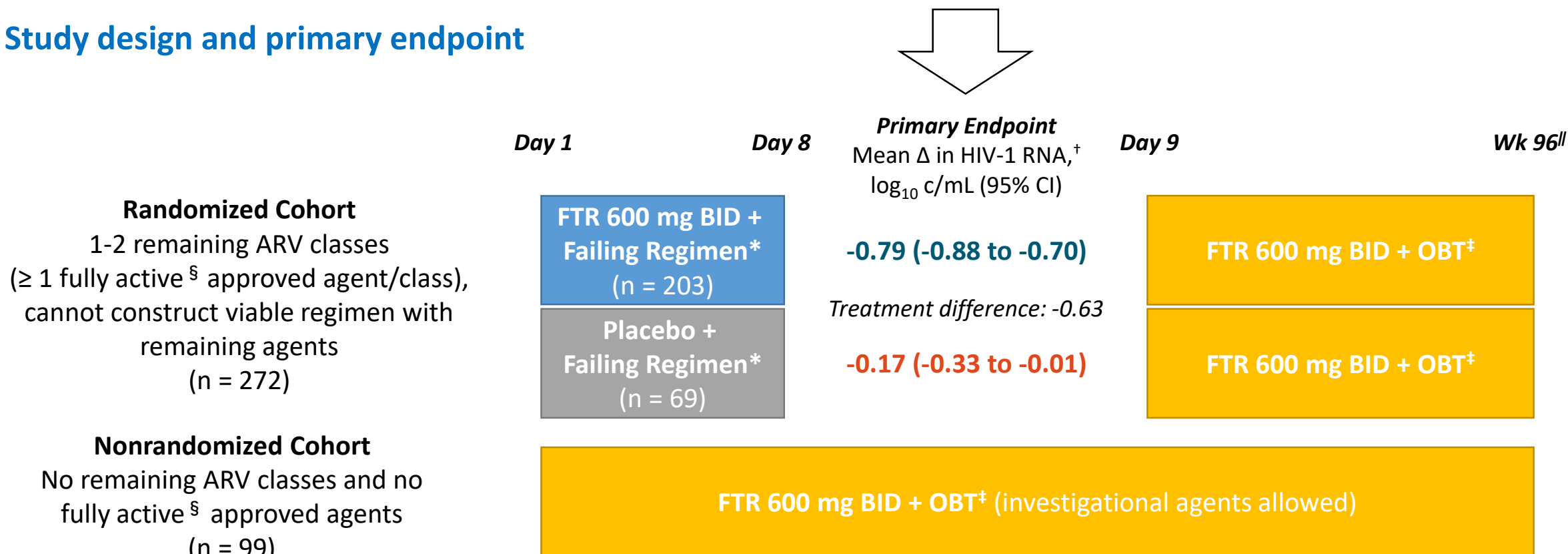
CCR5-Tropic MVC Resistance Was Not Predictive of Reduced Susceptibility to TMR in MOTIVATE-1 and -2

- No clinical samples from MOTIVATE were available, although gp120 sequences were accessible
- Sequences of envelopes from 5 participants who exhibited CCR5-tropic MVC resistance were used to regenerate functional envelopes
- All 5 exhibited resistance to MVC (and were CCR5-tropic), while 4/5 were fully susceptible to TMR
- The 1 envelope with reduced susceptibility to TMR contained an M426L polymorphism
- Site-directed mutagenesis reverting the virus back to M426 resulted in full sensitivity to TMR while MVC resistance remained

Envelope	TMR EC ₅₀ (nM)	MVC EC ₅₀ (nM)
MP5.7	1.00	>5000
MP35.2	1.00	>5000
MP49.20	1.56	>5000
MP53.36	0.51	>5000
MP11.38 (M426L)	412.7	>5000
MP11.38 (M426M; SDM)	0.98	>5000

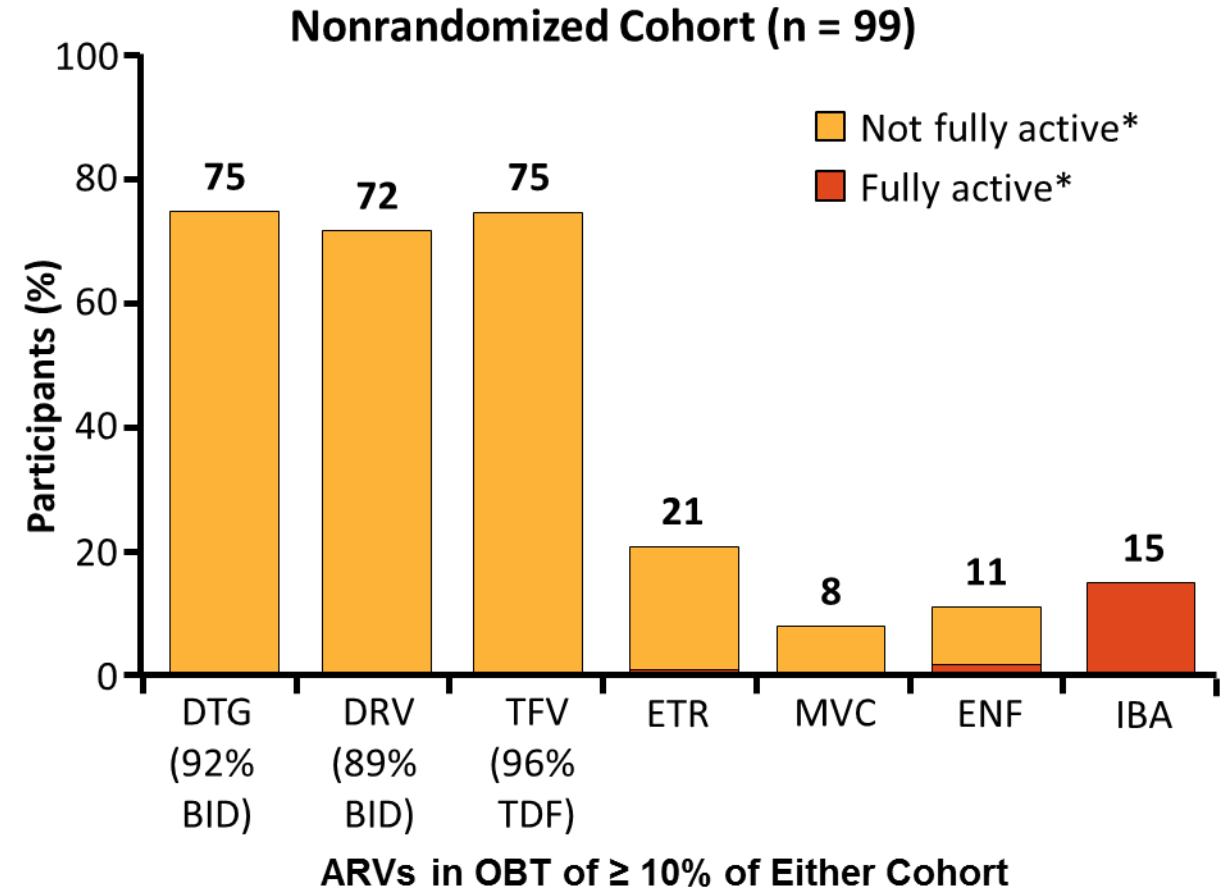
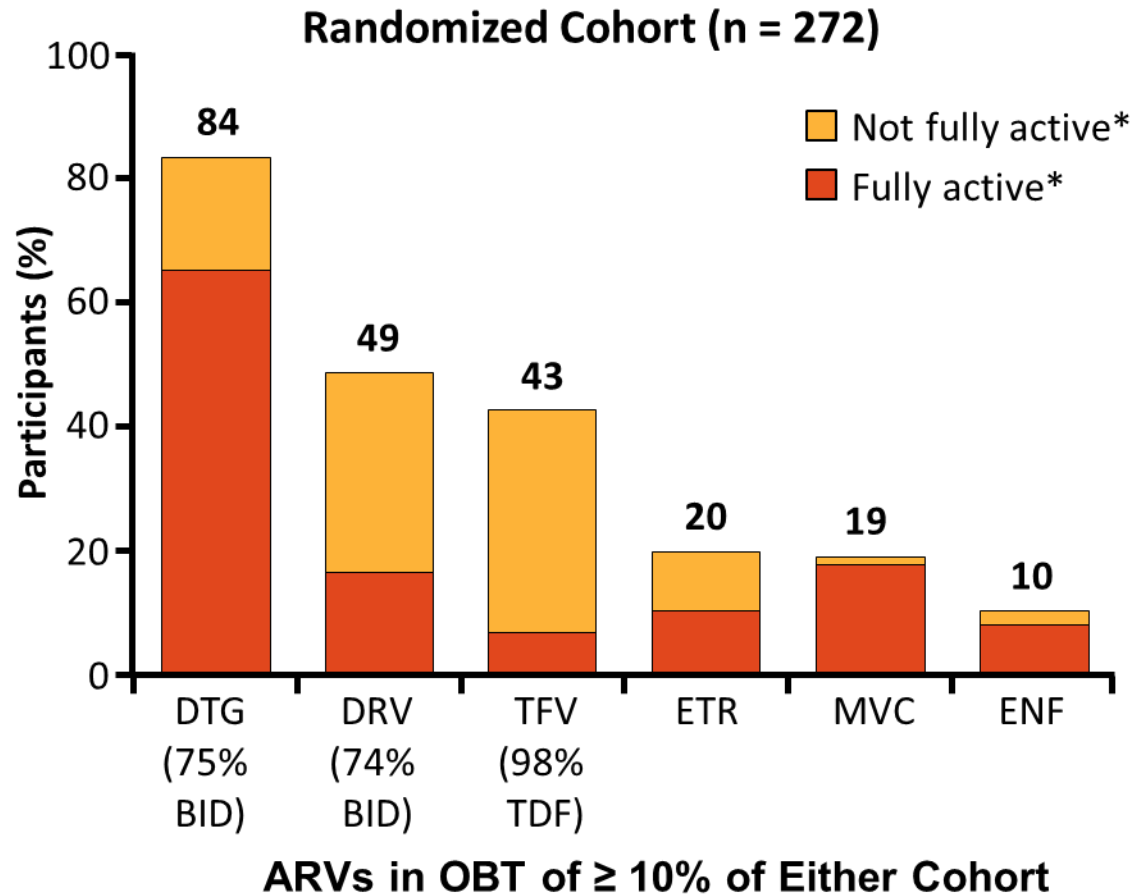
Fostemsavir in HTE individuals: the BRIGHTE study

Study design and primary endpoint



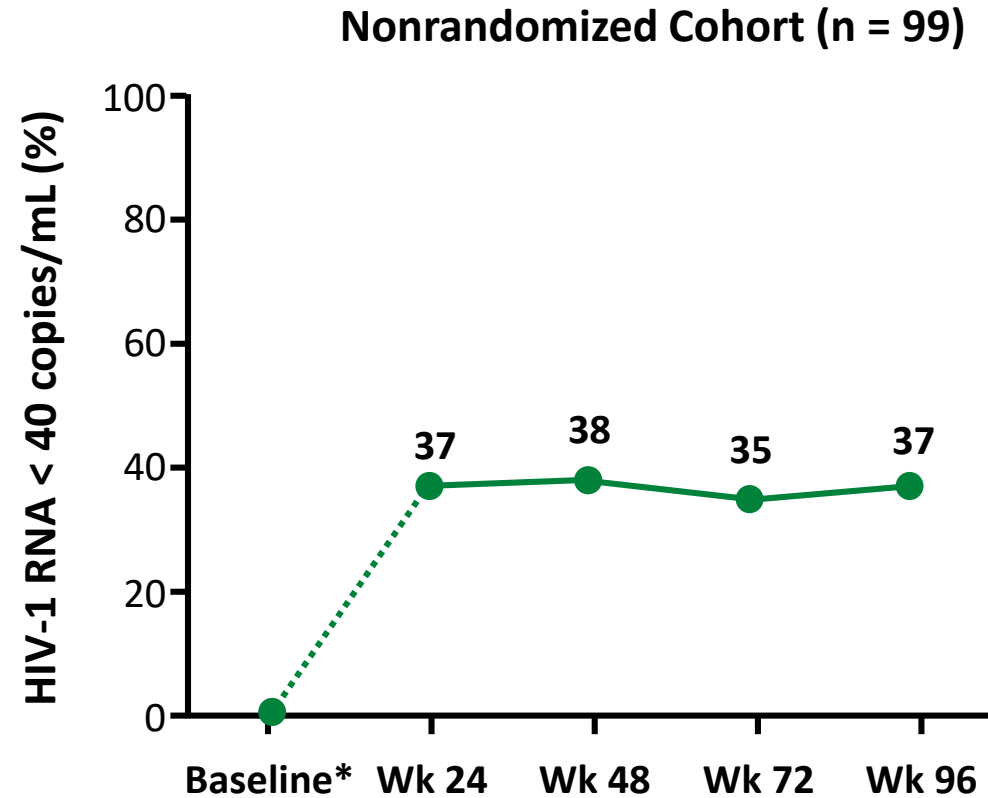
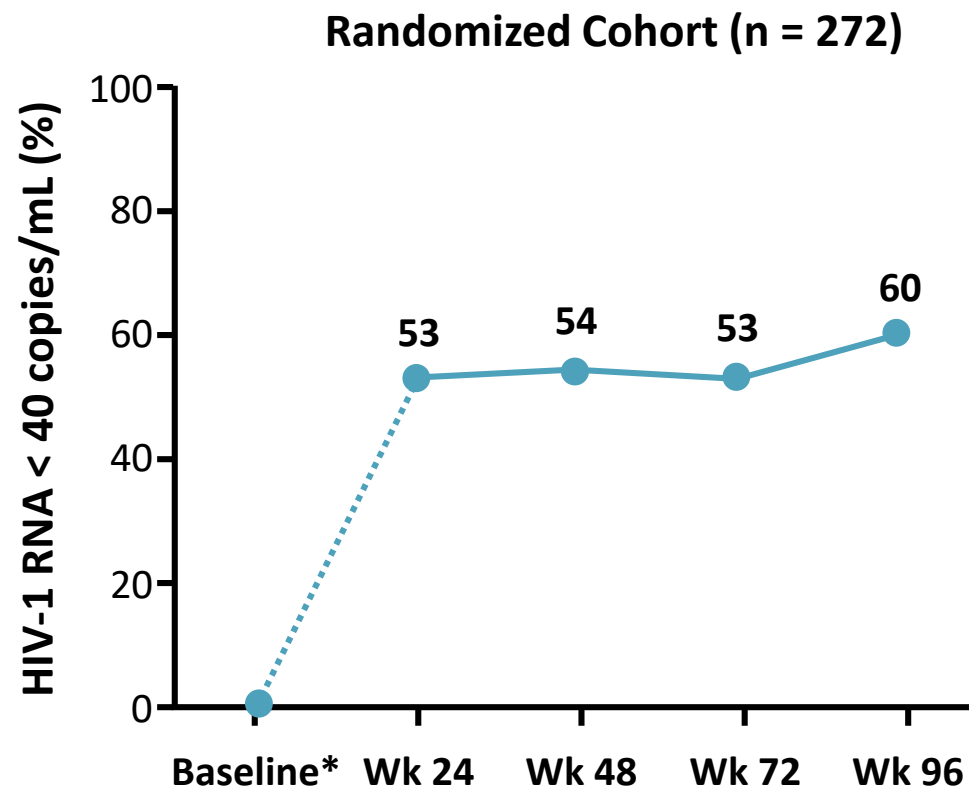
*Blinded. [†]Day 8 adjusted by Day 1. [‡]Open label. [§]No evidence of resistance; patient eligible for, tolerant of, willing to receive the ARV. ^{||}Measured from start of open-label treatment. Study conducted until another option, rollover study, or approved ARV available.

Fostemsavir in HTE individuals: the BRIGHT study



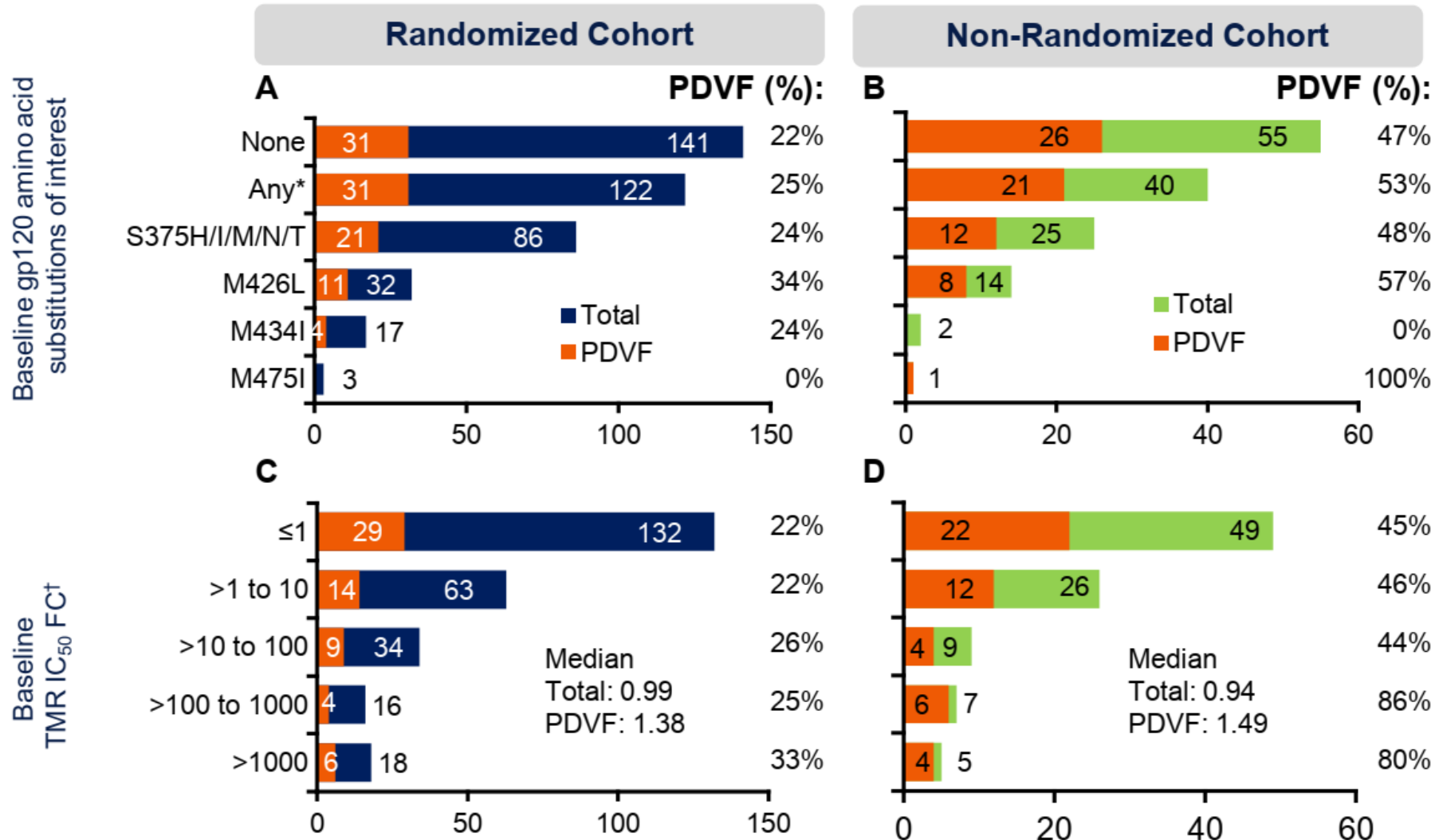
*Determined by commercial phenotype assays, historical resistance, eligibility, and tolerability.

Fostemsavir in HTE individuals: the BRIGHT study



*Snapshot analysis excluded baseline data. 1 patient had BL HIV-1 RNA < 40 copies/mL.

Fostemsavir in HTE individuals: the BRIGHTE study

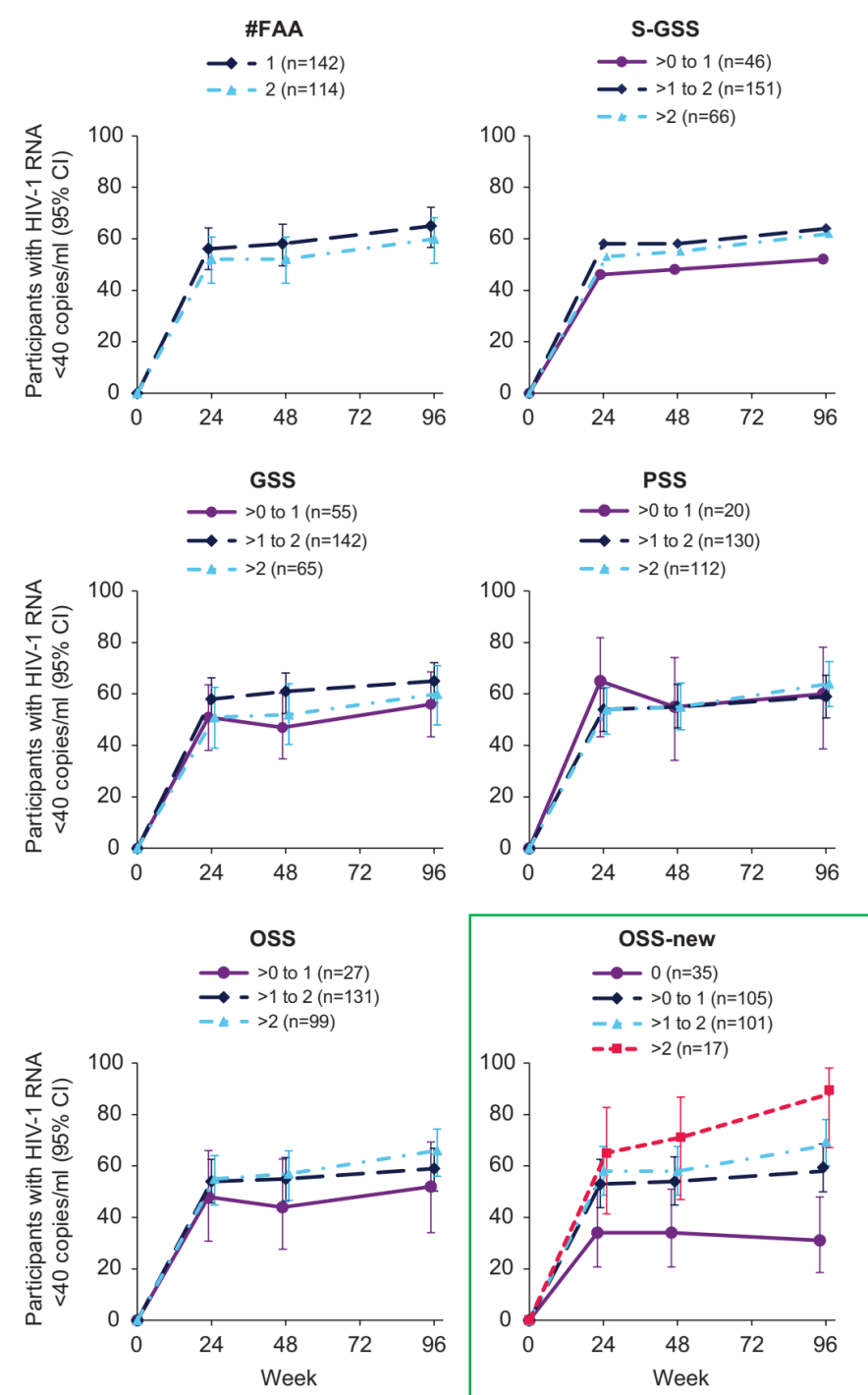


PDVF Through Week 96 for Randomized and Non-Randomized Cohorts by **Baseline gp120 Substitutions of Interest (A and B) and TMR IC₅₀ FC (C and D)**

- No correlation between baseline **genotypic** resistance and response
- Correlation between baseline **phenotypic** resistance (>100 FC IC₅₀) and response limited to the non-randomized cohort

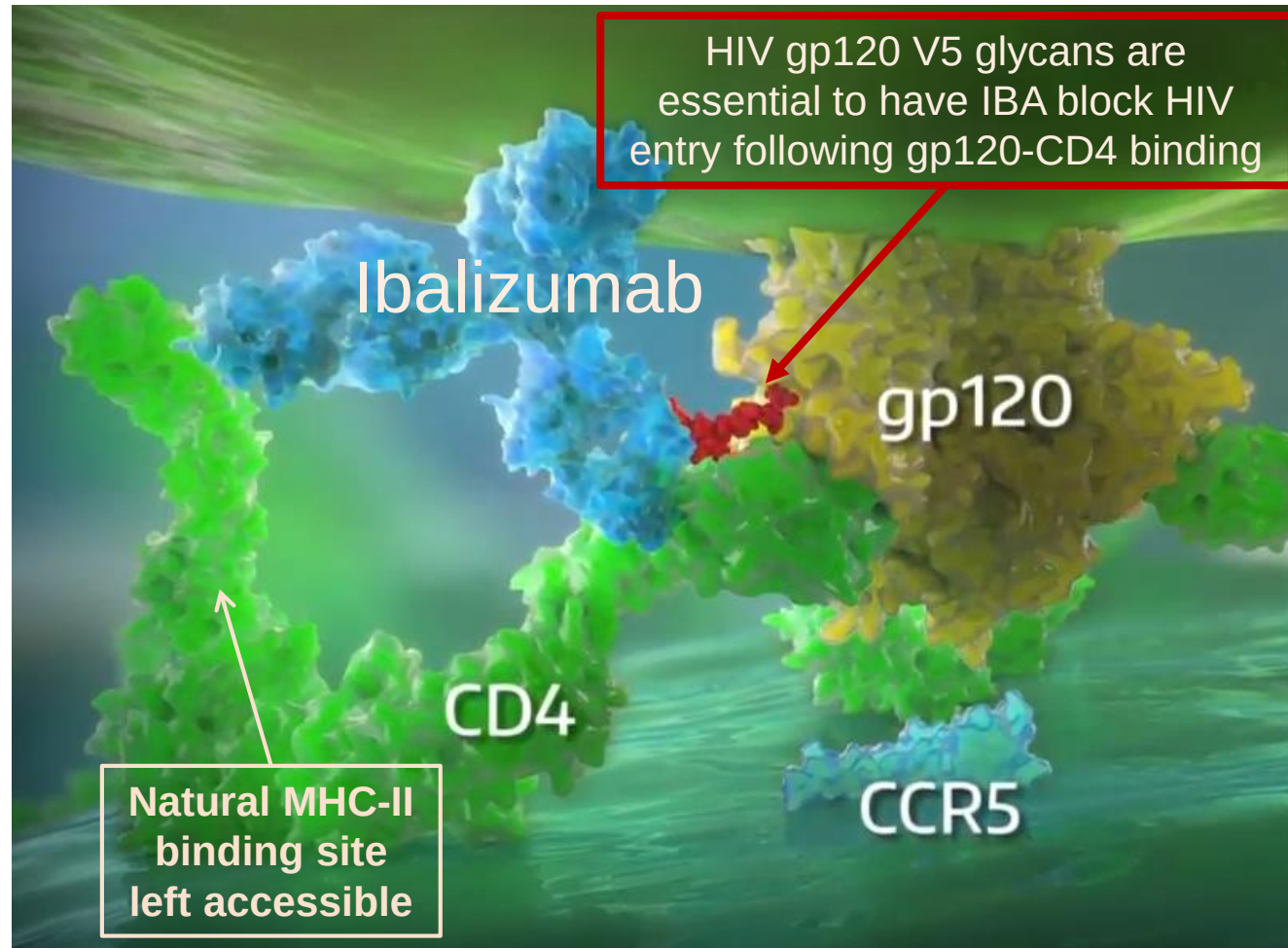
Fostemsavir in HTE individuals: the BRIGHT study

- VL <40 cp/mL by initial optimized background susceptibility scores
 - FAA**, fully active agents
 - S-GSS**, Stanford HIVdb genotypic susceptibility
 - GSS**, Monogram genotypic susceptibility
 - PSS**, Monogram phenotypic susceptibility
 - OSS**, overall susceptibility (combined GSS & PSS)
 - OSS-new**, OSS only for newly used drugs



New options for MDR HIV

Ibalizumab → targets CD4



New options for MDR HIV

Ibalizumab → targets CD4

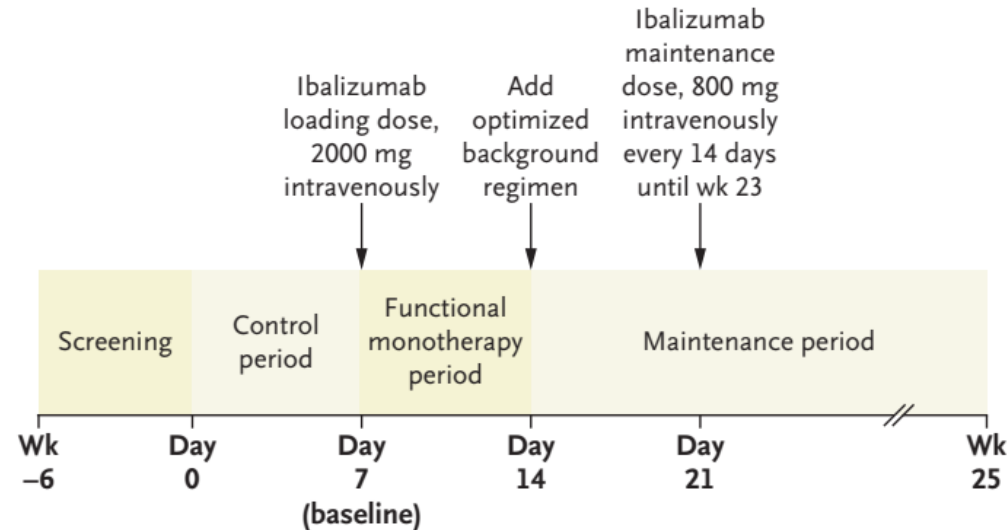
Known resistance to ≥1 drug in class — no. (%)

Nucleoside reverse-transcriptase inhibitor	37 (93)
Non-nucleoside reverse-transcriptase inhibitor	37 (93)
Protease inhibitor	36 (90)
Integrase inhibitor	27 (68)
Coreceptor antagonist†	33 (87)
Fusion inhibitor†	9 (24)

Known resistance to all drugs in class — no. (%)

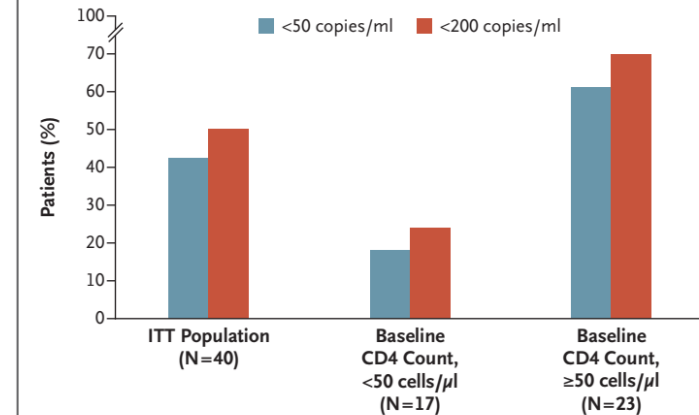
Nucleoside reverse-transcriptase inhibitor	26 (65)
Non-nucleoside reverse-transcriptase inhibitor	26 (65)
Protease inhibitor	25 (63)
Integrase inhibitor	19 (48)
Coreceptor antagonist†	33 (87)
Fusion inhibitor†	9 (24)

A Study Design

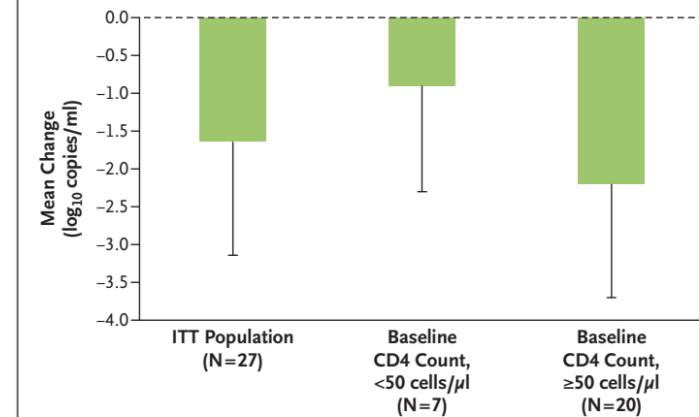


- TMB-301 study: 40 PLWH with extensively drug resistant HIV and VL >3 log (mean 4.5 ± 0.8 log)
- Primary end point: % patients with a decrease in VL ≥ 0.5 log from baseline (day 7) to day 14
- **On day 14, 33 (83%) had a decrease in VL ≥ 0.5 log₁₀ copies/ml from baseline**
 - Mean VL decrease: 1.1 log₁₀ copies/ml
- At week 25:
 - Mean decrease of 1.6 log₁₀ copies/ml from baseline
 - 43% VL <50 copies/ml; 50% VL <200 copies/ml
 - **10 patients had virologic failure or rebound**
 - **9 had a lower degree of susceptibility to ibalizumab than at baseline**

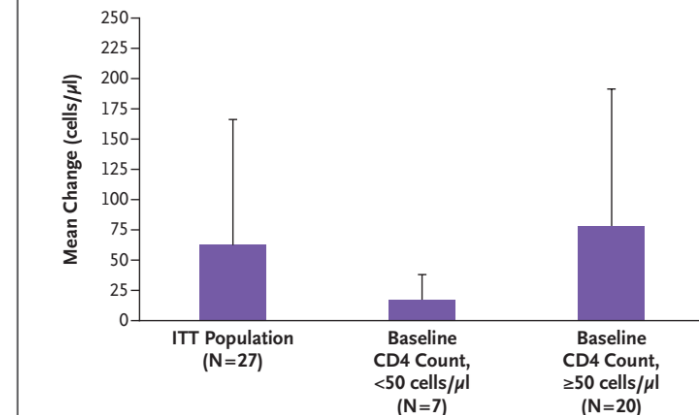
A HIV-1 Viral Load, According to CD4 Subgroup at Baseline



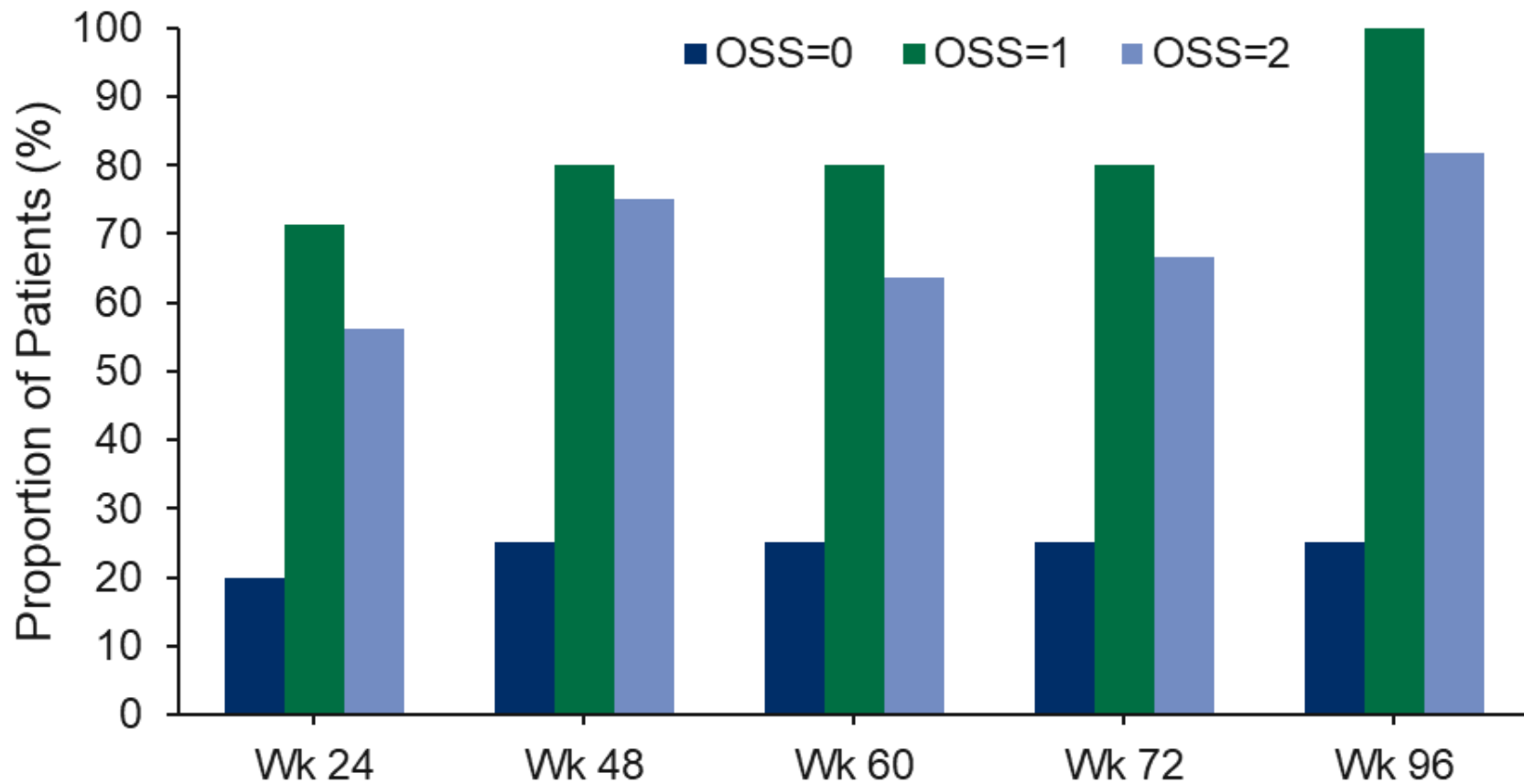
B Change from Baseline in Viral Load, According to CD4 Subgroup



C Change from Baseline in CD4 T-Cell Count, According to CD4 Subgroup



TMB-311: Proportion of patients with viral loads <50 cp/mL stratified by OSS (OT Population)



Overall susceptibility score (**OSS**) was calculated for each patient as the number of fully active ARVs (Stanford levels 1 and 2) in the OBR at baseline determined by viral genotype and phenotype (Monogram, San Francisco, CA) and historical results

Long-term Efficacy, Safety, and Durability of Ibalizumab-based Regimens in Subgroup of TMB-202 Participants

William Towner¹, MD, FACP, FIDSA, Edwin DeJesus², MD, FIDSA, Shannon Schrader³, MD, Jerome de Vente⁴, MD, Colleen McGary⁵, PhD, Mohammed Zogheib⁵, PharmD, MPH, Steven Weinheimer⁶, PhD, Pedro Mesquita⁷, PhD

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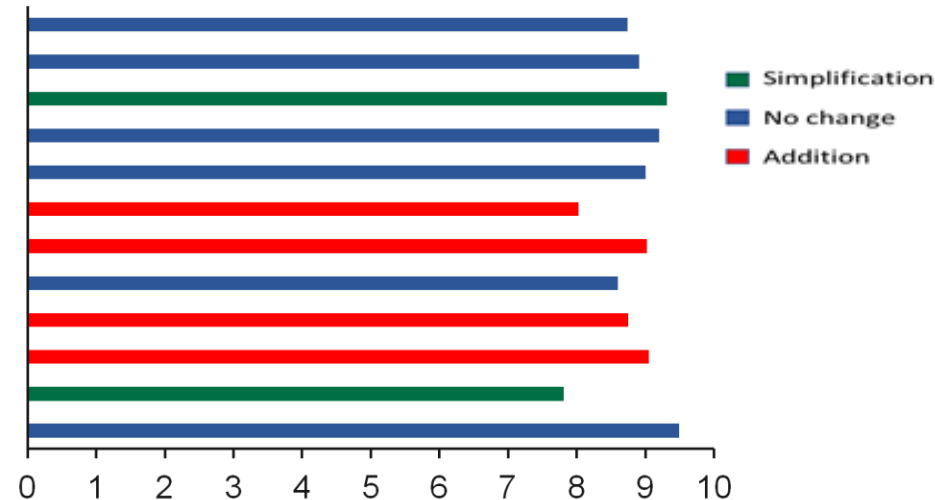


¹Kaiser Permanente Southern California; Pasadena, CA; ²Orlando Immunology Center, University of Central Florida College of Medicine; Orlando, FL; ³Research Access Network; Houston, Texas; ⁴Long Beach Education and Research Consultants; Long Beach, California; ⁵Syneos Health, Somerset, NJ; ⁶Taimed Biologics; Irvine, California; ⁷Theratechnologies Inc., Montreal, QC, Canada;

Results (cont'd)

Durability of Response

- ▶ Overall, the 12 patients remained on IBA for an average of 8.9 years (range 7.8-9.5)
- ▶ During this period, 8/12 did not require addition of new ARVs to their OBR to maintain suppression.



Clinical Characteristics and Outcomes of Coronavirus Disease 2019 Patients Who Received Compassionate-Use Leronlimab

Bryant Yang,¹ Jennifer A. Fulcher,^{1,2} Jenny Ahn,³ Marlene Berro,³ David Goodman-Meza,¹ Kush Dhody,⁴ Jonah B. Sacha,⁵ Arash Naeim,³ and Otto O. Yang^{1,6}

¹Division of Infectious Diseases, Department of Medicine, David Geffen School of Medicine at the University of California Los Angeles, Los Angeles, California, USA, ²Department of Medicine, VA Greater Los Angeles Healthcare System, Los Angeles, California, USA, ³Clinical and Translational Science Institute Office of Clinical Research, Department of Medicine, David Geffen School of Medicine at the University of California Los Angeles, Los Angeles, California, USA, ⁴Amarex Clinical Research, LLC, Germantown, Maryland, USA, ⁵Vaccine and Gene Therapy Institute and Oregon National Primate Research Center, Oregon Health & Science University, Beaverton, Oregon, USA, and ⁶Department of Microbiology, Immunology, and Molecular Genetics, David Geffen School of Medicine at the University of California Los Angeles, Los Angeles California, USA

Background. Leronlimab, a monoclonal antibody blocker of C-C chemokine receptor type 5 originally developed to treat human immunodeficiency virus infection, was administered as an open-label compassionate-use therapeutic for coronavirus disease 2019 (COVID-19).

Methods. Twenty-three hospitalized severe/critical COVID-19 patients received 700 mg leronlimab subcutaneously, repeated after 7 days in 17 of 23 patients still hospitalized. Eighteen of 23 received other experimental treatments, including convalescent plasma, hydroxychloroquine, steroids, and/or tocilizumab. Five of 23 received leronlimab after blinded, placebo-controlled trials of remdesivir, sarilumab, selinexor, or tocilizumab. Outcomes and results were extracted from medical records.

Results. Mean age was 69.5 ± 14.9 years; 20 had significant comorbidities. At baseline, 22 were receiving supplemental oxygen (3 high flow, 7 mechanical ventilation). Blood showed markedly elevated inflammatory markers (ferritin, D-dimer, C-reactive protein) and an elevated neutrophil-to-lymphocyte ratio. By day 30 after initial dosing, 17 were recovered, 2 were still hospitalized, and 4 had died. Of the 7 intubated at baseline, 4 were fully recovered off oxygen, 2 were still hospitalized, and 1 had died.

Conclusions. Leronlimab appeared safe and well tolerated. The high recovery rate suggested benefit, and those with lower inflammatory markers had better outcomes. Some, but not all, patients appeared to have dramatic clinical responses, indicating that unknown factors may determine responsiveness to leronlimab. Routine inflammatory and cell prognostic markers did not markedly change immediately after treatment, although interleukin-6 tended to fall. In some persons, C-reactive protein clearly dropped only after the second leronlimab dose, suggesting that a higher loading dose might be more effective. Future controlled trials will be informative.

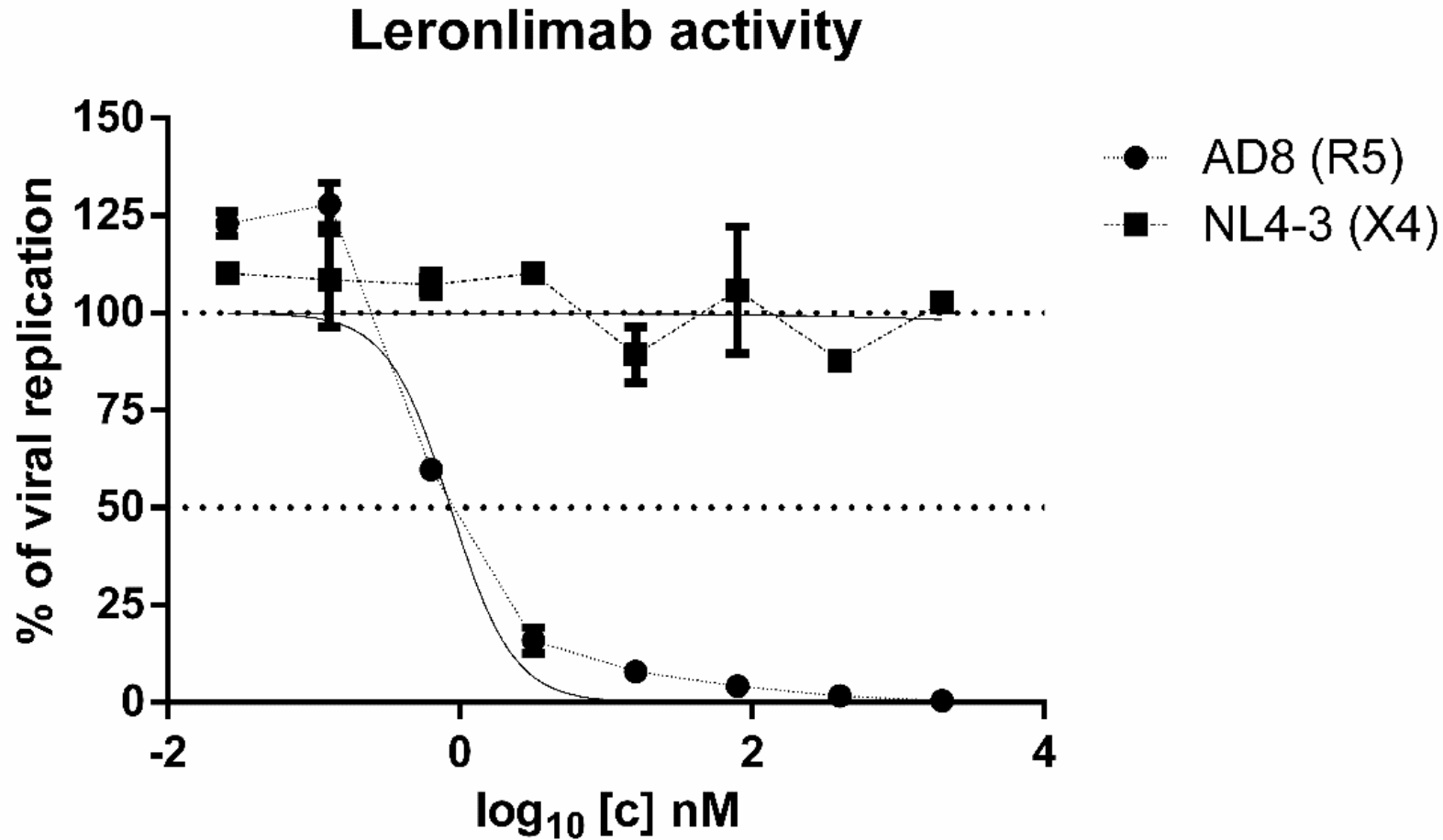
Leronlimab (PRO 140) *in vitro* activity against 4-class drug resistant HIV-1 from heavily treatment experienced subjects

Variable	Patient population (n=24)
Age (years), median (IQR)	48.0 (39.1 – 52.7)
Male gender, N (%)	19 (79%)
Calendar year of first HIV positive test, median (IQR)	1992 (1988 – 1993)
Years since first HIV positive test, median (IQR)	22.3 (19.3 – 25.2)
Calendar year of ART start, median (IQR)	1993 (1992 - 1997)
Years since ART start, median (IQR)	19.7 (15.6 – 22.9)
Months to ART start since first HIV positive test, median (IQR)	11.1 (1.2 – 48.7)
Previous diagnosis of AIDS, N (%)	14 (58%)
Nadir CD4 ⁺ T-cells (N/ μ L), median (IQR)	52 (9 - 184)
Zenith HIV-1 RNA ($\log_{10}$ copies/mL), median (IQR)	5.06 (4.36 – 5.60)
Duration (months) with detectable plasma HIV-1 RNA (>50 copies/mL) at the time of current analysis, median (IQR)	20.1 (4.4 – 55.8)
HIV-1 RNA ($\log_{10}$ copies/mL), median (IQR)	4.69 (4.11 – 5.20)
CD4 ⁺ T-cells (N/ μ L), median (IQR)	192 (65.5 - 252)
CD8 ⁺ T-cells (N/ μ L), median (IQR)	950 (656 - 1728)
CD4 ⁺ /CD8 ⁺ ratio, median (IQR)	0.14 (0.09 – 0.26)

S. Rusconi, Pharmacological Research 2022



Leronlimab (PRO 140) *in vitro* activity against 4-class drug resistant HIV-1 from heavily treatment experienced subjects



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Leronlimab (PRO 140) *in vitro* activity against 4-class drug resistant HIV-1 from heavily treatment experienced subjects

Table 1. Baseline HIV-1 RNA, CD4⁺ T cell counts Genotypic coreceptor tropism results obtained by Sanger and NGS and leronlimab *in vitro* susceptibility for the 9 phenotypically R5 pseudoviruses.

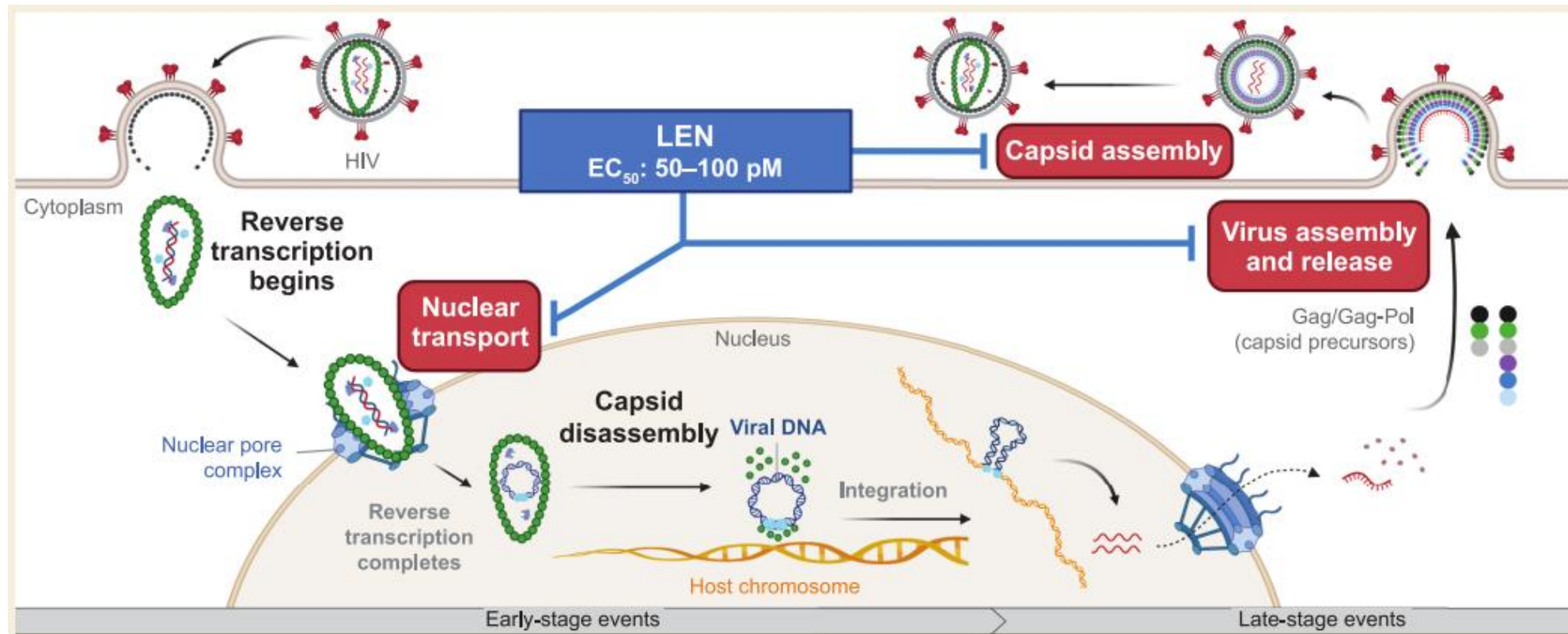
Patient ID	HIV-1 RNA (copies/mL)	CD4 ⁺ T-cells/ μ L	Percentage of non-R5 variants by NGS (FPR $\leq$ 3.5%)	FPR by Sanger (%)	IC ₅₀ (mean $\pm$ SD) PRO 140	Previous maraviroc exposure	Exposure to maraviroc at current analysis
25	1,104	240	32.7	4.4	0.3 $\pm$ 0.2	Yes	No
27	13,835	168	0.1	not done	1.2 $\pm$ 0.3	No	No
37	12,580	207	0.1	69.8	0.4 $\pm$ 0.3	Yes	Yes
55	17,888	326	0.0	54.7	0.7 $\pm$ 0.2	No	No
58	2,986	518	0.0	51.3	0.5 $\pm$ 0.3	No	No
82	373,713	942	83.8	54.7	0.4 $\pm$ 0.3	No	No
90	84,534	201	0.6	16.4	0.8 $\pm$ 0.6	No	No
117	6,718,473	34	0.1	51.2	0.3 $\pm$ 0.1	Yes	Yes

NGS, Next Generation Sequencing. FPR, False Positive Rate. IC₅₀, Half-maximal inhibitory concentration. SD, standard deviation.



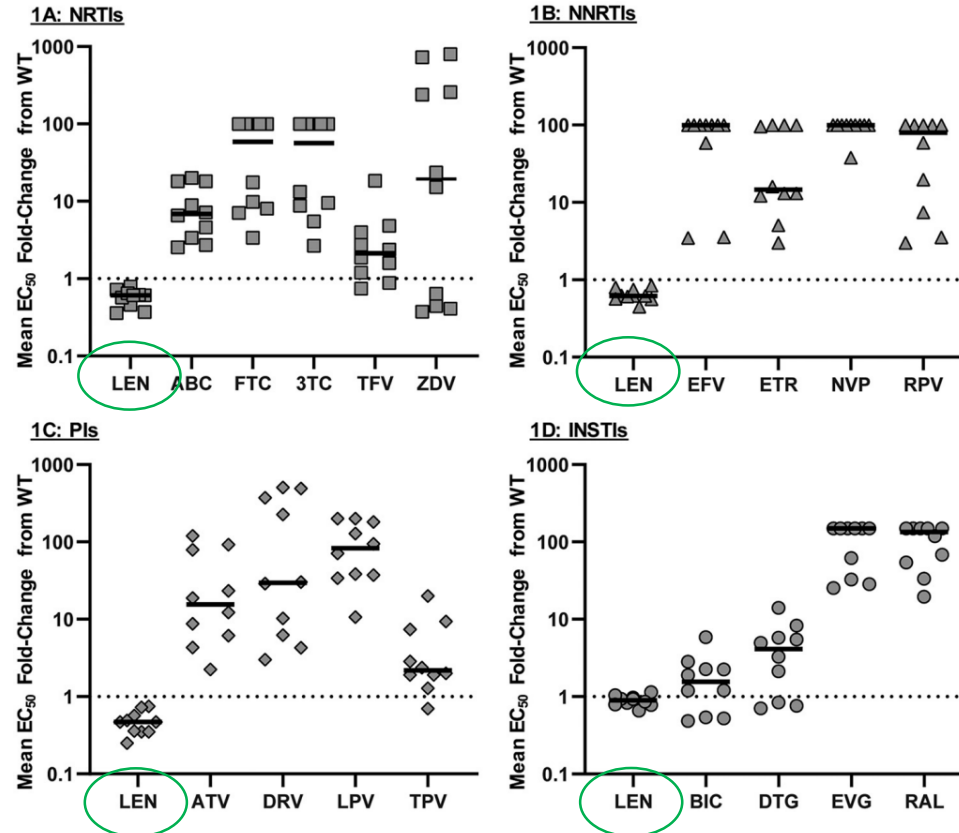
New options for MDR HIV

Lenacapavir → targets HIV capsid



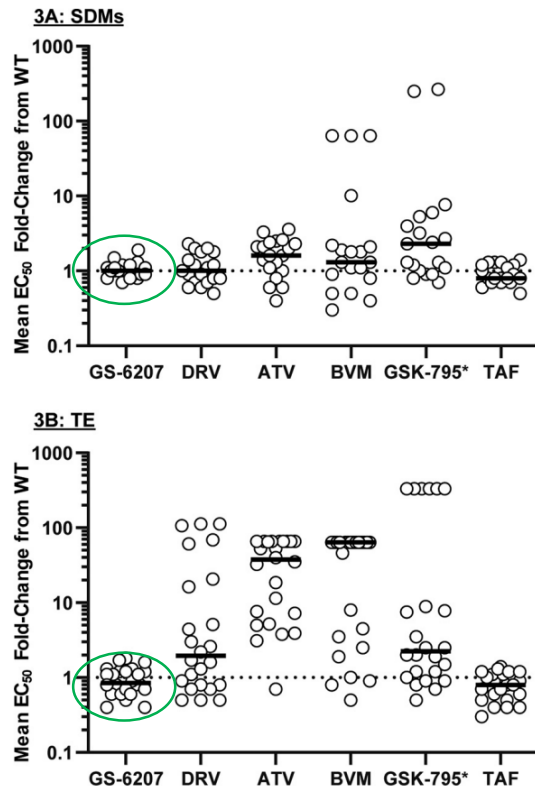
- First-in-class capsid (CA) inhibitor
- Picomolar potency (EC₅₀ = 50-100 pM)
- LEN inhibits CA-mediated nuclear entry of HIV DNA, HIV assembly and proper capsid formation

Lenacapavir



- Activity of LEN in mutants resistant to other drug classes
- HIV mutations were inserted into the pXXLAI clone, and the resulting mutants ($n = 40$) were evaluated using a 5-day antiviral assay

Lenacapavir



*The maturation inhibitor GSK3532795 was discontinued

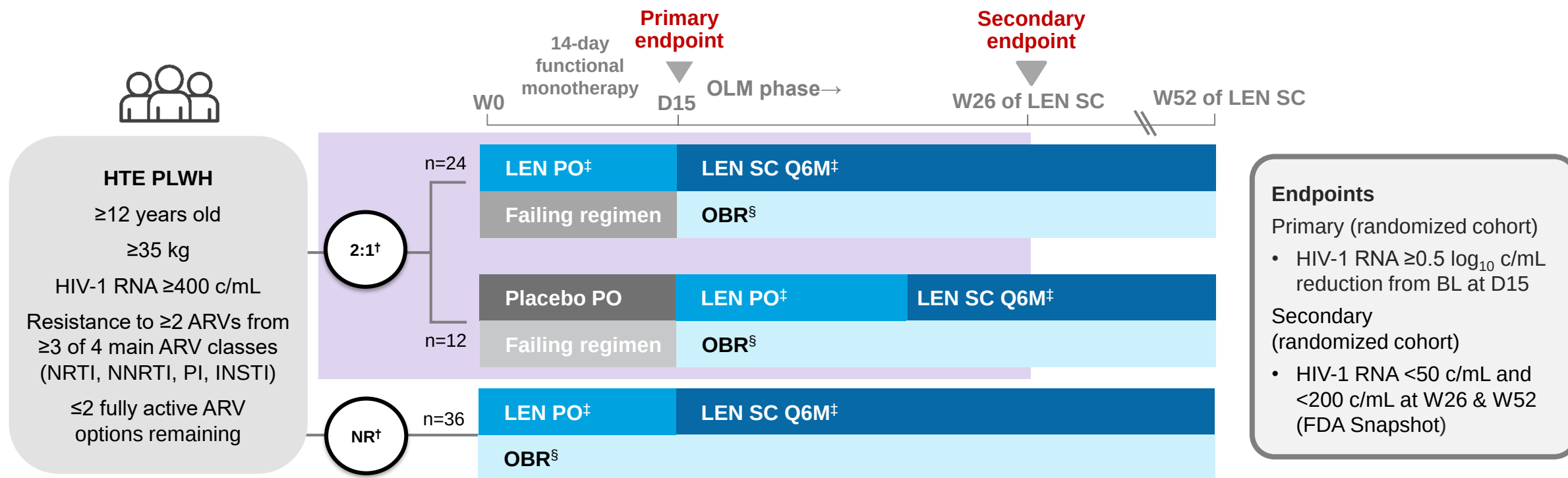
Site-directed
mutants

Clinical
isolates from
treatment
experienced
subjects

- Activity of LEN in mutants with **Gag cleavage site mutations**
- HIV mutations were inserted into the pXXLAI clone, and the resulting mutants ($n = 43$) were evaluated using a 5-day antiviral assay

LEN in Heavily Treatment-Experienced PLWH

Phase 2/3, blinded, placebo-controlled study to evaluate LEN as an add-on to a failing regimen in heavily treatment-experienced PLWH with MDR (N=72)



†Participants with <0.5 log₁₀ decline in HIV-1 RNA during screening entered the randomized cohort; participants with ≥0.5 log₁₀ decline in HIV-1 RNA during screening entered the non-randomized cohort

§Investigational agents, such as fostemsavir, were allowed; atazanavir (ATV), ATV/cobicistat, ATV/ritonavir, efavirenz, entecavir, nevirapine, tipranavir were not allowed

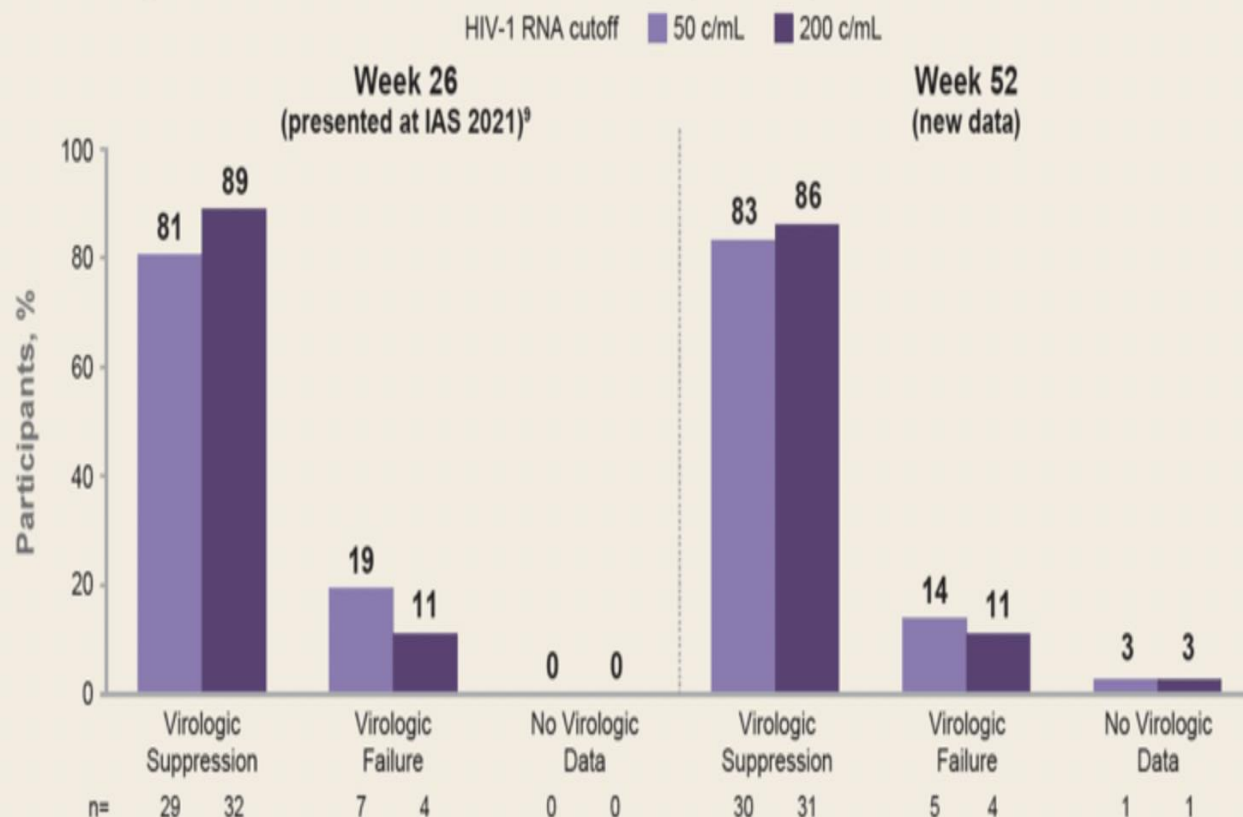
‡BL, baseline; D, day; HTE, heavily treatment-experienced; MDR, multidrug resistance; NR, nonrandomized; OBR, optimized background regimen; OLM, open-label maintenance; PO, by mouth; Q6M, every 6 months; SC, subcutaneous

Long-Acting Lenacapavir in People With Multidrug-Resistant HIV-1: Week 52 Results

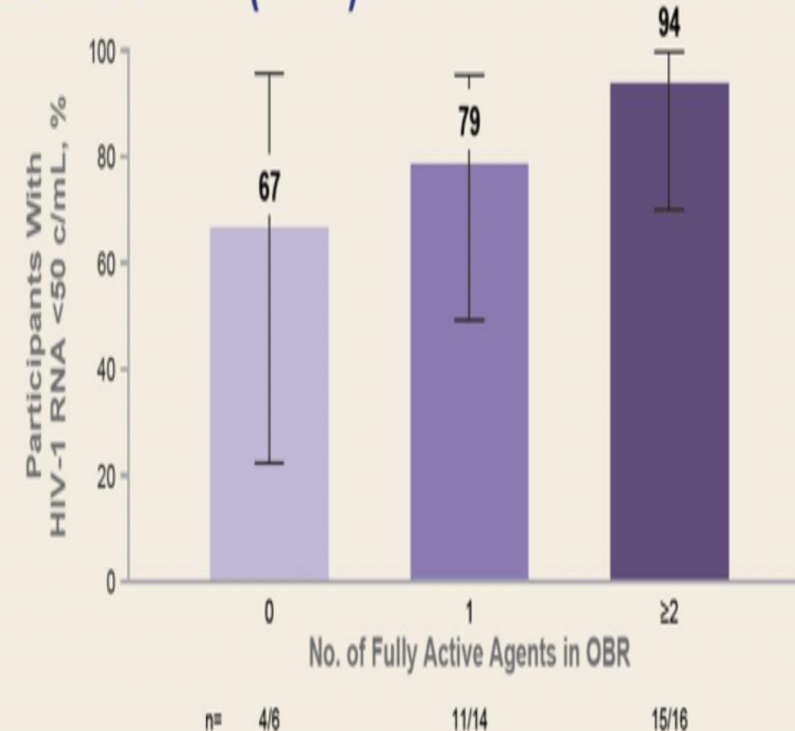
Oryema Ogburn,¹ Sonasa Segal-Maurer,² Cynthia Brinson,³ Ploechan Chetchotsak,⁴ Kenneth Lichtenstein,⁵ Joseph McGowan,⁶ Andrew A. Wozniak,⁷ Kimberly Workowski,⁸ Hui Wang,⁹ Nicolas Margot,¹ Hades Dwory-Sobel,¹ Martin S. Rhee,¹⁰ Jared M. Barton,¹ Jean-Michel Molina¹¹

¹Gilead University, Foster City, CA; ²New York Presbyterian Queens, NY; ³Central Texas Clinical Research, Austin, TX; ⁴Viragard Hospital, Khor Kaen, Thailand; ⁵Epsteinbaum Health Center, Palm Springs, CA; ⁶Belmont Health, Belmont, MA; ⁷Justi Medical Center, Bronx, NY; ⁸Emory University, Atlanta, GA; ⁹Gilead Sciences, Inc., Foster City, CA; ¹⁰Hopital Saint Louis, Paris, France

Efficacy in Randomized Cohort (n=36)



Efficacy by No. of Fully Active Agents in OBR at Week 52 in Randomized Cohort (n=36)



Long-Acting Lenacapavir in People With Multidrug-Resistant HIV-1: Week 52 Results

Oryema Ogburn,¹ Soana Segal-Maurer,² Cynthia Brinson,³ Ploechan Chetchotsak,⁴ Kenneth Lichtenstein,⁵ Joseph McGowan,⁶ Andrew A. Wozniak,⁷ Kimberly Workowski,⁸ Hui Wang,⁹ Nicolas Margot,¹ Hades Dwory-Sobel,¹ Martin S. Rhee,¹ Jared M. Barton,¹ Jean-Michel Molina¹⁰

¹Gilead University, Foster City, CA; ²New York Presbyterian Queens, NY; ³Central Texas Clinical Research, Austin, TX; ⁴Viragard Hospital, Nong Khai, Thailand; ⁵Eleusis Health Center, Palm Springs, CA; ⁶Belmont Health, Belmont, MA; ⁷Justi Medical Center, Brent, MI; ⁸Emory University, Atlanta, GA; ⁹Gilead Sciences, Inc., Foster City, CA; ¹⁰Hôpital Saint Louis, Paris, France

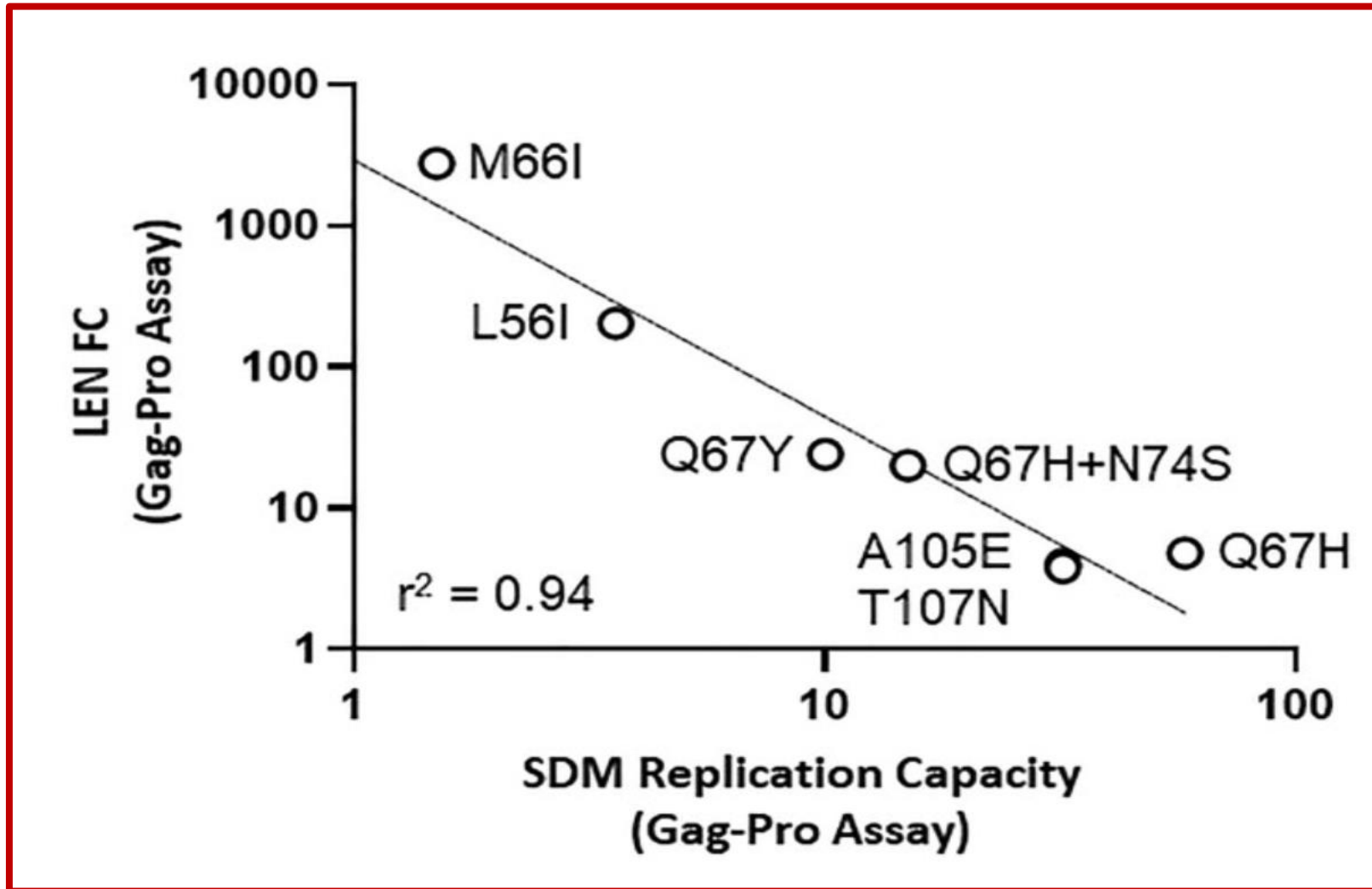
Emergent LEN Resistance*

n (%)	Randomized Cohort: n=36 (presented at IAS 2021, EACS 2021) ^{9,10}	Nonrandomized Cohort: n=36
Participants meeting criteria for resistance testing	11 (31)	10 (28)
Emergent LEN resistance [†]	4 (11)	4 (11)
M66I	4	2
Q67H/K/N	1	2
K70H/N/R/S	1	3
N74D/H/S	3	0
A105S/T	3	1
T107A/C/N	1	3

*Capsid genotypic and phenotypic resistance testing performed on any participants with confirmed HIV-1 RNA ≥ 50 c/mL and $< 1 \log_{10}$ HIV-1 RNA reduction from Day 1 at Week 4 visit, at any visit after achieving HIV-1 RNA < 50 c/mL and rebound to ≥ 50 c/mL, and at any visit with $> 1 \log_{10}$ increase from nadir; HIV-1, protease, reverse-transcriptase, and integrase genotypic and phenotypic testing were performed if rebound or suboptimal virologic response were confirmed;

[†]Developed during maintenance period (Week 4 [n=5], Week 10 [n=2], and Week 26 [n=1]).

Correlation between replication capacity and fold change in lenacapavir susceptibility in the Gag-Pro assay.



Long-Acting Lenacapavir in People With Multidrug-Resistant HIV-1: Week 52 Results

Oryema Ogbuagu,¹ Sorana Segal-Maurer,² Cynthia Brinson,² Ploechan Chetchotaisak,³ Kenneth Lichtenstein,⁴ Joseph McGowan,⁵ Andrew A. Wozniak,⁶ Kimberly Workowski,⁷ Hui Wang,⁸ Nicolas Margot,⁹ Hadas Dvory-Sobel,¹⁰ Martin S. Rhee,¹¹ Jared M. Barton,¹² Jean-Michel Molina¹³

¹Gilead University, New Haven, CT; ²New York Presbyterian Queens, NY; ³Central Texas Clinical Research, Austin, TX; ⁴Sirirajaporn Hospital, Nong Khai, Thailand; ⁵Eastman Health Center, Palm Springs, CA; ⁶Northwell Health, Manhasset, NY; ⁷Justi Medical Center, Bronx, NY; ⁸Emory University Atlanta, GA; ⁹Gilead Sciences, Inc., Foster City, CA; ¹⁰Hôpital Saint Louis, Paris, France

Adverse Events (excluding ISRs)*

≥10% Total in Any Grade, % (n)	Total LEN: N=72
Diarrhea	13 (9)
Nausea	13 (9)
COVID-19	11 (8)

*Serious adverse events (AEs) not related to study drug: malignant neoplasm and dizziness (n=1); abdominal pain, pancreatic mass, *Clostridium difficile* colitis, and angina pectoris (n=1); anal squamous cell carcinoma, proctalgia, impaired healing, and anal cancer (n=1); femoral neck fracture (n=1); COVID-19 (n=2); pneumonia (n=1); and septic shock, renal impairment, and shock (n=1). ISRs, injection-site reactions.

Incidence of ISRs Related to SC LEN*

ISR Types, %	After 1st SC Dose at Week 1 N=72	After 2nd SC Dose at Week 26 n=70	Median Duration, d
Swelling	26	13	12
Erythema	24	11	6
Pain	22	21	3
Nodule	22	11	180
Induration	11	10	118

*Only includes AEs related to LEN and excludes those not related to it.

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Grade 3 or 4 Laboratory Abnormalities

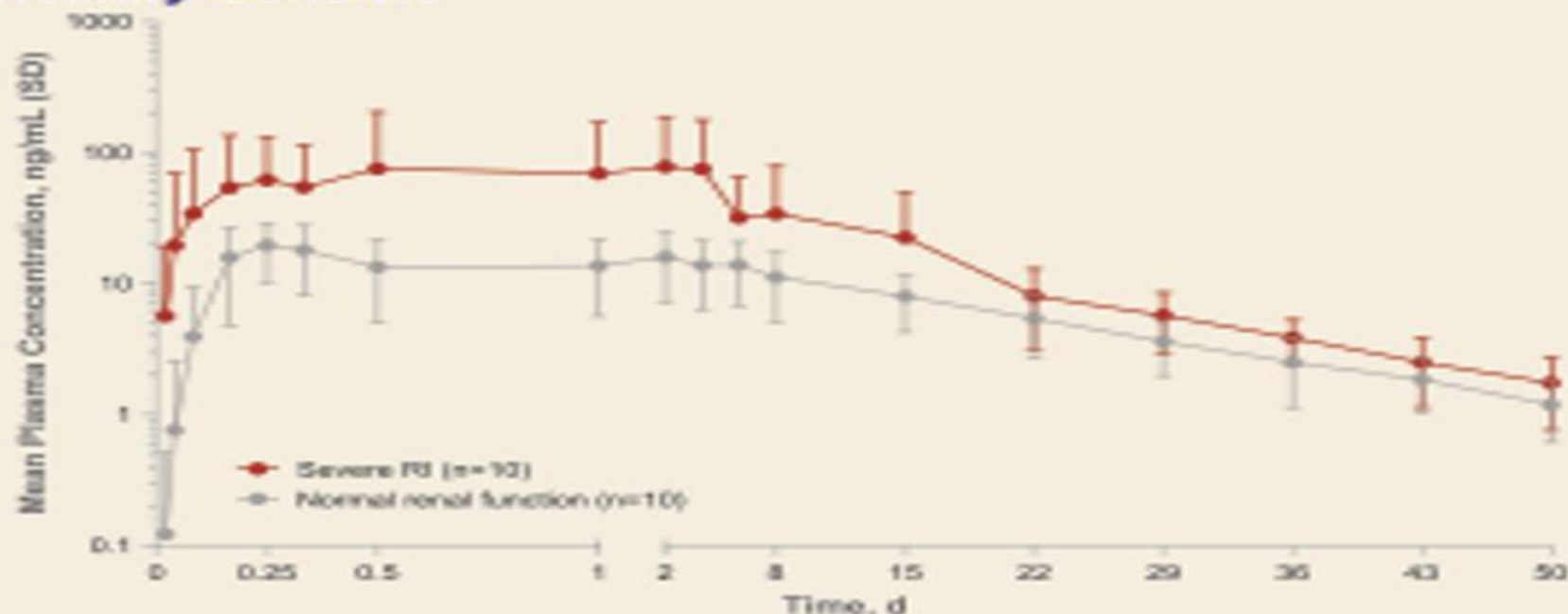
Laboratory Abnormality, % (n)	Total: N=72
Any Grade 3 or 4	29 (21)
≥5% in total	
Low creatinine clearance (eGFR)*	14 (10)
Elevated creatinine†	13 (9)
Glycosuria	6 (4)
Nonfasting/fasting hyperglycemia	6 (3)

*Per Division of AIDS scale, Grade 3 creatinine clearance is <60–30 mL/min or 30–<50% decrease from baseline; †Grade 3 creatinine is >1.8–<3.5 x upper limit of normal or increase to 1.5–<2.0 x baseline. eGFR, estimated glomerular filtration rate.

Pharmacokinetics of Lenacapavir in Participants With Severe Renal Impairment

Elijah J. Weber, Hiba Graham, Steve K. West, John Ling, Martin Rhee, Ramesh Palaparthi — Gilead Sciences, Inc., Foster City, CA

LEN PK in Participants With Severe RI and Matched Healthy Controls



PK Parameter*	Severe RI n=10	Normal Renal Function n=10	%CLSMR [90% CI]
AUC _{0-∞} , h·ng/mL	19,000 (28)	7400 (48)	164 (93.6, 360)
AUC ₀₋₂₄ , h·ng/mL	18,300 (101)	6640 (48)	169 (96.2, 377)
C _{max} , ng/mL	118 (129)	22.1 (44)	262 (112, 654)
T _{max} , h	8.00 (6.00, 24.0) [0.33 d]	8.00 (6.00, 8.00) [0.25 d]	—
t _{1/2} , h	234 (178, 354) [0.75 d]	318 (270, 367) [13.3 d]	—

*AUC from time 0 to ∞ = (AUC₀₋₂₄ + AUC from time 24 to last measured concentration) (AUC₀₋₂₄ and C_{max} presented as mean (% coefficient of variation), and time to C_{max} (T_{max}) and half-life (t_{1/2}) presented as median (quartiles 1, 3).

A New Dual Option with LEN

Capsid inhibitor
LEN



Islatravir

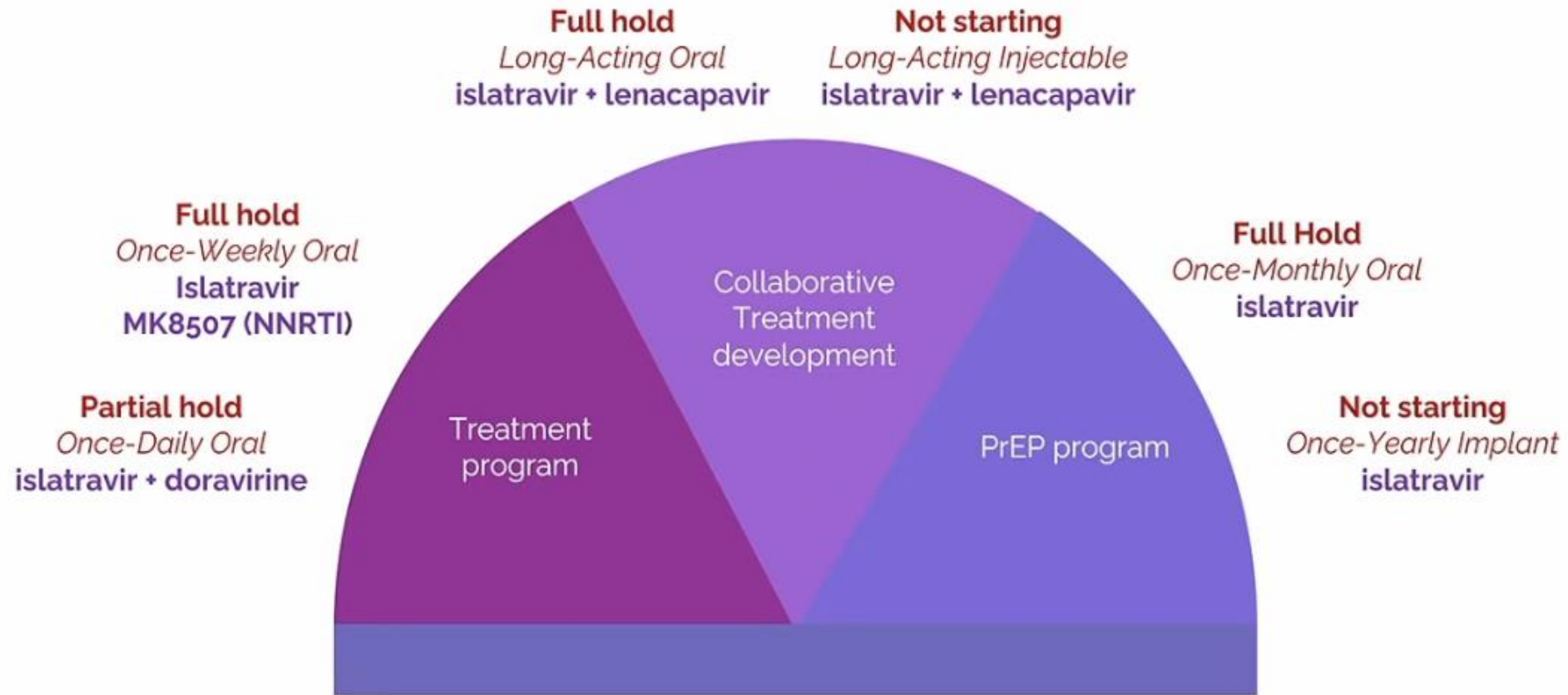
Oral capsid inhibitor
LEN



Islatravir oral

LEN and ISL are not approved for use by any regulatory body globally

Islatravir development : Current Status



A RANDOMIZED TRIAL OF DTG PLUS DRV/c AS A SWITCH STRATEGY IN SUBJECTS WITH MDR HIV-1 Poster-I02

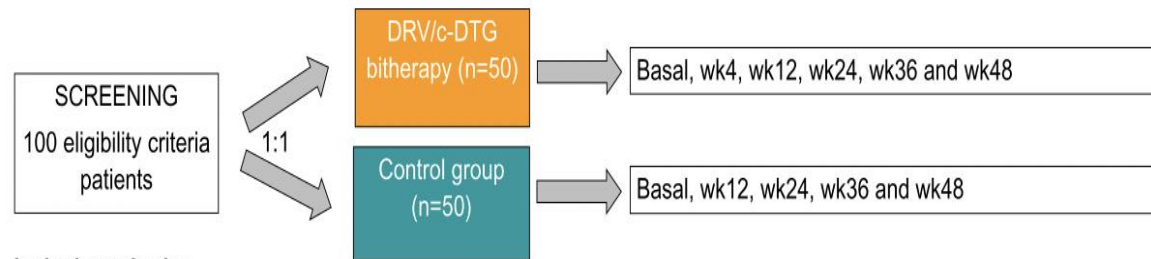
José R. Santos,¹ Pere Domingo,² Joaquín Portilla,³ Félix Gutiérrez,⁴ Arkaitz Imaz,⁵ Helem Vilchez,⁶ Adrià Curran,⁷ Nieves Valcarce-Pardeiro,⁸ Antoni Payeras,⁹ Enrique Bernal,¹⁰ Marta Montero-Alonso,¹¹ Miguel Yzuskí,¹² and R. Paredes¹³ on behalf of the 2D Study

¹Hospital Germans Trias i Pujol, ²Hospital de la Santa Creu i Sant Pau, ³Hospital General Universitario de Alicante, ⁴Hospital G. U. de Elche, ⁵Hospital Universitari de Bellvitge, ⁶Hospital Universitario Son Espases, ⁷Hospital Vall d'Hebron, ⁸Hospital Arquitecto Marcide, ⁹Hospital Son Llàtzer, ¹⁰Hospital General Universitario Reina Sofia de Murcia, ¹¹Hospital Universitario y Politécnico La Fe, ¹²Hospital General Nuestra Señora del Prado

METHODS

Study design and patients

- Study design: This is an open label, multicenter, randomized pilot study to evaluate the efficacy and safety of switching to DTG+DRV/c as a 2-pill once-daily simplification strategy in well-suppressed and highly ART-experienced patients harboring archived DRM against at least 2 antiretroviral classes.
- Subjects who meet all eligibility criteria were randomly assigned to:
 - Experimental group: Bithery based on DTG (50 mg QD) plus DRV/c (800/150 mg QD).
 - Control group: Continuation of their current stable ART.

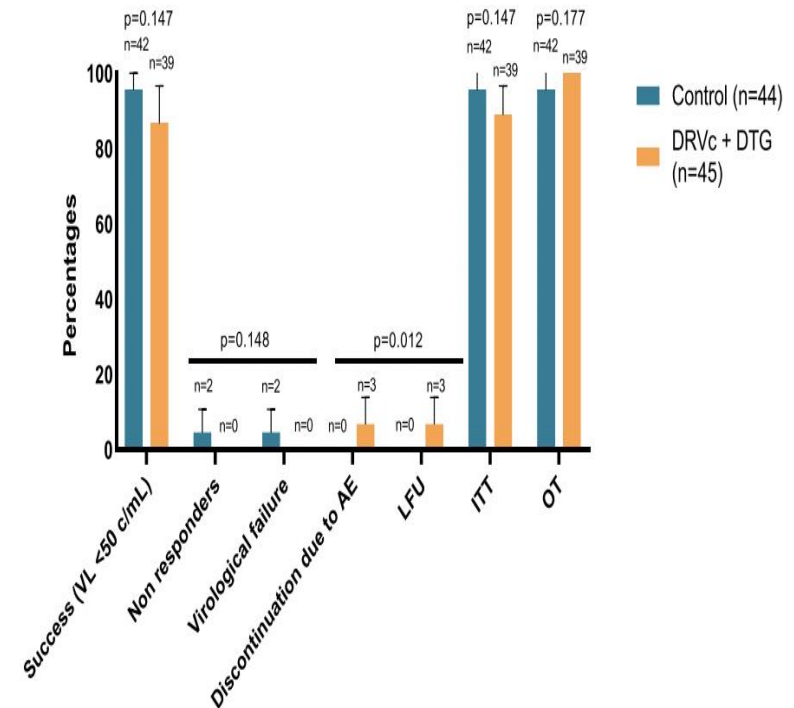


Inclusion criteria:

CONCLUSIONS

- Dual therapy with DTG+DRV/c maintains viral suppression in highly treatment experienced subjects with multiclass drug resistance, as long as they retain DRV- and INSTI-susceptible HIV-1.
- More frequently observed AEs in the DRV/c+DTG arm has to be interpreted with caution due to the open label design.
- AES leading to treatment discontinuation were unfrequent.

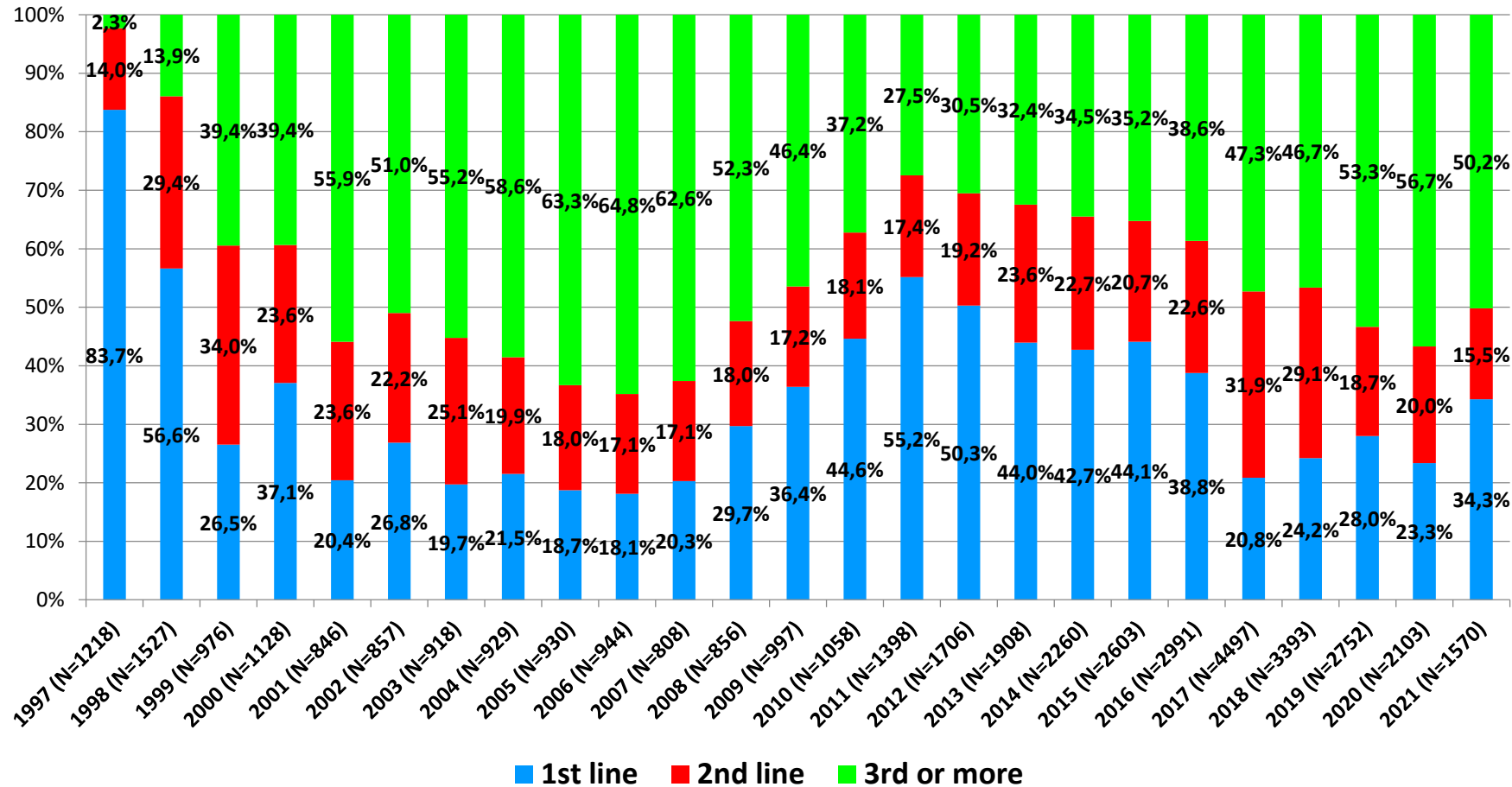
Figure 1. Virological outcome at week 48 according to on-treatment analysis



VL, viral load; LFU, loss to follow-up; ITT: M = F, intention to treat: missing equal to failure; OT: M = F, On-treatment: missing censored



Proportion of patients under different ART regimen lines according to calendar year



BIC+DRV or DTG+DRV		
Years of ART strata n	percent	
0-4	38	0,9%
5-9	44	1,3%
10-14	16	1,4%
15-19	6	1,4%
20-24	17	2,9%

Take-home Message

- HTE subjects are a challenging population, especially those with a pan-resistant virus
- New options with specific MoA are available (DOR, FTR, IBA), others will hopefully (LEN, ISL, LER..bNABS...MI...) come
- We need for new tools for correctly managing these options
- We need a multidisciplinary approach for managing HTE PLWH with the ultimate goal to achieve and maintain undetectability and possibly to simplify complex regimens