

Update on 2 drug regimens

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Disclosures

- Gilead – grants, honoraria
- Merck – grants, honoraria
- ViiV Healthcare - honoraria

Outline

Oral 2DR

- ART naives - GEMINI 1 & 2
- ART switch – TENGO
- ART switch - SWORD

Long acting injectable 2DR

- ATLAS & FLAIR combined analysis

Islatravir (MK-8591) plus doravirine 2DR

Conclusions

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Long acting injectable 2DR

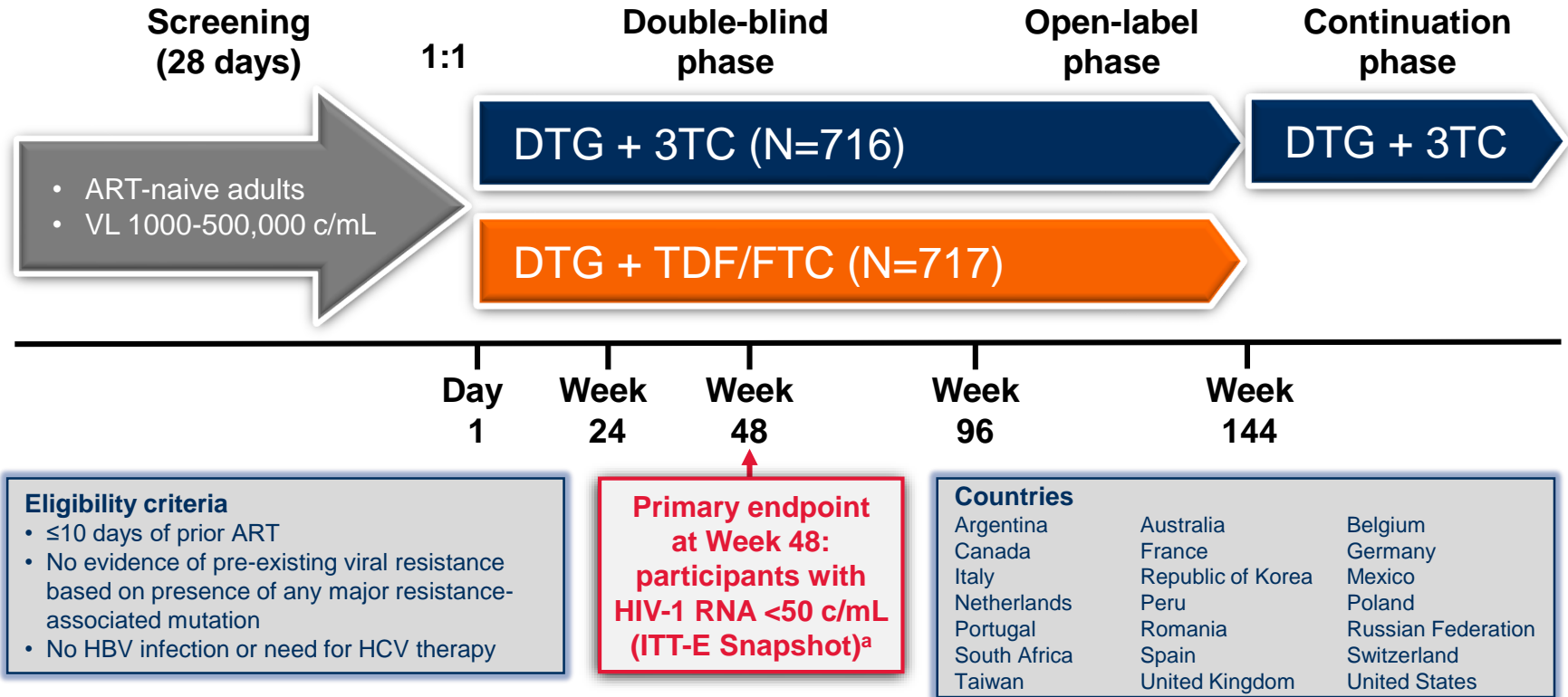
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Islatravir (MK-1598) plus doravirine 2DR

Conclusions

GEMINI-1 and -2 Phase III Study Designs

Identically designed, randomized, double-blind, parallel-group, multicenter, non-inferiority studies



Baseline stratification factors: plasma HIV-1 RNA (≤100,000 vs >100,000 c/mL) and CD4+ cell count (≤200 vs >200 cells/mm³).

^a–10% non-inferiority margin for individual studies.

Cahn et al. *Lancet*. 2018 [Epub ahead of print].

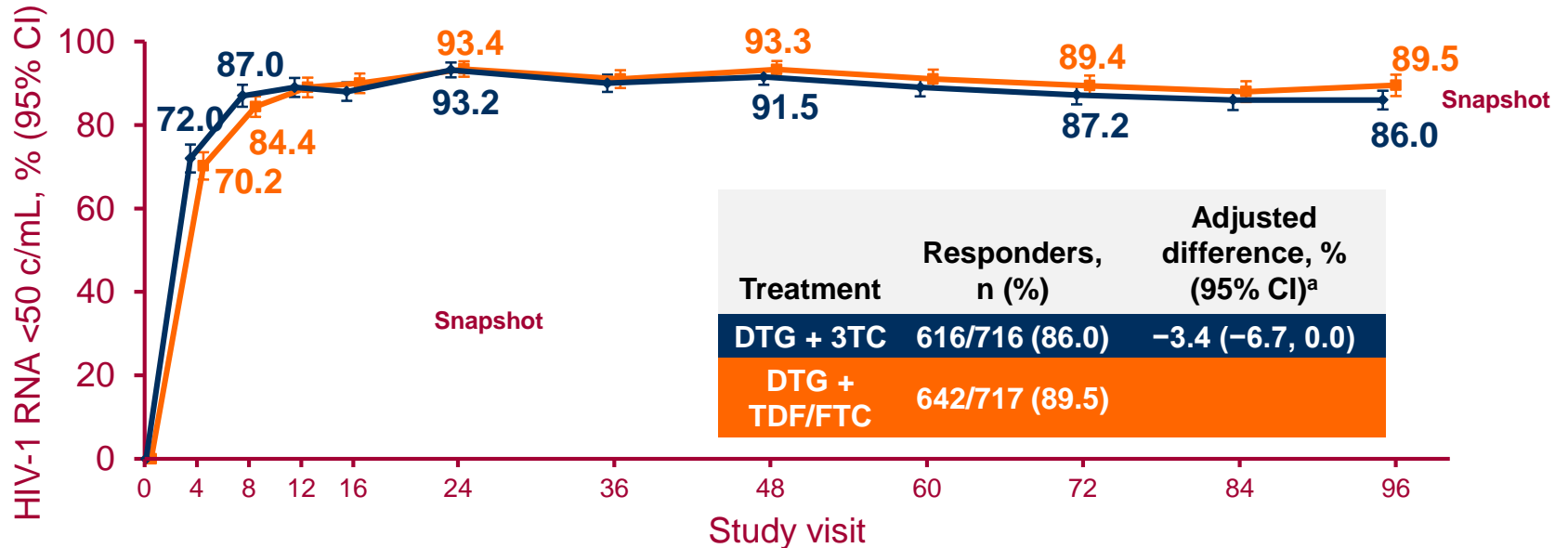
Demographic and Baseline Characteristics for the Pooled GEMINI-1 and -2 Population

Characteristic	DTG + 3TC (N=716)	DTG + TDF/FTC (N=717)
Age, median (range), y	32.0 (18-72)	33.0 (18-70)
≥50 y, n (%)	65 (9)	80 (11)
Female, n (%)	113 (16)	98 (14)
Race, n (%)		
White	480 (67)	497 (69)
African American/African heritage	99 (14)	76 (11)
Asian	71 (10)	72 (10)
Other	66 (9)	72 (10)
Ethnicity, n (%)		
Hispanic or Latino	215 (30)	232 (32)
Not Hispanic or Latino	501 (70)	485 (68)
HIV-1 RNA, median (range), log₁₀ c/mL	4.43 (1.59-6.27)	4.46 (2.11-6.37)
≤100,000	576 (80)	564 (79)
>100,000	140 (20)	153 (21)
>250,000	51 (7)	46 (6)
>400,000	18 (3)	24 (3)
>500,000 ^a	13 (2)	15 (2)
CD4+ cell count, median (range), cells/mm³	427.0 (19-1399)	438.0 (19-1497)
≤200	63 (9)	55 (8)
>200	653 (91)	662 (92)

^aParticipants were required to have HIV-1 RNA ≤500,000 c/mL at screening. Other than 1 participant enrolled without meeting study entry criteria, these participants had an observed increase in HIV-1 RNA between screening and baseline.

Cahn et al. *Lancet*. 2018 [Epub ahead of print].

GEMINI: primary endpoint (snapshot HIV-RNA <50 c/mL) at week 96



Non-inferiority criteria were met for GEMINI-1, GEMINI-2, and the pooled analysis^b

^aBased on Cochran-Mantel-Haenszel stratified analysis adjusting for the following baseline stratification factors: plasma HIV-1 RNA ($\leq 100,000$ vs $> 100,000$ c/mL), CD4+ cell count (≤ 200 vs > 200 cells/mm³), and study (GEMINI-1 vs GEMINI-2). The upper limit of the 95% CI for the pooled analysis was 0.0007%.

^bIn GEMINI-1, HIV-1 RNA <50 c/mL (95% CI) was achieved in 300/356 participants (84.3% [80.5-88.1]) in the DTG + 3TC group and 320/358 (89.4% [86.2-92.6]) in the DTG + TDF/FTC group (adjusted treatment difference [95% CI], -4.9% [-9.8, 0.03]). In GEMINI-2, the corresponding values were 316/360 (87.8% [84.4-91.2]) and 322/359 (89.7% [86.5-92.8]), respectively (adjusted treatment difference [95% CI], -1.8% [-6.4, 2.7]).

GEMINI 1 & 2: treatment emergent resistance at confirmed virological failure

Variable, n (%)	GEMINI-1		GEMINI-2		Pooled	
	DTG + 3TC (N=356)	DTG + TDF/FTC (N=358)	DTG + 3TC (N=360)	DTG + TDF/FTC (N=359)	DTG + 3TC (N=716)	DTG + TDF/FTC (N=717)
Week 48 CVW	4 (1.1)	2 (0.6)	2 (0.6)	2 (0.6)	6 (0.8)	4 (0.6)
Week 96 CVW	5 (1.4)	4 (1.1) ^a	6 (1.7)	3 (0.8)	11 (1.5)	7 (1.0) ^a
Treatment-emergent resistance	0	0	0	0	0	0

^aOne participant met the criteria for CVW at Week 12 but was not reported at the Week 48 analysis because of a laboratory reporting error identified after the Week 48 analysis.

GEMINI 1 & 2:

Adverse event profile

n (%)	DTG + 3TC (N=716)	DTG + TDF/FTC (N=717)
Any AE	591 (83)	609 (85)
AEs occurring in ≥10% of participants in either group		
Nasopharyngitis	71 (10)	114 (16)
Diarrhea	89 (12)	93 (13)
Headache	79 (11)	87 (12)
Drug-related AEs^a		
Any Grade 2-5 drug-related AEs	140 (20)	179 (25)
Grade 2-5 drug-related AEs occurring in ≥1% of participants	50 (7)	57 (8)
Headache	8 (1)	8 (1)
AEs leading to withdrawal from the study		
AEs of interest leading to withdrawal from the study	24 (3)	23 (3)
Neuropsychiatric	10 (1)	5 (1)
Renal-related	2 (<1)	7 (1)
Osteoporosis	0	2 (<1)
Any serious AE^b	64 (9)	67 (9)

- Increased weight was reported as an AE in 13 (1.8%) participants treated with DTG + 3TC and in 10 (1.4%) treated with DTG + TDF/FTC
 - Overall mean change from baseline was 3.1 kg in the DTG + 3TC group and 2.1 kg in the DTG +TDF/FTC group

^aRelative risk (95% CI) for the DTG + 3TC vs DTG + TDF/FTC group was 0.78 (0.64, 0.95).

^b3 deaths (acute myocardial infarction, n=1; Burkitt's lymphoma, n=1; coronary artery disease, n=1), 1 in GEMINI-1 and 2 in GEMINI-2; all were in the DTG + 3TC group and were considered unrelated to the study drug regimen.

GEMINI 1 & 2: VL decline through 48 Weeks: Pooled ITT-E Population

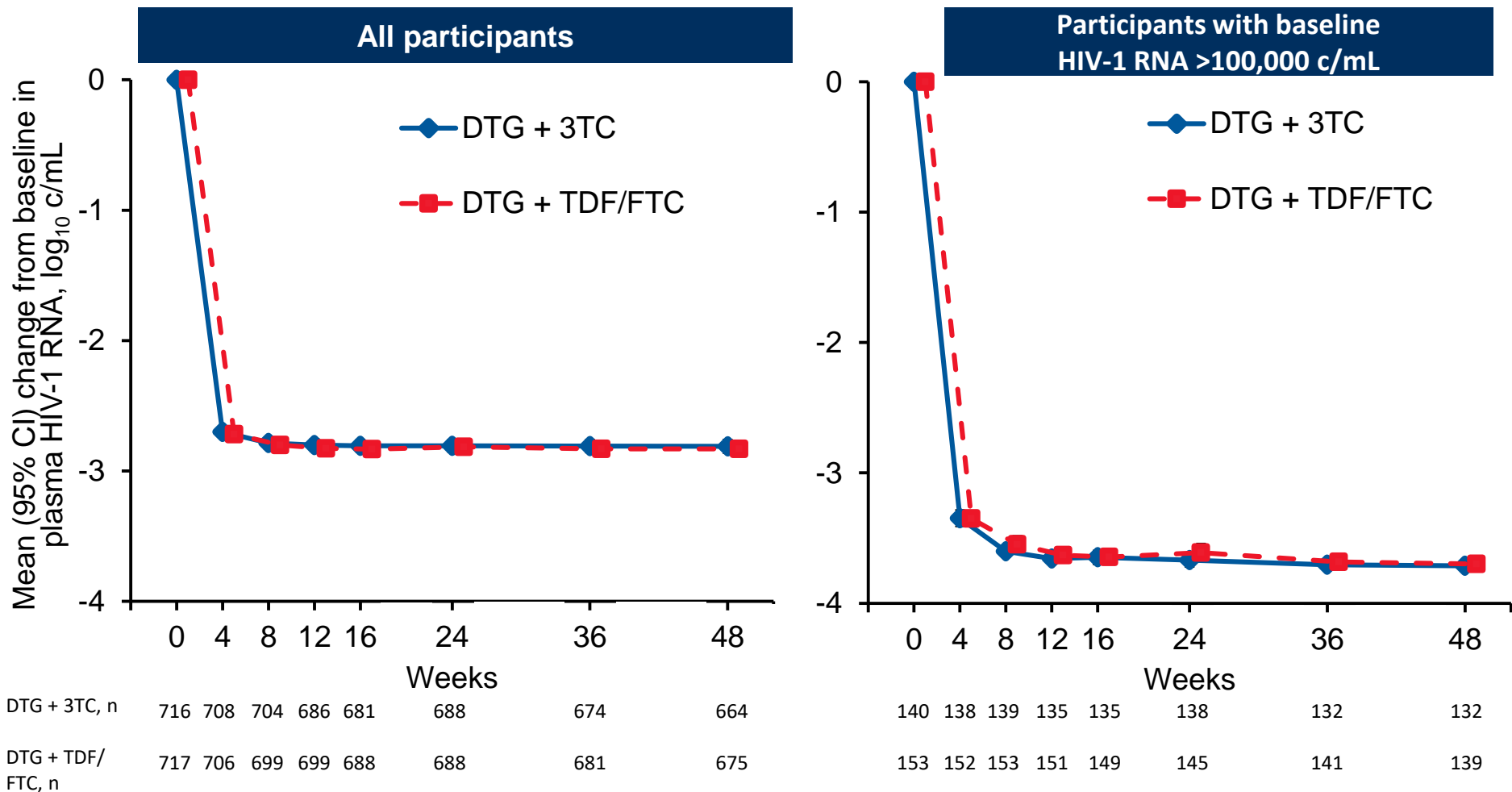
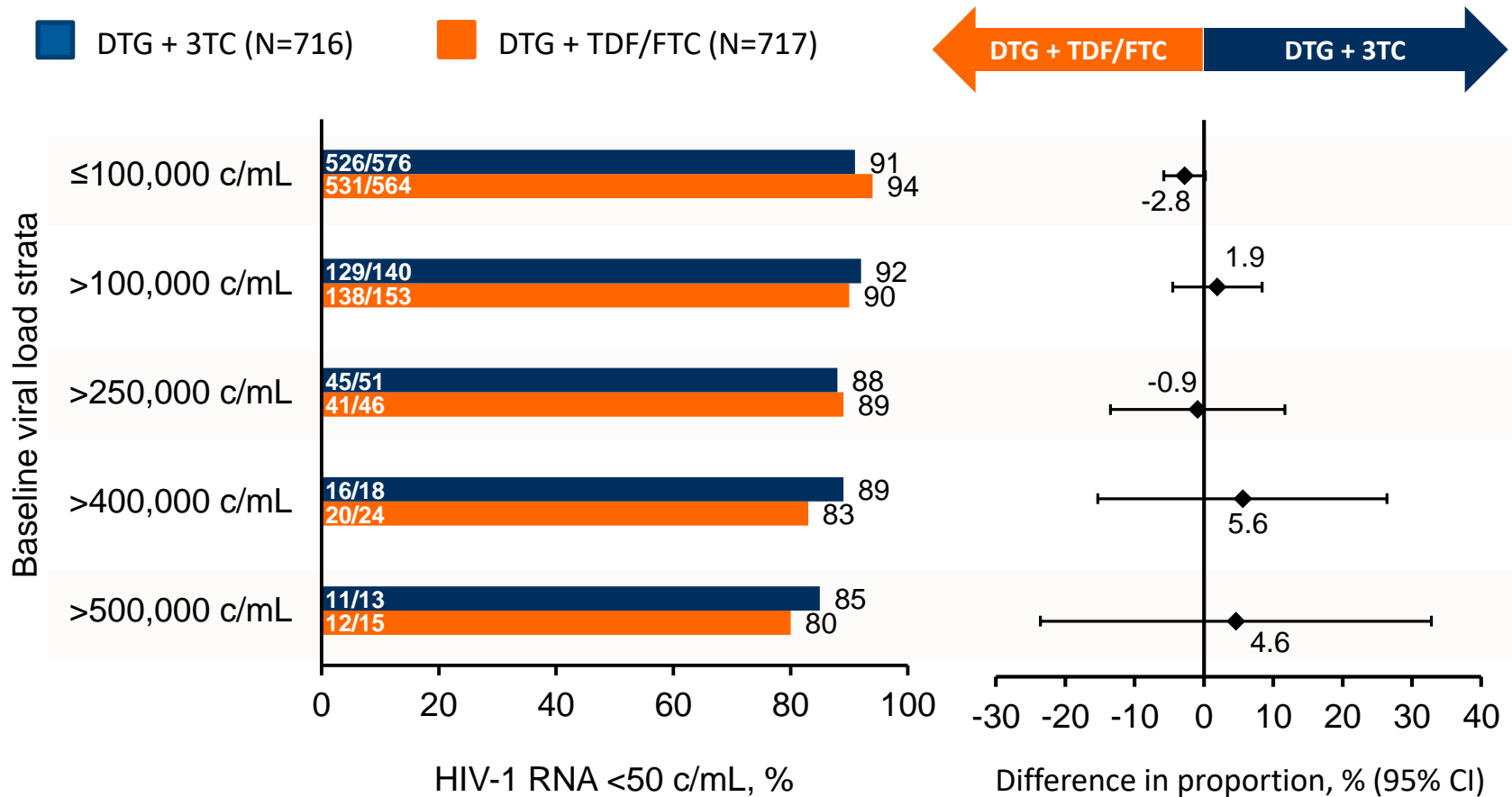


Figure on the left reproduced from Cahn et al. *Lancet*. 2018 [Epub ahead of print]. With permission from Elsevier.

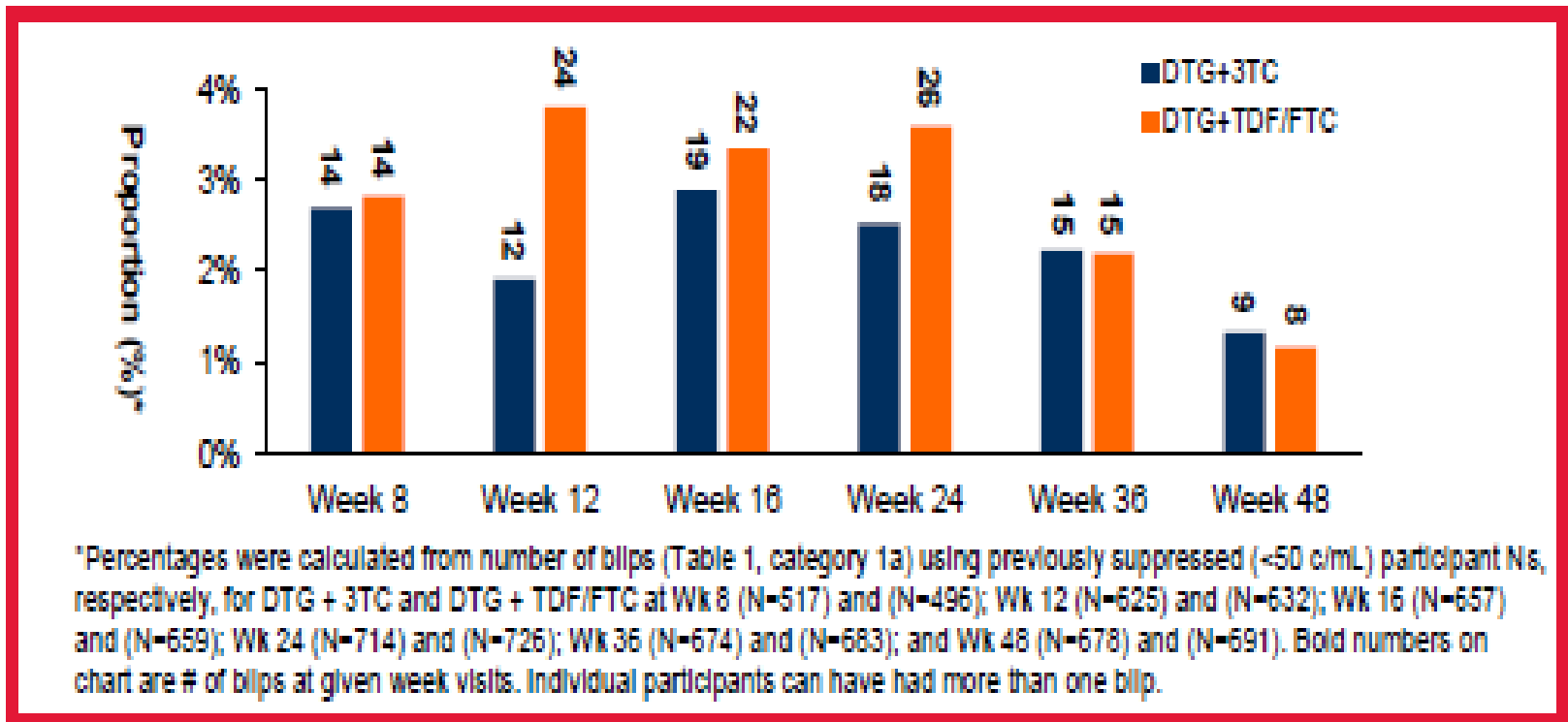
GEMINI 1 & 2: Proportion of participants with VL <50 c/mL at Week 48 (snapshot analysis) by baseline VL: Pooled ITT-E Population



Cahn et al. *Lancet*. 2018 [Epub ahead of print].

Eron et al. HIV DART and Emerging Viruses 2018; Miami, FL. Oral Presentation #7.

GEMINI 1 & 2: VL blip frequencies and number by visit week



- Similar 'blip' frequencies were seen across arms by visit week.
- Cumulative occurrences: DTG + 3TC (N=87); DTG + TDF/FTC (N=109).

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- ART switch – TENGO
- ART switch - SWORD

LA injectable 2DR

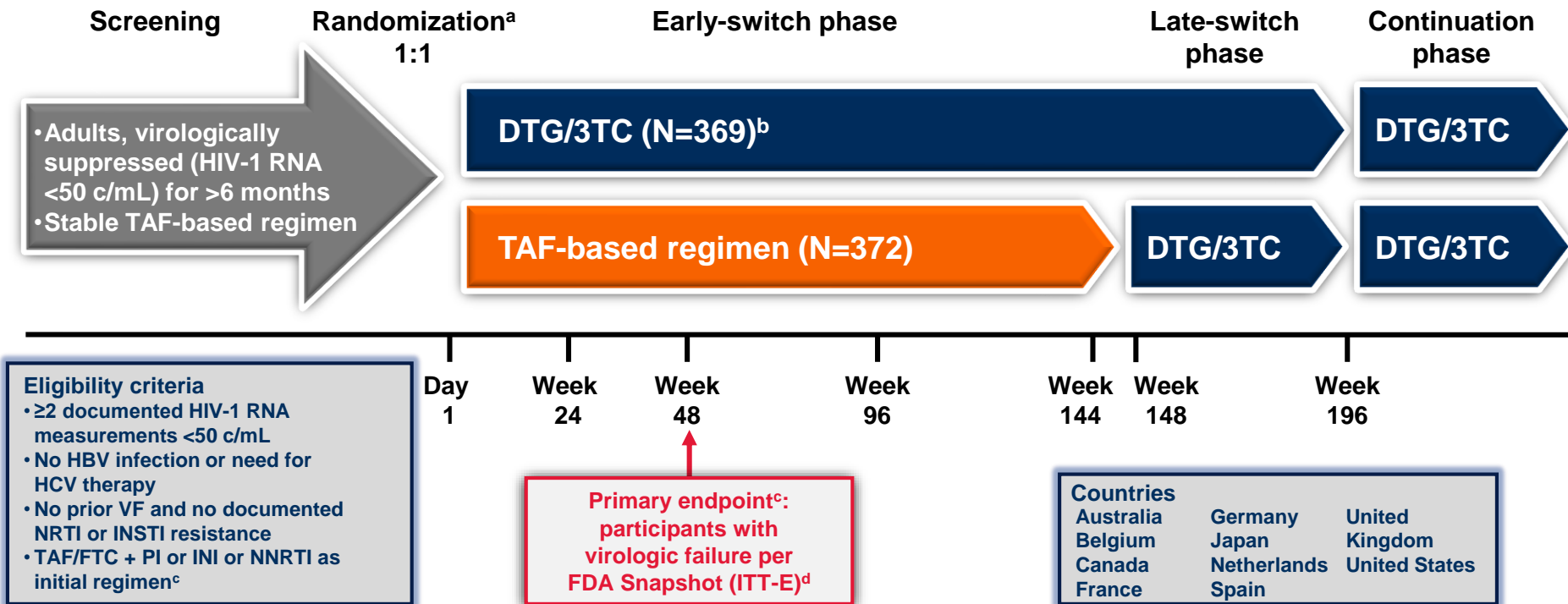
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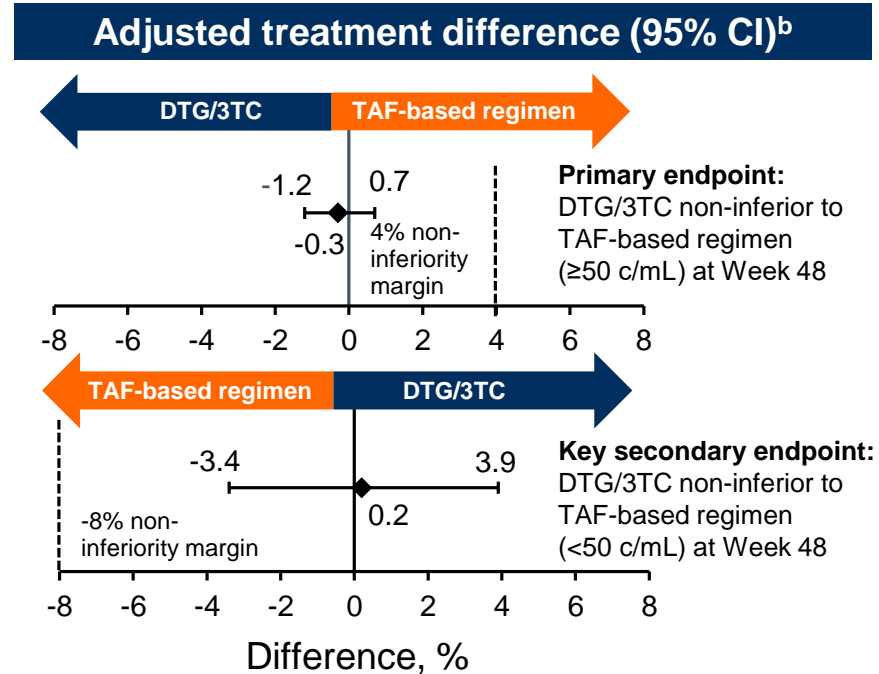
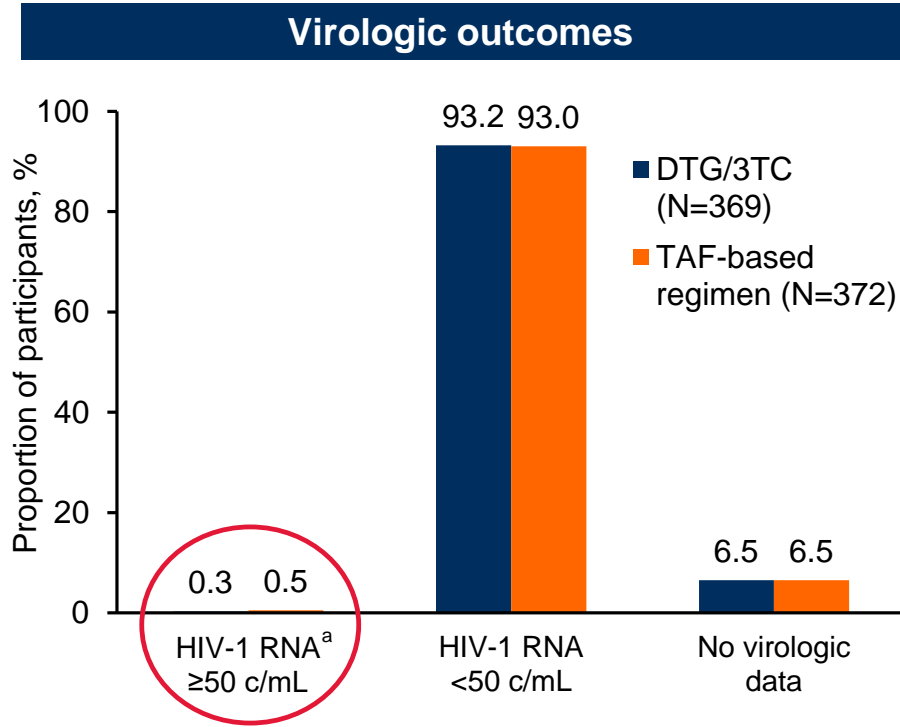
TANGO Phase III Study Design

Randomized, open-label, multicenter, parallel-group, non-inferiority study



^aStratified by baseline third agent class (PI, INI, or NNRTI). ^bTwo patients excluded who were randomized but not exposed to study drug. ^cParticipants with initial TDF treatment who switched to TAF ≥3 months before screening, with no changes to other drugs in their regimen, were also eligible. ^d4% non-inferiority margin. ^eIncludes participants who changed a background therapy component or discontinued study treatment for lack of efficacy before Week 48, or who had HIV-1 RNA ≥50 c/mL in the 48-week window.

TANGO: primary endpoint (snapshot HIV-RNA <50 c/mL) at week 48



- In the per-protocol population, 0/352 participants in the DTG/3TC group and 2/358 participants in the TAF-based regimen group had HIV-1 RNA ≥ 50 c/mL at Week 48 (adjusted difference, -0.6 ; 95% CI, -1.3 to 0.2)^b

^aPrimary endpoint (Snapshot virologic non-response, ITT-E). ^bBased on Cochran-Mantel-Haenszel stratified analysis adjusting for baseline third agent class.

TANGO:

Treatment emergent resistance at confirmed virological failure

n (%)	DTG/3TC (N=369)	TAF-based regimen (N=372)
Confirmed virologic withdrawal (CVW) ^a	0	1 (<1) ^b
Observed resistance mutation at failure ^c	0	0

^aOne assessment with HIV-1 RNA ≥ 200 c/mL after Day 1 with an immediately prior HIV-1 RNA ≥ 50 c/mL.

^bTreatment interrupted before suspected virologic withdrawal (VL, 38,042 c/mL) and resumed 3 weeks before VL retest (297 c/mL).

^cPlasma HIV-1 RNA resistance genotype at failure is compared with baseline PBMC pro-viral resistance genotype.

TANGO: Adverse event profile

n (%)	DTG/3TC (N=369)	TAF-based regimen (N=371)
Any AE	295 (80)	292 (79)
Nasopharyngitis	43 (12)	41 (11)
Upper respiratory tract infection	31 (8)	32 (9)
Diarrhea	30 (8)	26 (7)
Headache	24 (7)	17 (5)
Syphilis	24 (7)	13 (4)
Back pain	21 (6)	28 (8)
Fatigue	20 (5)	3 (1)
Bronchitis	8 (2)	20 (5)
Any drug-related Grade 2-5 AE	17 (5)	3 (1)
Drug-related Grade 2-5 AEs occurring in $\geq 0.5\%$^a		
Insomnia	4 (1)	0
Constipation	2 (1)	1 (<1)
Flatulence	2 (1)	0
Headache	2 (1)	0
AEs leading to withdrawal from the study	13 (4)^b	2 (1)
Drug-related AEs leading to withdrawal from the study	9 (2)	1 (<1)
Any SAE^c	21 (6)^b	16 (4)

- At Week 48, a similar adjusted mean increase from baseline in weight of 0.8 kg was observed in both treatment groups
- Increased weight was reported as an AE in 3 (1%) participants treated with DTG/3TC and in 6 (2%) treated with a TAF-based regimen

SAE, serious adverse event.

^aAll drug-related AEs were of grade 2. ^bOne fatal AE occurred (homicide). ^cNo SAEs were drug related.

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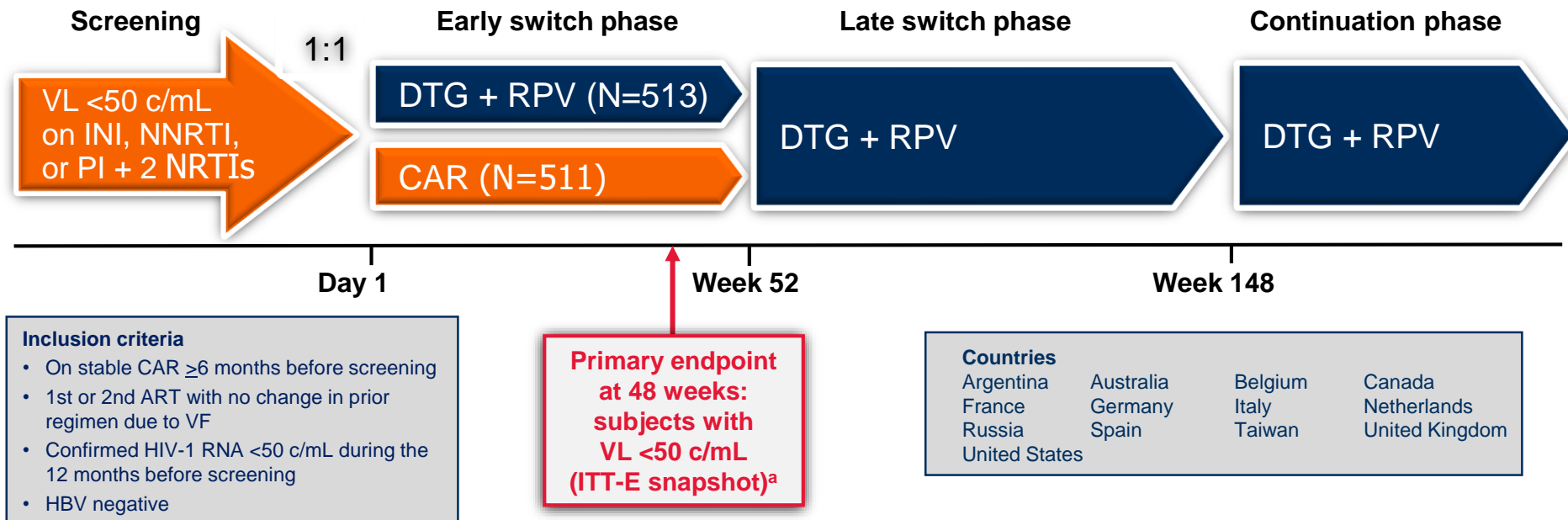
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SWORD 1 & 2 Phase III Study Design

Identically designed, randomized, multicenter, open-label, parallel-group, non-inferiority studies



^a-8% non-inferiority margin for pooled data. -10% non-inferiority margin for individual studies

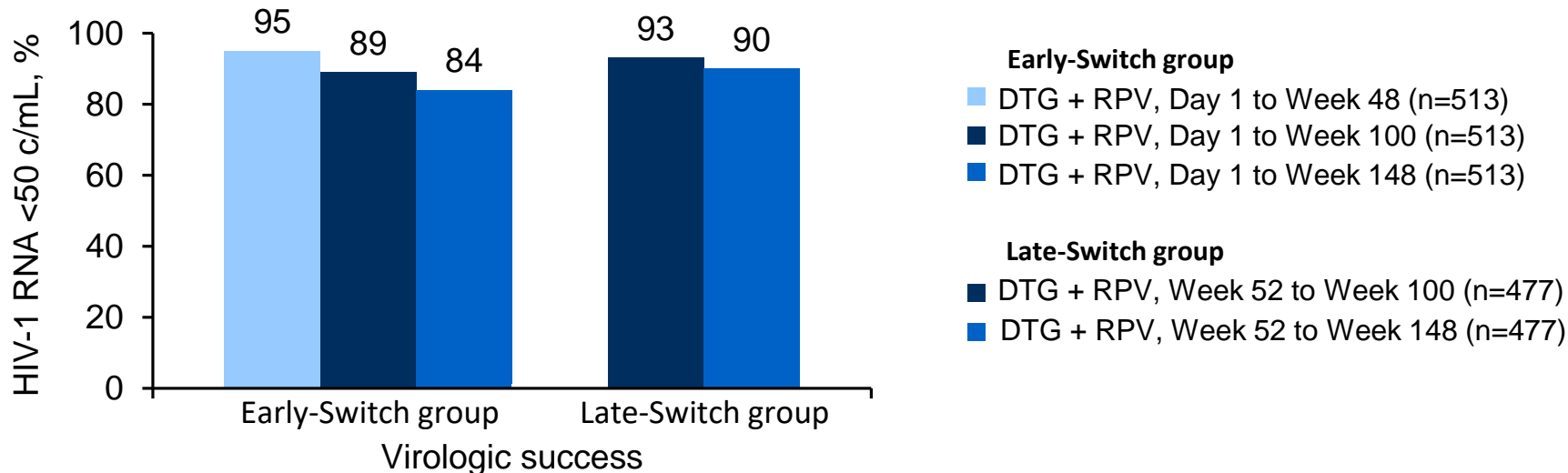
SWORD 1 & 2: HIV-1 RNA <50 c/mL (FDA snapshot) at weeks 48, 100, and 148

Study Disposition

- Overall, 990 participants received DTG + RPV treatment (Early-Switch group, n=513; Late-Switch group, n=477)

Virologic Efficacy

- Through 148 weeks of treatment, DTG + RPV continued to be efficacious in the Early-Switch group
- Virologic efficacy in the Late-Switch group at Week 148 was similar to that in the Early-Switch group at Week 100



SWORD 1 & 2: confirmed virologic failure from week 100-148 in those exposed to DTG + RPV

- Through Week 148, there was a low number of confirmed virologic withdrawals (CVWs) across study populations who received DTG + RPV (1%; 11/990)
- Three CVWs occurred between the Week 100 analysis and Week 148

Week of failure	Previous regimen	Viral loads, copies/mL ^c	Resistance mutations ^b				Fold change	
			Baseline (GenoSure ^d)		Confirmed virologic withdrawal		DTG	RPV
			NNRTI	INSTI	NNRTI	INSTI		
W112	RAL/TDF/FTC	<u>118</u> ; 230; 324	None	E157Q, G193E, T97T/A	M230M/L	E157Q, G193E	1.47	2
W112	DRV, RTV, TDF/FTC	<u>148</u> ; 219; 307	ND		ND		ND	ND
W136 ^e	EFV/TDF/FTC	<u>4294</u> ; 7247; 40,020; 3378	NR ^f		E138A, L100L/I	NR ^f	—	4.14

ND, not determined; NR, not reported. ^aThere was an additional participant with confirmed virologic withdrawal at Week 160. However, blood collected with the first elevated viral load was collected as a protocol deviation on the day of DTG + RPV reinitiation after treatment interruption due to injuries sustained in a motor vehicle accident. ^bShading represents participants with NNRTI RAMs. ^cUnderlined value denotes viral load when participant met virologic withdrawal. ^dHIV-1 baseline resistance testing was performed on integrated HIV-1 proviral DNA using GenoSure Archive[®] assay (Monogram Biosciences, South San Francisco, CA). On-study resistance testing used standard plasma-based genotypic and phenotypic resistance testing. ^eParticipants in the Late-Switch group. ^fSample failed testing.

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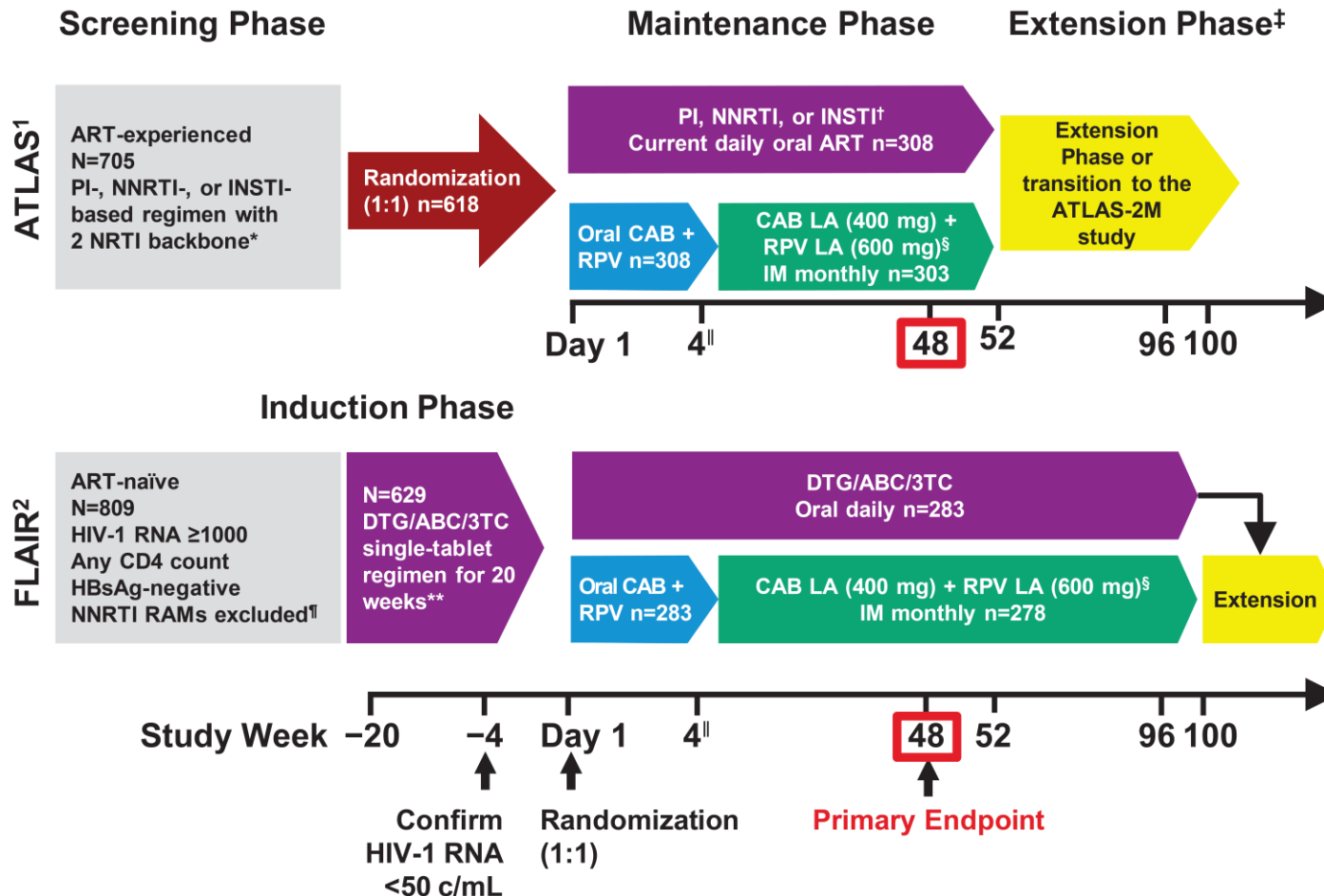
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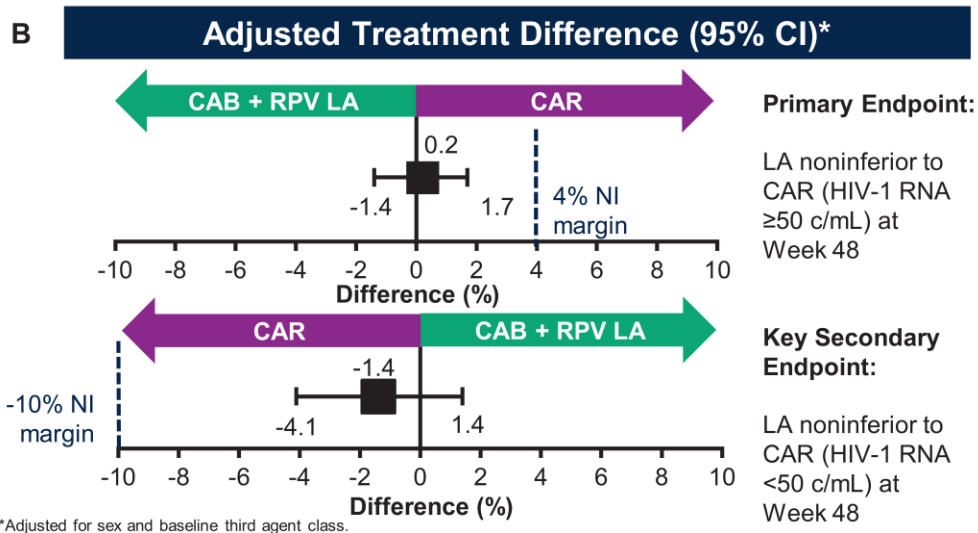
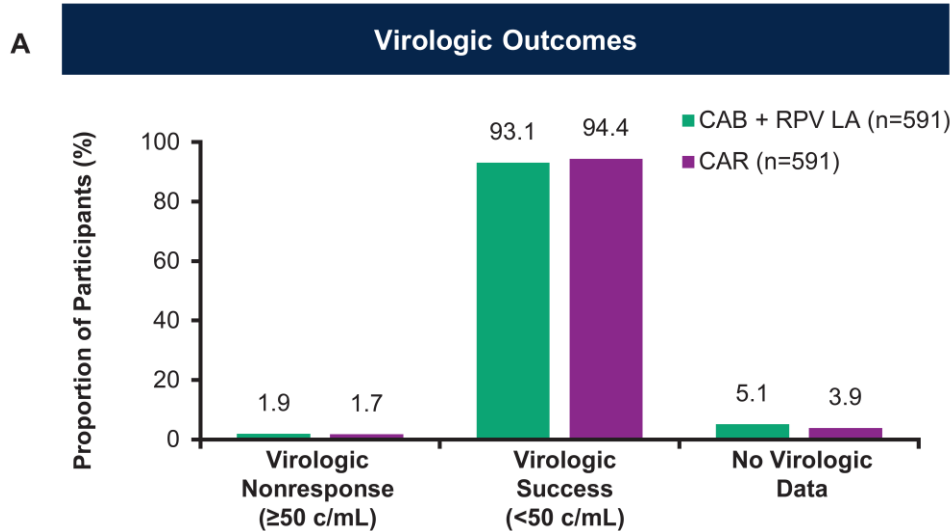
Conclusions

ATLAS and FLAIR Study Design



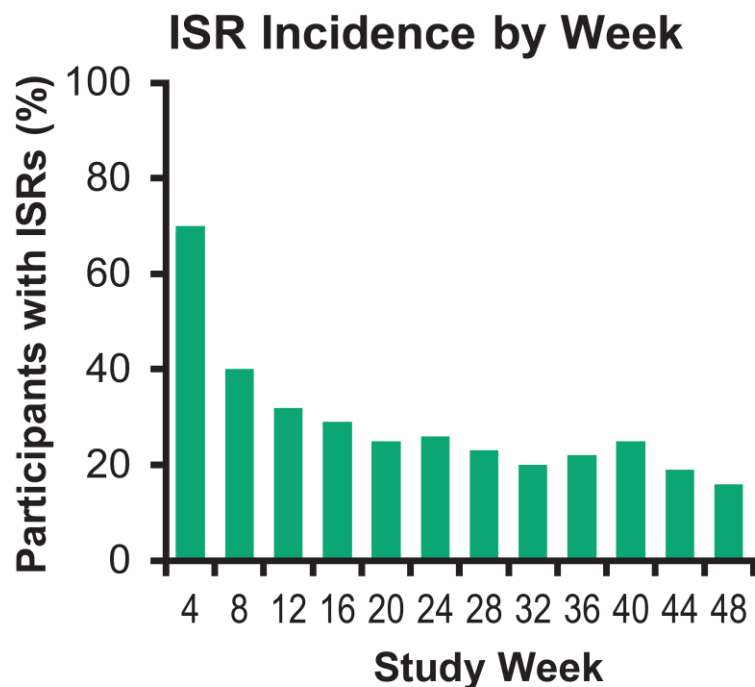
*Uninterrupted ART ≥ 6 months and VL <50 c/mL at Screening, 2 × VL <50 c/mL ≤12 months; †INSTI-based regimen capped at 40% of enrollment; ‡Triumeq excluded from study; ‡Optional switch to CAB + RPV LA at Week 52 for those on CAR; §Participants who withdraw/complete IM CAB + RPV LA must complete 52 weeks of follow-up; ¶Participants received initial loading dose of CAB LA (600 mg) and RPV LA (900 mg) at Week 4. From Week 8 onwards, participants received CAB LA (400 mg) + RPV LA (600 mg) injections every 4 weeks; ¶NNRTI RAMs, but not K103N, were exclusionary; **DTG plus two alternative non-ABC NRTIs was permitted if participant was intolerant or HLA-B*5701-positive.

ATLAS and FLAIR combined analysis: primary endpoint (ITT-E)



Pooled Injection Site Reactions

- The majority (99%, 3628/3663) of ISRs were Grade 1–2 and most (88%) resolved within ≤ 7 days



Event	CAB + RPV LA N=591
Participants receiving injections, n	581
Injections given, n	14682
ISR events, n (%)	3663 (24.9)
Pain	3087 (21.0)
Nodule	140 (1.0)
Induration	136 (0.9)
Swelling	86 (0.6)
Grade 3 ISR pain	32 (0.2)
Median duration of ISRs, days	3
Participants with ISR leading to withdrawal, n (%)	6 (1)

Bars represent incidence of onset ISRs relative to the most recent IM injection visit.

Overton et al. IAS 2019; Mexico City, Mexico. Poster MOPEB257.

Press Release 22 August 2019: [ViiV reports ATLAS-2M has met its primary endpoint.](#)

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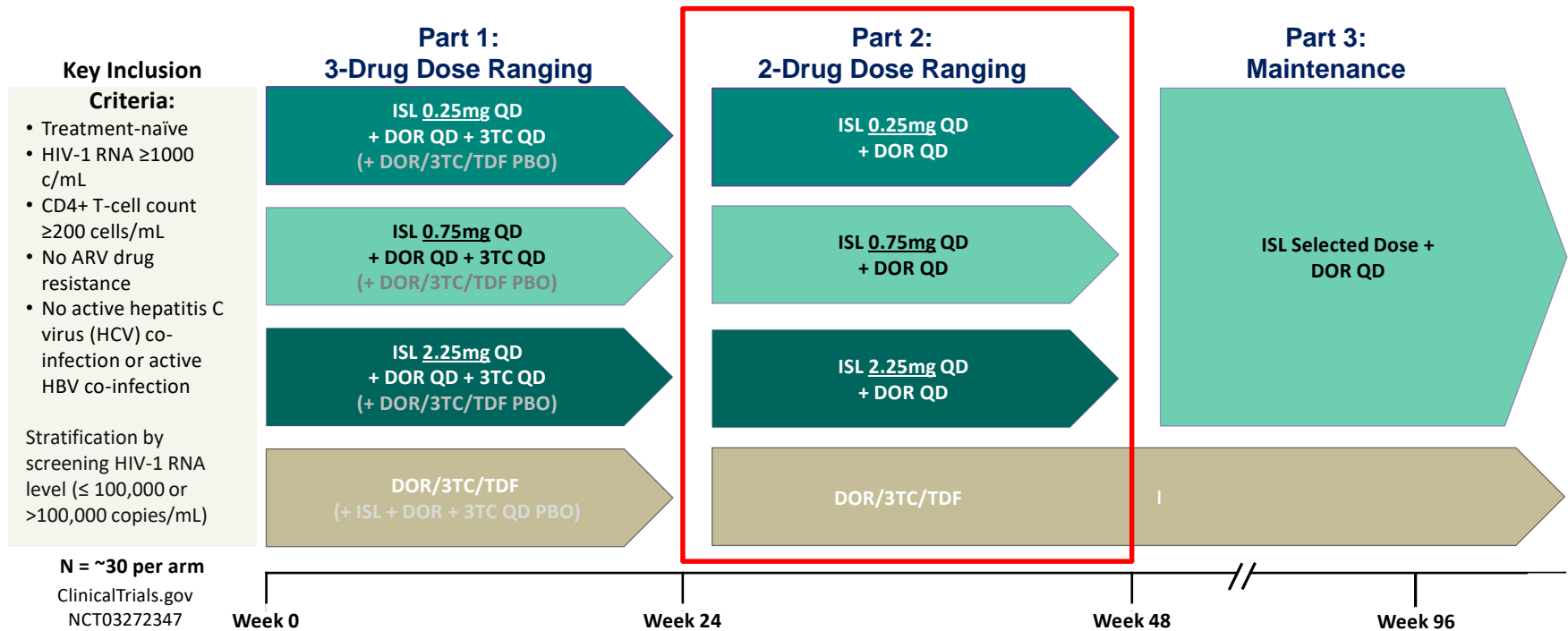
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Islatravir plus doravirine 2DR

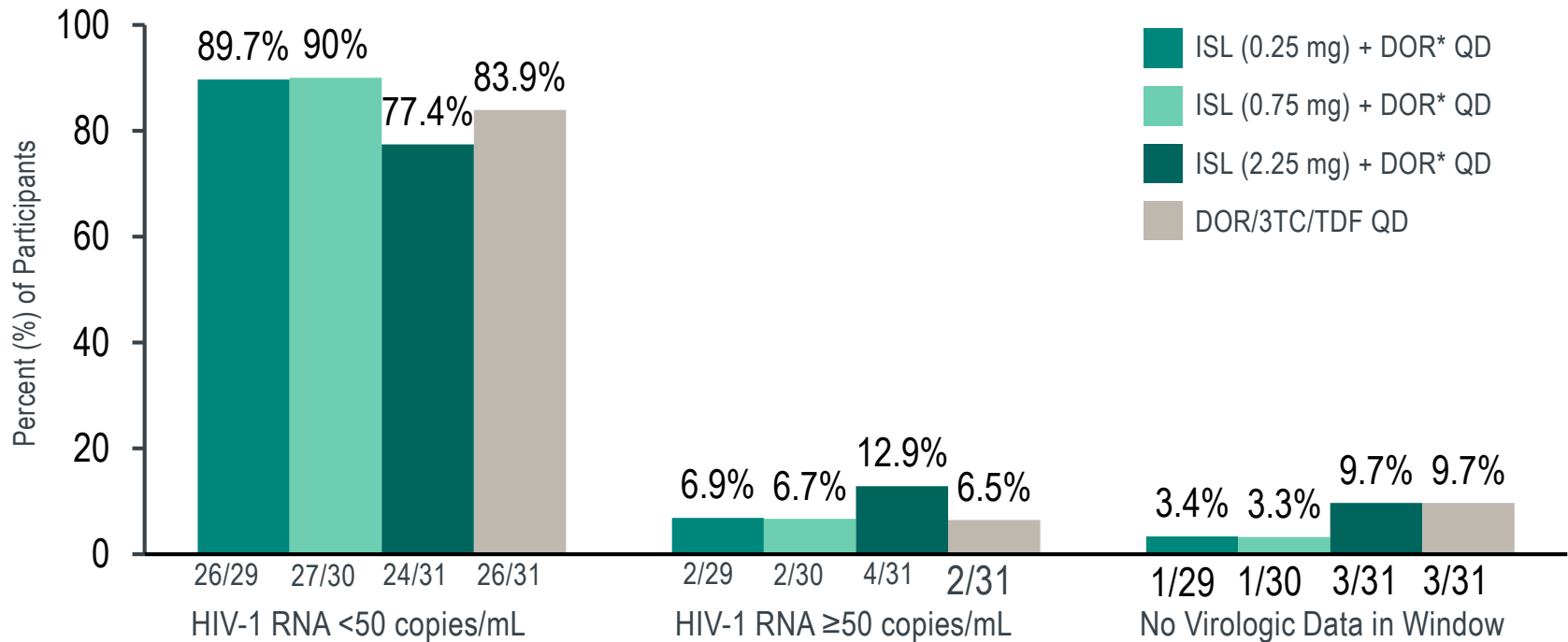
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MK-8591 Protocol 011: Phase 2 Dose Ranging Trial of ISL + DOR



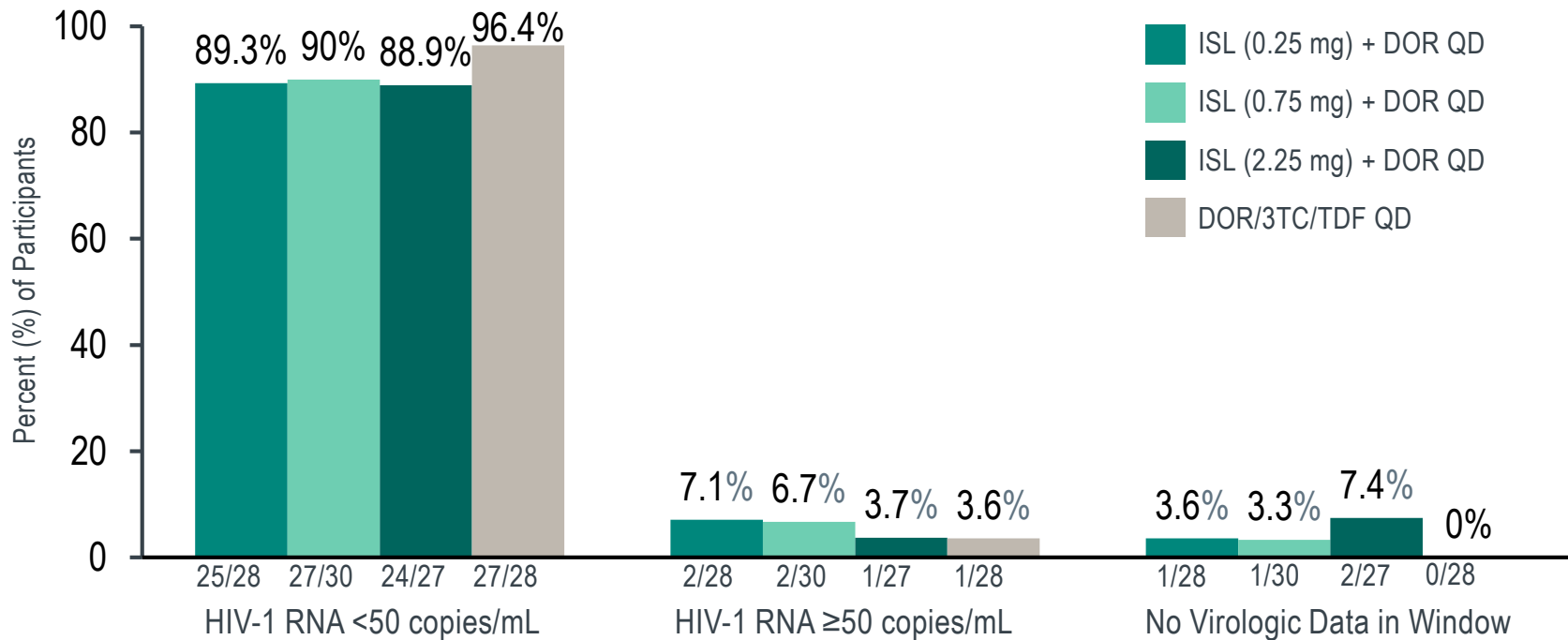
After 24 weeks of dosing in Part 1, participants who are virologically suppressed (HIV-1 RNA < 50 copies/mL) at the Week 20 visit and have not met any viral failure criteria are eligible to switch to Part 2 of the trial at Week 24. Participants with HIV-1 RNA levels ≥ 50 copies/mL at Week 20 will remain in Part 1 until the HIV-1 RNA is < 50 copies/mL and they have not met any of the viral failure criteria, at which point they transition to Part 2 at their next visit.

Virologic outcomes through week 48 (FDA snapshot approach)



*Participants initially received ISL+DOR+3TC and switched to ISL+ DOR during the week 24-48 period of the study.

Virologic outcomes 24 weeks after entering Part 2 (FDA snapshot approach)



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2DR DTG-3TC is non-inferior to 3DR over 96 weeks in ART naives

- More data required in use in women, high baseline VLs, M184V/I resistance (GART and proviral DNA) and test & treat

2DR switch to DTG/RPV or DTG/3TC appears safe and effective

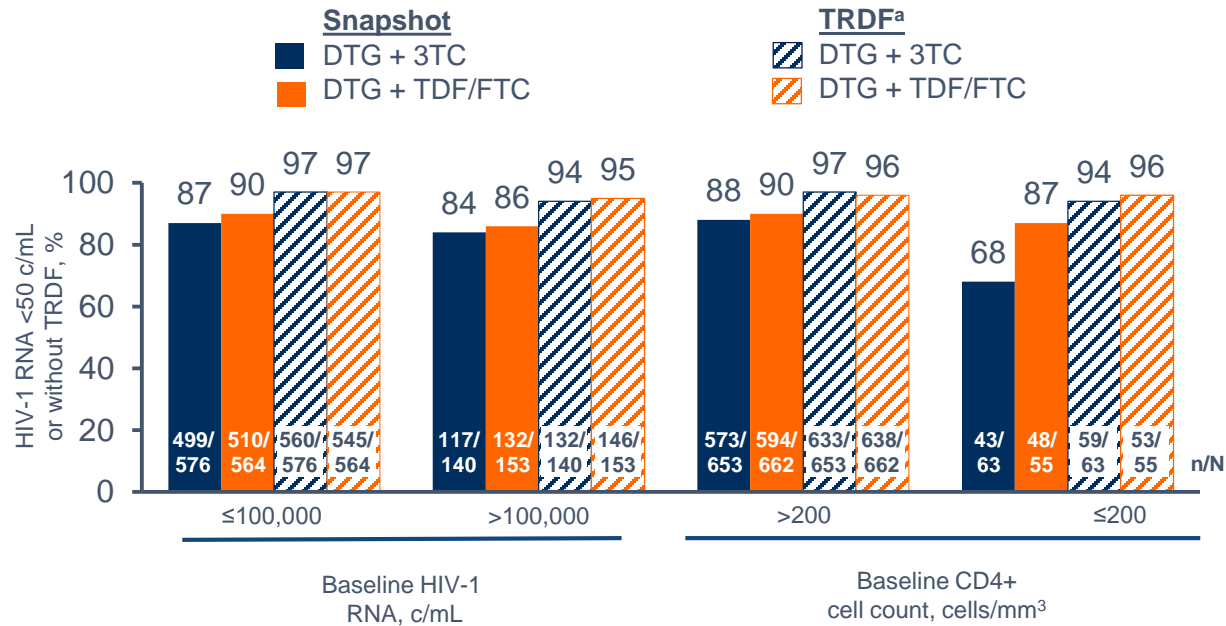
Another drug company exploring 2DR is intriguing

- But what can Merck partner with Islatravir?

Are we at a point when we can ask ‘why 3DR’?

Back up slides

GEMINI 1 & 2: proportion of participants with HIV RNA <50c/mL by baseline VL and CD4+ count: Snapshot and TRDF analysis



- At Week 96, there were 3 confirmed virologic withdrawals in the DTG + 3TC group and 2 in the DTG + TDF/FTC group in the CD4 ≤ 200 stratum

TRDF, treatment-related discontinuation equals failure. ^aTRDF was a pre-planned analysis at Week 96.

SWORD 1 & 2: Adverse Events with onset through Week 52

	Early switch phase ^a	
	DTG + RPV (n=513) n (%)	CAR (n=511) n (%)
Any AE	395 (77)	364 (71)
AEs occurring in ≥5% of subjects in either group		
Nasopharyngitis	49 (10)	50 (10)
Headache	41 (8)	23 (5)
Upper respiratory tract infection	24 (5)	37 (7)
Diarrhea	32 (6)	27 (5)
Back pain	15 (3)	31 (6)
Any Serious AEs¹	27 (5)	21 (4)
Drug-related AEs		
Grades 1-2	89 (17)	8 (2)
Grades 3-4	8 (2)	1 (<1)
AEs leading to withdrawal from the study	21 (4)	3 (<1)
CNS AEs leading to withdrawal	9 (2)	1 (<1)

^aData pooled across SWORD-1 and SWORD-2. ^bTwo deaths in the study, both unrelated to study drug. DTG + RPV, Kaposi's sarcoma (n=1); CAR, lung cancer (n=1).

Resistance-Associated Mutations

- CVWs with resistance-associated mutations (RAMS) were low across both groups and were detected in 6 (0.6%) participants
 - RAMs were not detected at baseline in 4 of these participants; in 2 participants, GenoSure testing was not available at baseline
 - No participants had INSTI RAMs at CVW time point; 1 participant with only polymorphic INSTI V151V/I (no impact on DTG susceptibility) had INSTI RAM mixtures N155N/H and G163G/R at baseline