

Effectiveness of Dolutegravir + Lamivudine in Real-world Studies in People With HIV-1 With M184V/I Mutations: A Systematic Review and Meta-analysis

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- Using real-world data from people with HIV-1 (PWH), a systematic literature review and a meta-analysis were performed to investigate the impact of historical or archived M184V/I on the effectiveness of dolutegravir + lamivudine (DTG + 3TC) in real-world switch populations; a sensitivity analysis was performed using data from randomized controlled trials (RCTs) identified via a targeted literature review
- Virologic failure (VF) incidence was low, and no treatment-emergent INSTI resistance mutations were reported in populations with M184V/I that switched to DTG + 3TC, providing reassurance that M184V/I may have a limited impact on the efficacy of DTG + 3TC in PWH considering treatment change when drug resistance-associated mutations (RAMs) are unknown or inadvertently missed

Introduction

- M184V/I is the most common RAM selected by 3TC¹
- Clinical development phase 3 RCTs excluded participants with known or suspected RAMs
- The presence of archived M184V/I mutations in phase 3 trials evaluating switch to DTG/3TC (TANGO, n=4; SALSA, n=5)^{2,3} did not impact virologic efficacy
- Absence of historical resistance results or availability of prior genotype (pooled TANGO/SALSA analysis, n=294) also had no impact on results⁴
- In clinical practice, prior history of resistance is not always available when considering treatment options • Real-world evidence (RWE) can help address the knowledge gap of whether switching to DTG + 3TC is safe in real-world clinical practice when full treatment history or historical genotype results are not available

Figure 1. PRISMA Flow Charts for (A) RWE Studies and (B) RCTs





- This meta-analysis describes VF at Weeks 24, 48, and 96 using real-world data from PWH receiving DTG + 3TC in a suppressed switch setting, with historical RNA- or archived proviral DNA-detected M184V/I mutation
- A sensitivity analysis was performed using RCT data

Methods

- A systematic literature review was performed following Preferred Reporting Items for Systematic Reviews and Meta-Analyses guidelines (Figure 1A)
- Embase[®], Ovid MEDLINE[®], MEDLINE[®] In-Process, and Cochrane library (January 2013-March 2022) and relevant conference archives (2016-2021) were searched for real-world studies reporting virologic outcomes for PWH receiving DTG + 3TC
- A targeted literature review was performed to identify RCTs assessing M184V/I impact on DTG + 3TC efficacy (Figure 1B)
- Studies were screened for suppressed switch populations reporting M184V/I mutations before DTG + 3TC initiation
- For the primary objective, common- and random-effects model analyses were conducted using RWE studies
- Random-effects models provide estimates that are more generalizable to the overall population of interest
- Common-effects (or fixed-effects) models assume that the included studies are the population of interest and are more informative when zero VF events are observed
- For the secondary objective, sensitivity analyses were performed using RCT data
- In both RWE and RCT data sets, base analyses were performed using studies with identical VF definitions; sensitivity analyses were performed using all studies regardless of VF definition to maximize sample size

Results

VF Outcomes in RWE Studies and RCTs

- Of 3492 publications and 198 conference abstracts identified via systematic literature review, 5 real-world studies met all search criteria and were analyzed (Table)
- The targeted literature review also identified 5 relevant RCTs
- Proportions of PWH with historical M184V/I estimated to have VF at Weeks 24, 48, and 96 were low in

Figure 2. Meta-analysis Estimates of Proportions of VF at Weeks 24, 48, and 96 in PWH With Reported M184V/I Receiving DTG + 3TC From (A) Systematic Literature Review–Identified RWE Studies and (B) Targeted Literature Review–Identified RCTs, Inclusive of All VF Definitions

A. RWE studies

Week 24	Study Ev Hocqueloux 2021 Santoro 2021 Borghetti 2021	vents 1 2 0	Total 105 36 45		Proportion 0.01 0.06 0.00	95% Cl (0.00-0.05) (0.01-0.19)
	Random-effects model	0			0.00	(0.00-0.08) (0.00-0.14)
	Heterogeneity: /² = 56%, τ² = 0.0063, <i>P</i> =	=0.10		0 0.2 0.4 0.6 0.8 1 Proportion of individuals with VF		
	Study Ev	vents	Total		Proportion	95% CI
Week 48	Hocqueloux 2021 Hidalgo-Tenorio 2019 Santoro 2021 Galizzi 2020 Borghetti 2021	2 1 2 1 1	105 4 36 47 45		0.02 0.25 0.06 0.02 0.02	(0.00-0.07) (0.01-0.81) (0.01-0.19) (0.00-0.11) (0.00-0.12)
	Random-effects model		237		0.03	(0.01-0.08)
	Heterogeneity: $/^2 = 25\%$, $\tau^2 = 0$, $P=0.25$			0 0.2 0.4 0.6 0.8 1 Proportion of individuals with VF		
Week 96	Study Ev	vents	Total		Proportion	95% CI
	Hocqueloux 2021 Santoro 2021 Borghetti 2021	2 3 2	105 36 45		0.02 0.08 0.04	(0.00-0.07) (0.02-0.22) (0.01-0.15)
	Random-effects model		186		0.04	(0.01-0.17)
	Heterogeneity: $/^2 = 27\%$, $\tau^2 = 0$, $P=0.26$			0 0.2 0.4 0.6 0.8 1 Proportion of individuals with VF		(, ,
B. R	CTs					
B. R	CTs Study DOLULAM TANGO ART PRO	vents 0 0 0	Total 17 4 17		Proportion 0.00 0.00 0.00	95% Cl (0.00-0.20) (0.00-0.60) (0.00-0.20)
Neek 24	CTs Study DOLULAM TANGO ART PRO Common-effects model	vents 0 0 0	Total 17 4 17 38		Proportion 0.00 0.00 0.00 0.00	95% Cl (0.00-0.20) (0.00-0.60) (0.00-0.20) (0.00-0.03)
Week 24	CTsStudyExDOLULAM TANGO ART PROExCommon-effects model Heterogeneity: $/^2 = 0\%$, $\tau^2 = 0$, $P=1.00$	vents 0 0 0	Total 17 4 17 38	0 0.2 0.4 0.6 0.8 1 Proportion of individuals with VF	Proportion 0.00 0.00 0.00 0.00	95% Cl (0.00-0.20) (0.00-0.60) (0.00-0.20) (0.00-0.03)
Neek 24	CTsStudyExDOLULAM TANGO ART PROExCommon-effects model Heterogeneity: $/^2 = 0\%$, $\tau^2 = 0$, $P=1.00$ StudyEx	vents 0 0 0	Total 17 4 17 38 Total	0 0.2 0.4 0.6 0.8 1 Proportion of individuals with VF	Proportion 0.00 0.00 0.00 0.00 0.00	95% Cl (0.00-0.20) (0.00-0.60) (0.00-0.20) (0.00-0.03) 95% Cl
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Week 48 Week 24 B	StudyExDOLULAM TANGO ART PROExCommon-effects model Heterogeneity: $/^2 = 0\%$, $\tau^2 = 0$, $P=1.00$ StudyExSOLAR 3D DOLULAM SALSA TANGO ART PRORandom-effects model	vents 0 0 0 0 vents 1 0 1 0 0	Total 17 4 17 38 38 Total 50 17 5 4 17 5 4 17 93	0 0.2 0.4 0.6 0.8 1 Proportion of individuals with VF	Proportion 0.00 0.00 0.00 0.00 0.00 Proportion 0.02 0.00 0.20 0.00 0.20 0.00 0.00 0.00 0.00	95% Cl (0.00-0.20) (0.00-0.60) (0.00-0.20) (0.00-0.03) 95% Cl (0.00-0.11) (0.00-0.20) (0.01-0.72) (0.00-0.60) (0.00-0.20) (0.00-0.20)
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k 96 Week 48 Week 24 B	CTsStudyExDOLULAM TANGO ART PROExCommon-effects model Heterogeneity: $/^2 = 0\%$, $\tau^2 = 0$, $P=1.00$ StudyExSOLAR 3D DOLULAM SALSA TANGO ART PROExRandom-effects model Heterogeneity: $/^2 = 11\%$, $\tau^2 = <0.0001$, F StudyExDOLULAM ART PROExDOLULAM ART PROEx	vents 0 0 0 0 vents 1 0 1 0 0 2 =0.34	Total 17 4 17 38 Total 50 17 50 17 50 17 93 Total 17 93	0 0.2 0.4 0.6 0.8 1 Proportion of individuals with VF	Proportion 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.02 0.00 0.20 0.00 0.00 0.00 0.00 0.00 0.00 0.01	95% Cl (0.00-0.20) (0.00-0.60) (0.00-0.20) (0.00-0.03) 95% Cl (0.00-0.11) (0.00-0.20) (0.01-0.72) (0.00-0.20) (0.00-0.20) (0.00-0.20) (0.00-0.20) (0.00-0.20)
Veek 96 Week 48 Week 24 B	CTsStudyExDOLULAM TANGO ART PRODOLULAM TANGOCommon-effects modelHeterogeneity: $l^2 = 0\%$, $\tau^2 = 0$, $P=1.00$ StudyExSOLAR 3D DOLULAM SALSA TANGO ART PRORandom-effects modelHeterogeneity: $l^2 = 11\%$, $\tau^2 = <0.0001$, P StudyExDOLULAM ART PROStudyExDOLULAM ART PROCommon-effects modelCommon-effects model	vents 0 0 0 0 vents 1 0 1 0 0 2=0.34	Total 17 4 17 38 Total 50 17 5 4 17 5 4 17 93 93 Total 17 34	0 0.2 0.4 0.6 0.8 1 Proportion of individuals with VF	Proportion 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.02 0.00 0.02 0.00 0.00 0.00 0.00 0.00 0.01	95% Cl (0.00-0.20) (0.00-0.60) (0.00-0.20) (0.00-0.03) 95% Cl (0.00-0.11) (0.00-0.20) (0.01-0.72) (0.00-0.20) (0.00-0.20) (0.00-0.20) (0.00-0.20) (0.00-0.20) (0.00-0.20) (0.00-0.20)

real-world and RCT analyses based on reported VF outcomes at each time point

- Real-world: 3/186 (1.61%), 7/237 (2.95%), and 7/186 (3.76%), respectively
- RCT: 0/38 (0%), 2/93 (2.15%), and 0/34 (0%), respectively
- No treatment-emergent resistance mutations were reported
- Including all studies regardless of VF definition increased sample sizes without significantly impacting estimates

Table. Summary of VF Definitions and Outcomes for PWH With M184V/I RAMs Receiving DTG + 3TC in RWE Studies and RCTs

	Proportion with	M191)//Lidoptification	VF time point	VF	
Study (cohort)	M184V/I	method	week	n/N (%)	VF definition
RWE studies					
Hocqueloux	105/695 (15.11%)	RNA and proviral DNA genotypes (pooling both)	24	1/105 (0.95)	2 consecutive confirmed VL >50 c/r
2021			48	2/105 (1.90)	or 1 VL >200 c/mL
(Dat AIDS) ⁵			96	2/105 (1.90)	
Santoro 2021	36/533 (6.75%)	RNA and proviral DNA	24	2/36 (5.56)	2 consecutive confirmed VL >50 c/n
(LAMRES)⁰		genotypes	48	2/36 (5.56)	or 1 VL >200 c/mL
			96	3/36 (8.33)	
Borghetti 2021	48/669 (7.17%) ^a	Historical genotypes; does not	24	0/45	1 VL ≥1000 c/mL or 2 consecutive
(ODOACRE) ^{7,8}		specify RNA or proviral DNA	48	1/45 (2.22)	VL ≥50 c/mL
			96	2/45 (4.44)	
Galizzi 2020	47/174	Either RNA or proviral	24	—	2 consecutive confirmed VL >50 c/mL
(NR) ⁹	(27.01%) ^b	DNA genotypes at baseline	48	1/47 (2.13)	or 1 VL >50 c/mL followed by ART
		(before switch)	96	—	modification of 1 VL >1000 c/mL
Hidalgo-Tenorio	4/178 (2.25%)	Baseline RNA genotype	24		2 consecutive VL >50 c/mL
2019			48	1/4 (25.00)	
			96		
RCTs					
ART PRO ¹¹	17/41 (41.46%)	Proviral DNA genotype	24	0/17	VL ≥50 c/mL
			48	0/17	
			96	0/17	
SOLAR 3D ¹²	50/100 (50.00%)	Historical genotypes; does not	24	—	VL ≥50 c/mL
		specify RNA or proviral DNA	48	1/50 (2.00)	
			96		
TANGO ²	4/322 (1.24%)	Proviral DNA genotype	24	0/4 ^c	VL ≥50 c/mL
			48	0/4	
			96		
DOLULAM ¹³	17/27 (62.96%)	RNA and proviral DNA	24	0/17	VL >50 c/mL
		genotypes	48	0/17	
			96	0/17	
SALSA ³	5/192 (2.60%)	Proviral DNA genotype	24		VL ≥40 c/mL
			48	1/5 (20.00) ^d	
			96	_	

NR, not reported. aCohort reference reporting the proportion with VF for individuals with M184V/I was used for analysis (n=45 individuals with M184V/I).8 bAssumption: n=60 PWH with M184V/I were reported out of N=220 total PWH with available pre-switch genotype resistance data across 2 groups but not reported for DTG + 3TC specifically. Table n with M184V/I was calculated according to the proportion of PWH in the DTG + 3TC (n=174) vs other group (n=46). Assumption: Week 24 was not reported, but reports described no VF to Week 48. dVFs and discontinuations were not directly reported; study reported n (%) with VL <40 c/mL and TND, and here the participant had VL <40 c/mL with qualitative target detected (TD) outcome.

VF Estimates

- Random-effects models are associated with greater uncertainty vs common-effects models but can be used to estimate results for the wider population of interest based on the sample of studies used in the analysis • Common-effects (or fixed-effects) models assume that the included studies are the population of interest and can be more appropriate and informative when zero VF events are observed
- RWE common-effects models estimated the proportions (95% CI) of individuals with VF were 0.01 (0.00-0.03) at Week 24, 0.03 (0.01-0.06) at Week 48, and 0.04 (0.02-0.08) at Week 96; random-effects estimates are in Figure 2A
- RCT common-effects models estimated the proportions (95% CI) of individuals with VF were 0.01 (0.00-0.05) at Week 48; the random-effects estimate for this time point is in Figure 2B
- Common-effects models better represented Week 24 and Week 96 data consisting of zero observed events each (Figure 2B); random-effects models estimated Week 24 and Week 96 proportions (95% CI) were 0.00 (0.00-0.00)

Proportions were log-transformed, or arcsine-transformed if any studies reported zero events.

Conclusions

• Overall, pre-switch M184V/I prevalence was low in PWH in RWE studies

- Real-world studies of PWH with historical or archived M184V/I receiving DTG + 3TC identified low incidence of VF through 96 weeks and no reported cases of INSTI treatment-emergent mutations; these findings were consistent with results from RCTs
- Genotypic data at the time of VF were unavailable and the occurrence of resistance mutations to 3TC or DTG at failure could not be described
- This meta-analysis provides reassuring data on outcomes with DTG + 3TC in PWH with incomplete history or in cases where M184V/I was inadvertently missed

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