

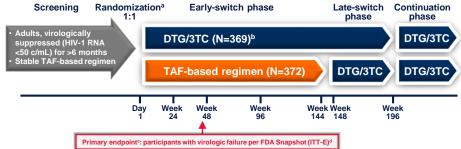
SWITCHING FROM A 3-DRUG TENOFOVIR ALAFENAMIDE (TAF)-BASED REGIMEN TO A 2-DRUG DOLUTEGRAVIR/LAMIVUDINE (2DR, DTG/3TC FDC) WAS NOT ASSOCIATED WITH A HIGHER FREQUENCY OF INTERMITTENT VIREMIA IN SUPPRESSED PATIENTS IN THE TANGO STUDY

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Introduction

- TANGO is a 200-week, phase III, randomized, open-label trial to evaluate efficacy and safety of switching from a TAF-based regimen to a 2DR of DTG/3TC in HIV-1-infected adults with HIV-1 RNA <50 c/mL and without prior virologic failure or historical NRTI or INSTI major resistance mutations at study entry (Figure 1)¹
- Switching to DTG/3TC was non-inferior to continuing a TAF-based regimen through Week 48 using a 4% non-inferiority margin for Snapshot virologic failure (HIV-1 RNA ≥50 c/mL)¹
- We assessed elevated viral loads (VLs) through 48 weeks of therapy overall, by study visit and in a subset of participants with archived resistance mutations





^aStratified by baseline third agent class (PI, INST), or NNRTI), ^b2 participants excluded who were randomized but not exposed to study drug. ^c4% noninferiority margin, ^dIncludes 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

Methods

- Elevated viral loads (EVLs. HIV-1 RNA ≥50 c/mL) for exposed participants (intention-to-treat– exposed [ITT-E population]) with at least one post-baseline, on-treatment plasma HIV-1 RNA were assessed in 2 major categories (Table 1): (1) with only VL ≥50 to <200 c/mL, or (2) at least one VL ≥200 c/mL
- · Each of these major categories was further divided into a single non-consecutive occurrence or >2 consecutive occurrence sub-categories
- A "blip" is defined here as a VL of 50 to <200 c/mL with adjacent values <50 c/mL
- Confirmed virologic withdrawal (CVW) criteria were defined as 2 consecutive on-treatment VL ≥50 c/mL with the second VL ≥200 c/mL
- Proviral DNA genotyping was conducted retrospectively on baseline whole blood samples using GenoSure Archive assay by Monogram Biosciences

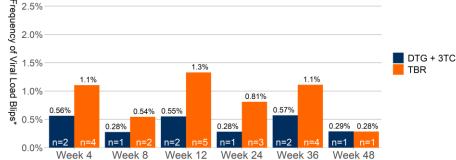
Results

- The proportion of participants with EVLs through 48 weeks of treatment (Table 1) was low and comparable across groups; most frequently observed VL rebounds were in Category 1a
- The proportion of Category 1a participants with blips by study visit is shown in Figure 2
- By Week 48, no participant in the DTG/3TC group met CVW criteria
- One participant on TAF-based regimen without any pre-existing resistance met CVW criteria · No blips occurred before CVW and no emergent resistance was observed at failure

Elevated VL categories for participants in the TT-E population	DTG/3TC FDC (N=369) n (%)	TAF-based regimen (N=372) n (%)
1. Participants with VLs between 50 to <200 c/mL and no VL ≥200 c/mL	11 (3%)	22 (6%)
 VLs between 50 to <200 c/mL with adjacent values <50 c/mL ("blips") 	9 (2%)	18 (5%)
1b. ≥2 consecutive VLs between 50 to <200 c/mL	2 (<1%)	4 (1%)
2. Participants with at least one VL ≥200 c/mL	3 (<1%)	3 (<1%)
2a. A single VL ≥200 c/mL and no 2 consecutive VL ≥50 c/mL	3 (<1%)	1 (<1%)
2b. ≥2 consecutive VLs ≥50 c/mL with at least one VL ≥200 c/mL	0	2ª (<1%)
Total (all categories)	14 (4%)	25 (7%)

Table 1 Summary of Participants With Flevated Viral Load Categories Through Week 48

Figure 2. Frequency of Viral Load Blips in Category 1a Participants by Visit Through Week 48



*Percentages were calculated from number of blins in Category 1a participants (Table 1) using post-baseline previously suppressed (<50 c/ml) pt visit Ns respectively for DTG/3TC and DTG + TDF/FTC at: Wk4 (N=355) and (N=362); Wk8 (N=361) and (N=367); Wk 12 (N=362) and (N=376); Wk 24 (N=355) and (N=370); Wk 36 (N=350) and (N=360); Wk 48 (N=348) and (N=351). Numbers on the bottom of each bar represent # of blips at given week visit. Individual participants can have had more than one blin

- The occurrences of viral blips at each visit by treatment group over 48 weeks were similar (Figure 2)
- The prevalence of archived, pre-existing M184V/I or K65R/E/N was very low (Table 2)
- No EVLs were observed among these participants

3.0%

- Through Week 48, the frequency of participants who experienced EVLs with archived major NRTI or INSTI resistance was very low in either treatment group
- In the DTG/3TC group, one participant with archived INSTI mutation mixture Q148Q/R had 2 consecutive VLs between 50 to <200 c/mL during the Week 4 window and a blip at Week 12; the participant withdrew consent and discontinued the study with VL <50 c/mL at the withdrawal visit

Table 2. Summary of Prevalen
Rebound by Pre-existing Resi

Baseline genotypic data available	DTG/3TC (N=322)		TAF-based regimen (N=321)	
Resistance class	Mutation prevalence ^a	Experienced elevated VL ^b	Mutation prevalence ^a	Experienced elevated VL ^b
Any major resistance ^c	81 (25%)	5 (2%)	88 (27%)	9 (3%)
No major resistance	241 (75%)	7 (2%)	233 (73%)	13 (4%)
Major NRTI class – any	25 (8%)	1 (<%)	17 (5%)	1 (<%)
M184V or M184I	4 (1%)	0	3 (<1%)	0
K65R or K65E or K65N	0	0	2 (<1%)	0
Any TAM ^d	9 (3%)	0	5 (2%)	0
Other	12 (4%)	1 (<1%)	7 (2%)	1 (<1%)
No M184V/I	318 (99%)	12 (4%)	318 (>99%)	22 (7%)
Major INSTI – any ^e	3 (<1%)	1 (<1%)	5 (2%)	0
Q148Q/R	2 (<1%)	1 (<1%)	1 (<1%)	0
Y143Y/H	0	0	2 (<1%)	0
Y143Y/C	1 (<1%)	0	0	0
R263R/K	0	0	2 (<1%)	0

wild-type and specific mutations.

Conclusions

- The occurrences of blips by visit were similar across treatment groups
- occurrences were similar
- The frequency of archived, pre-existing M184V/I or K65R/E/N was very low and did not appear to increase the risk of elevated viral loads in either treatment group, with no participants exhibiting intermittent viremia through Week 48
- In participants with or without pre-existing resistance, intermittent viremia was infrequent
- Switching from a 3-drug TAF-based regimen to a DTG/3TC 2DR was not associated with a higher frequency of intermittent viremia

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Reference: 1. van Wyk J, Ajana F, Bisshop F, et al. Switching to DTG/3TC fixed-dose combination (FDC) is non-inferior to continuing a TAF-based regimen in maintaining virologic suppression through 48 weeks (TANGO study). Presented at: 10th IAS Conference on HIV Science; July 21-24, 2019; Mexico City, Mexico. Slides WEAB0403LB.



ce of Archived Mutations and Participants With Viral stance at Baseline

abPercentage is based on N - number of participants with baseline proviral DNA genotypic data available from the proviral DNA resistance analysis population (PRAP). PRAP is based on the ITT-E population for whom there are: (1) available proviral DNA genotyping data; (2) at least one post-baseline on-treatment HIV-1 RNA viral load result available; and (3) reason for withdrawal is not protocol deviation. Includes major NRTI, NNRTI, PI, and INSTI resistance-associated mutations based on IAS 2019. dTAM - thymidine analogue mutation. Participants with archived major INSTIs all had mixtures of

• All other categories of VL ≥50 c/mL occurred infrequently in all groups, and the