

HIV Drug Resistance Mutations Associated with Virologic Failure in Women on Efavirenz-based ART following PROMISE Study

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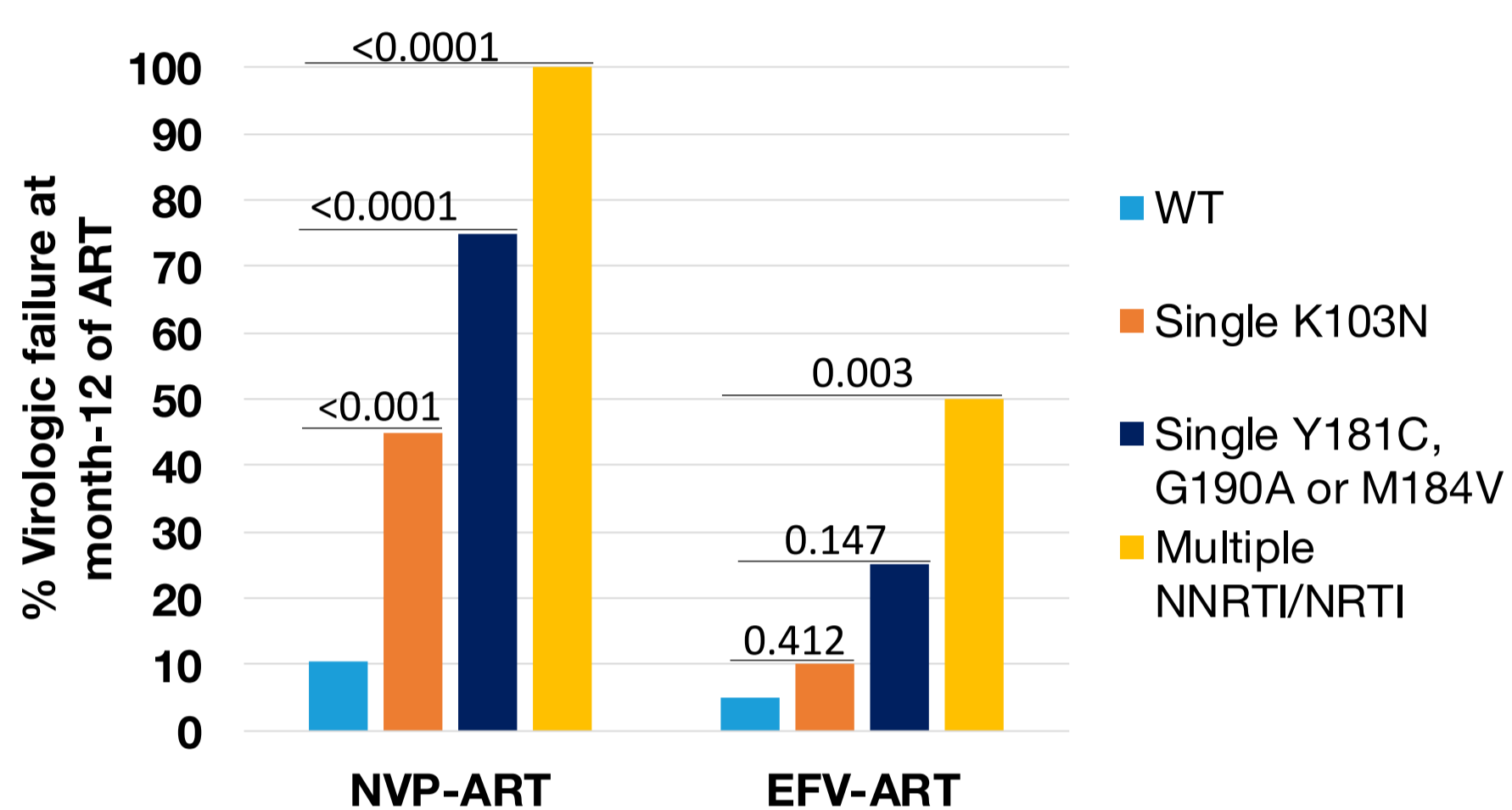
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Introduction

BACKGROUND

- Efavirenz-based antiretroviral therapy (EFV-ART) is a WHO-recommended 1st-line ART
- Pre-treatment HIV-drug resistance (PDR) to 1st-line non-nucleoside reverse transcriptase inhibitor (NNRTI) is increasing in low-resource communities due to transmitted and selected mutations, particularly in women, in part from treatment to prevent mother-to-child HIV transmission (PMTCT)
- Studies of Kenyans initiating NNRTI-ART from 2006-14:
 - PDR increased to 11% and to >20% in women 18-24yo
 - In 2014, drugs switched: NVP to EFV and ZDV to TDF
- Virologic failure (VF) varied by ART regimen (Figure 1):

Figure 1. VF at month-12 of NVP+ZDV- or EFV+TDF-ART by number of pre-ART DR mutations detected by OLA vs. wild type



OBJECTIVE OF THIS STUDY

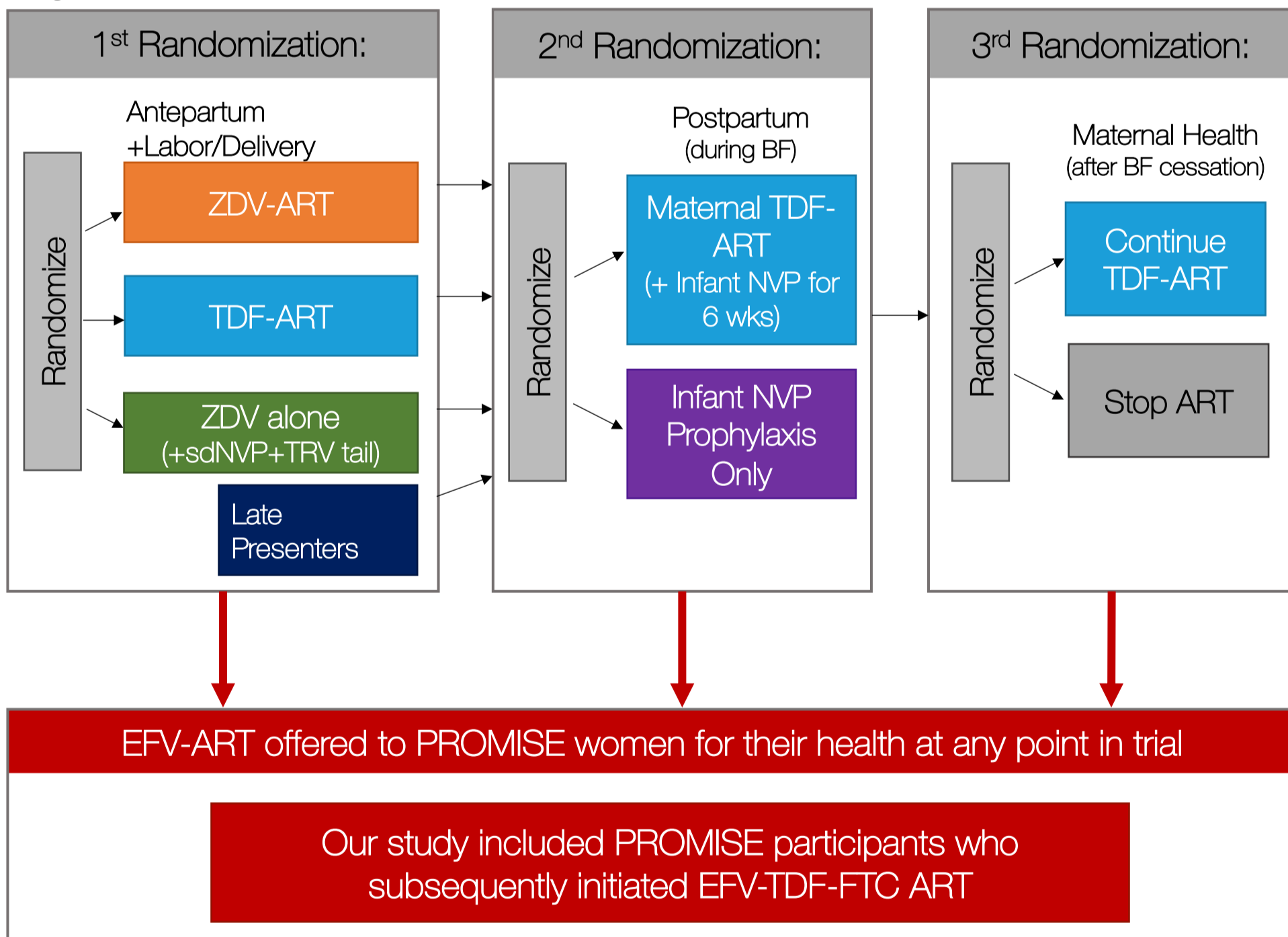
Assess if pre-EFV-ART DR in PROMISE Study participants who subsequently initiated EFV-ART is associated with VF

Methods

SUBJECTS

Participants in PROMISE study 1077BF (of strategies for PMTCT; Figure 2) who subsequently started EFV-ART for their own health; initiated EFV-ART at any point during the study

Figure 2. PROMISE randomization schema



INCLUSION CRITERIA

- PROMISE women who initiated EFV-ART and had:
 - Enrollment plasma HIV RNA ≥ 400 c/mL available
 - Plasma available just prior to EFV-ART initiation
 - Plasma HIV RNA known at month-6 and -12 of EFV-ART

METHODS & ANALYSES

- Genotyped PR & RT by consensus sequencing (CS)
- Illumina sequencing with "Primer ID" technology for minority variants
- Phylogenetic and bioinformatic analyses used for quality assurance
- Virologic failure was defined as HIV RNA > 400 c/mL
- NRTI- & NNRTI-associated mutations with Stanford Database score > 10 were analyzed (Table 1)
- Rates of EFV-ART VF compared by pre-EFV genotype and PROMISE antepartum (AP) treatment arm (Fisher's Exact test)

Table 1. Mutations included as DRMs in the analyses

NRTI:	Mutations
NRTI:	M41L, K65R*, D67N, K70_, L74I*, V75I, M184_, T215_, K219_
NNRTI:	Mutations
NNRTI:	A98G, L100I, K101_, K103_, V106_, V108I, Y181C, Y188_, G190_, H221Y, P225H, M230L*, K238T

*DRMs not found in any of the women in this study

Results

Figure 3. Pre-EFV DR and VF by month-6 or -12 of EFV-ART by clinical site and overall

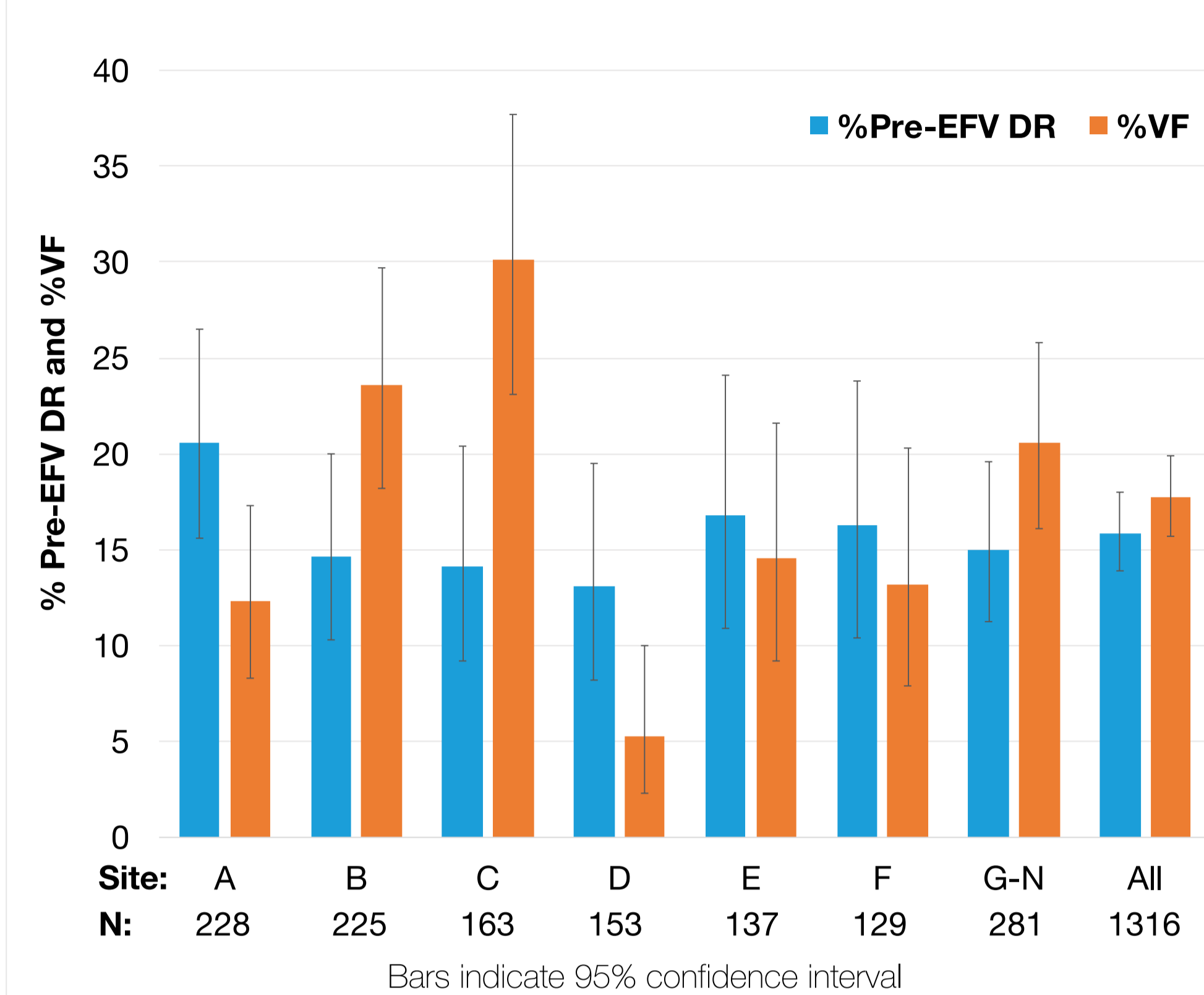


Figure 4. VF by pre-EFV genotype across clinical sites and overall

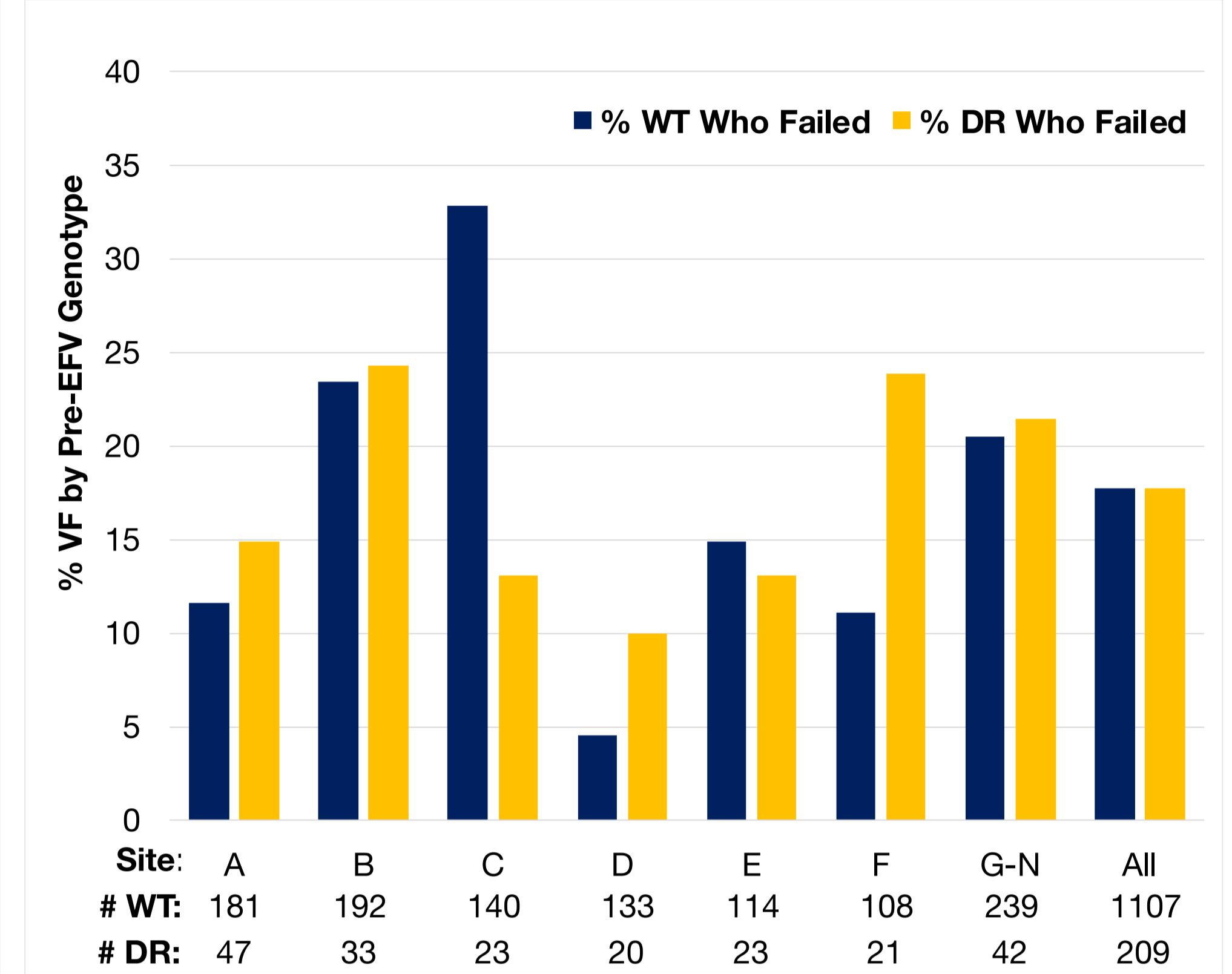


Table 3. VF by pre-EFV genotype

Pre-EFV Genotype	Total # Women	# (%) VF	p-value
WT	1,107	196 (17.7)	reference
1 NRTI only	13	0 (0)	0.2362
K65R only	0	0 (--)	N/A
K70R only	4	0 (0)	1.0000
M184V only	1	0 (0)	1.0000
≥ 2 NRTI only	0	0 (--)	N/A
1 NNRTI only	169	26 (15.3)	0.5897
K103N only	97	18 (18.6)	0.8918
Y181C only	8	1 (12.5)	1.0000
G190A only	5	0 (0)	1.0000
≥ 2 NNRTI only	19	4 (21.1)	0.7674
NRTI & NNRTI	8	7 (87.5)	<0.0001

N/A = not analyzed

Table 5. VF within previous antepartum treatment arm by pre-EFV genotype

AP Treatment Arm	Pre-EFV Genotype	Total # (%) Women	# (%) VF on EFV-ART	p-value
ZDV-ART (ZDV-3TC/LPV-RTV)	Total	582	119 (20.5)	N/A
	WT	496 (85.4)	104 (21.0)	Reference
	Any DRM	85 (14.6)	15 (13.8)	0.5618
	NRTI DRM only	5 (0.9)	0 (0.0)	0.5888
	NNRTI DRM(s) only	75 (12.9)	11 (14.7)	0.2786
NRTI & NNRTI	5 (0.9)	4 (80.0)	0.0086**	
TDF-ART (TDF-FTC/LPV-RTV)	Total	181	27 (14.8)	N/A
	WT	149 (81.9)	16 (10.7)	Reference
	Any DRM	33 (18.1)	11 (31.3)	0.0024**
	NRTI DRM only	0 (0.0)	0 (--)	1.0000
	NNRTI DRM(s) only	31 (17.1)	10 (32.3)	0.0043**
NRTI & NNRTI	1 (0.6)	1 (100)	0.1133	
ZDV Alone (+sdNVP+TRV tail)	Total	553	87 (15.7)	N/A
	WT	461 (83.3)	76 (16.5)	Reference
	Any DRM	92 (16.6)	11 (10.0)	0.3468
	NRTI DRM only	8 (1.4)	0 (0.0)	0.3649
	NNRTI DRM(s) only	82 (14.8)	9 (11.0)	0.2493
NRTI & NNRTI	2 (0.4)	2 (100)	0.0281*	

*p < 0.05; **p < 0.01; N/A = not analyzed

Table 4. VF by previous PMTCT treatment arm: ZDV alone vs ART

Antepartum Treatment Arm	Total # Women	# (%) VF	p-value
ZDV Alone	553	87 (15.7)	Reference
ART (TDF or ZDV)	763	146 (19.1)	0.1941

Summary & Conclusions

SUMMARY

- Pre-EFV-ART DR was detected in 209/1,316 women (15.9%, 95% CI: 13.9-18%)
- VF was detected in 233/1,316 women (17.7%, 95% CI: 15.7-19.9%)
- Pre-EFV DR prevalence and rates of VF varied across the 14 clinical sites (IQR = 11.2-18.8% and IQR = 12.4-28.5%, respectively)
- Rates of VF by 6-12 months of EFV-ART (Table 2) did not differ by pre-EFV genotype by CS (17.7% vs. 17.7%, p=1.0)
- Single or multiple NRTI or NNRTI DRMs were not associated with VF compared to no DRM (p=0.40 and p=0.77, respectively)
- Pre-EFV DR to multiple drug classes (≥ 1 NRTI & ≥ 1 NNRTI DRMs) was associated with increased VF (p<0.0001)
- Past randomization to ART vs ZDV in PROMISE did not significantly impact rates of VF (p=0.1941)
- Any Pre-EFV DR found after past randomization to TDF-ART was associated with VF whereas only multi-class PDR was associated with VF for ZDV-ART
- Illumina sequencing analyses of minority variants in women with wild-type pre-EFV genotype and VF is ongoing

CONCLUSIONS

- Pre-EFV-ART drug resistance data from PROMISE trial aligns with previous observations in Kenya:
 - Single NRTI or NNRTI DRMs do not significantly impact rates of VF to EFV-ART
 - EFV-ART does not appear to suppress replication of multi-class drug-resistant HIV
- This study suggests:
 - EFV-ART is a more potent regimen compared to NVP-ART
 - Detection of multi-class PDR could extend the effective use of EFV-ART in resource limited settings

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