Drug resistance interpretation: PR HIVDB 9.5.1 (2023-11-05)

PI Major Mutations: None PLAccessory Mutations:

PR Other Mutations: L10I are . 113V are . E35D are . M36I are . R41K are . 164V are

Protease Inhibitors

atazanavir/r (ATV/r) Susceptible darunavir/r (DRV/r) Susceptible lopinavir/r (LPV/r) Susceptible

## Other

. L10(V are polymorphic, PI-selected accessory mutations that increase the replication of viruses with other PI-resistance mutations.

Mutation scoring: PR

NRTI Mutations:

No drug resistance mutations were found for PI.

Drug resistance interpretation: RT

K65R on 500 568G on K70KT 1 mm, 6 200 M184V on K219KQ K815 Q on

NNRTI Mutations: K103N --- G190A ---

RT Other Mutations: P4PS 128 - 139KN 128 - 139KN 128 - 1282C - 139KN 128 - 1282C -

Nucleoside Reverse Transcriptase Inhibitors

Non-nucleoside Reverse Transcriptase Inhibitors abacavir (ABC) High-Level Resistance doravirine (DOR) Susceptible High-Level Resistance zidovudine (AZT) Susceptible efavirenz (EFV) emtricitabine (FTC) High-Level Resistance etravirine (ETR) Potential Low-Level Resistance lamivudine (3TC) High-Level Resistance nevirapine (NVP) High-Level Resistance tenofovir (TDF) High-Level Resistance rilpivirine (RPV) Low-Level Resistance

## RT comments

# NRTI

- K65R confers intermediate reductions in susceptibility to TDF, ABC, and 3TC/FTC. It increases AZT susceptibility. In NRTI-naive patients with K65R, TDF+3TC+DTG is usually highly effective and more effective than AZT/3TC/DTG. However, in patients receiving TDF+3TC+DTG.
- 5686 is a polymorphic mutation that is often selected in combination with K63R. It partially restores the replication defect associated with K65R.
- K70/E/Q/N/T/S/G cause low-leve resistance to ABC and TDF.
- M184V/I cause high-level in vitro resistance to 3TC and Iow/intermediate resistance to ABC (3-fold reduced susceptibility). M184V/I are not contraindications to continued treatment with 3TC or FTC because they increase susceptibility to AZT and TDF and are associated with clinically significant reductions in HIV-1 replication.
- K219E/Q/N/R are accessory TAMS that usually occur in combination with multiple other TAMs.

NNRTI

Other

- . K103N is a non-polymorphic mutation that confers high-level reductions in NVP and EPV susceptibility. It is the most commonly transmitted DRM.
- 6190A is a non-polymorphic mutation that causes high-level resistance to NVP and intermediate resistance to EFV. It does not significantly reduce susceptibility to RPV, ETR, or DOR.

- V179D/E are somewhat polymorphic accessory NNRTI-selected mutation. In combination with other NNRTI DRMs, they appear to contribute low-levels of reduced susceptibility. V179F is a non-polymorphic mutation selected in combination with Y181C in persons receiving ETR. Alone it has little effect on NNRTI. susceptibility, however in combination with Y181C it is associated with high-level reductions in ETR and RPV susceptibility. V179L is a rare non-polymorphic mutation disted as a RPV-associated resistance mutation by the FDA package insert. Its effects on NNRTI susceptibility have not been well studied. V179A is an unusual mutation at this position.
- . This virus is predicted to have low-level reduced susceptibility to RPV. The use of the combination of CAB/RPV should be considered to be relatively contraindicated.

Mutation scoring: RT

HIVDB 9.5.1 (2023-11-05)

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Drug resistance muti	ation score:	s of NRTI:		Download CSV		
Rule	ABC =	AZT ≑	FTC ÷	зтс ≑	TDF 0	
K65R	45	-10	30	30	50	
K70KT	15	0	10	10	15	
M184V	15	-10	60	60	-10	
K219KQ	5	10	0	0	5	
K70KT + M184V	0	0	0	0	10	
K65R + S68G	0	0	0	0	5	
Total	80	-10	100	100	75	

Drug resistance mutation scores of NNRTI:

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Rule	DOR ÷	EFV ≑	ETR ≑	NVP ≑	RPV ≑
K103N	0	60	0	60	0
G190A	0	45	10	60	15
Total	0	105	10	120	15