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

PI Major Mutations:

None

PI Accessory Mutations: None

115V mm - K20R mm - E35D mm - M36I mm - R41K mm - R57K mm - 162V mm - L63P mm - H69K mm - V75I mm - L89M mm - R57K m PR Other Mutations:

Protease Inhibitors

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

PR comments

Other

. K20R is a highly polymorphic PI-selected accessory mutation that increases replication fitness in viruses with PI-resistance mutations.

Mutation scoring: PR

No drug resistance mutations were found for PI.

Drug resistance interpretation: RT

K219KE 1-40%, 6, 37% NRTI Mutations: NNRTI Mutations: K103N ***....

RT Other Mutations:

Nucleoside Reverse Transcriptase Inhibitors Non-nucleoside Reverse Transcriptase Inhibitors abacavir (ABC) Susceptible doravirine (DOR) Potential Low-Level Resistance

Susceptible zidovudine (AZT) efavirenz (EFV) High-Level Resistance emtricitabine (FTC) etravirine (ETR) Susceptible Susceptible lamivudine (3TC) Susceptible High-Level Resistance nevirapine (NVP) tenofovir (TDF) Susceptible rilpivirine (RPV) Susceptible

RT comments

NRTI

. K219E/Q/N/R are accessory TAMS that usually occur in combination with multiple other TAMs.

NNRTI

. K103N is a non-polymorphic mutation that confers high-level reductions in NVP and EFV susceptibility. It is the most commonly transmitted DRM.

. V179I is a polymorphic mutation that is frequently selected in persons receiving ETR and RPV. However, it has little, if any, direct effect on NNRTI susceptibility.

Mutation scoring: RT

HIVDB 9.5.1 (2023-11-05)

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rug resistan	ce mutation	Download	CSV ~		
Rule	ABC ÷	AZT ≑	FTC ÷	3TC ≑	TDF ÷
K219KE	5	10	0	0	5

Drug resista	nce mutation	Download CSV			
Rule	DOR ÷	EFV ‡	ETR ÷	NVP ÷	RPV ≑
K103N	0	60	0	60	0