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

PI Major Mutations:

None

None

PI Accessory Mutations:

PR Other Mutations: L100 cm. + L13V cm. + E35D cm. + M36I cm. + R41K cm. + K45R cm. + L63T cm. + I64L - L00 cm. + H69Q cm. + L89M cm.

Protease Inhibitors

atazanavir/r (ATV/r) Susceptible darunavir/r (DRV/r) Susceptible fosamprenavir/r (FPV/r) Susceptible indinavir/r (IDV/r) Susceptible lopinavir/r (LPV/r) Susceptible nelfinavir (NFV) Susceptible saquinavir/r (SQV/r) Susceptible tipranavir/r (TPV/r) Susceptible

PR comments

Other

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

No drug resistance mutations were found for PI.

Drug resistance interpretation: RT

Mutation scoring: PR

NRTI Mutations:

None

K103N - V108I -

NNRTI Mutations: RT Other Mutations:

K20R -- V35IM -- W179I -- T200A -- V207A -- V207

Nucleoside Reverse Transcriptase Inhibitors abacavir (ABC) Susceptible zidovudine (AZT) Susceptible stavudine (D4T) Susceptible didanosine (DDI) Susceptible emtricitabine (FTC) Susceptible lamivudine (3TC) Susceptible tenofovir (TDF) Susceptible

Non-nucleoside Reverse Transcriptase Inhibitors

doravirine (DOR) Potential Low-Level Resistance efavirenz (EFV) High-Level Resistance etravirine (ETR) Susceptible nevirapine (NVP) High-Level Resistance rilpivirine (RPV) Susceptible

RT comments

NNRTI

- K103N is a non-polymorphic mutation that confers high-level reductions in NVP and EFV susceptibility. It is the most commonly transmitted DRM.
- V108I is a relatively non-polymorphic accessory mutation selected in vitro and/or in vivo with each of the NNRTIs. It appears to contribute to reduced susceptibility to most NNRTIs only in combination with other NNRTI-resistance mutations.

Other

. 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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No drug resistance mutations were found for NRTI.

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Orug resistance mutation scores of NNRT1:			Download CSV		•	
Rule	DOR ÷	EFV ≑	ETR ≑	NVP ≑	RPV	÷
V108I	10	10	0	15	0	
K103N	0	60	0	60	0	
Total	10	70	0	75	0	