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

PI Major Mutations: None

PI Accessory Mutations: None

PR Other Mutations: 113V == . E35D == . M36I == . R41K == . K45R == . K57K == . L63T == . 164M == . H69K == . L89M =

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 Susceptible nelfinavir (NFV) saquinavir/r (SQV/r) Susceptible tipranavir/r (TPV/r) Susceptible

HIVDB 9.5.1 (2023-11-05) Mutation scoring: PR

No drug resistance mutations were found for PI.

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

NRTI Mutations:

K103N *** NNRTI Mutations:

RT Other Mutations:

Nucleoside Reverse Transcriptase Inhibitors

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

Non-nucleoside Reverse Transcriptase Inhibitors

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

RT comments

NNRTI

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

Other

Mutation scoring: RT

. 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.

No drug resistance mutations were found for NRTI.

Drug resistance mutation scores of NNRTI:

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Rule	DOR ÷	EFV ÷	ETR ÷	NVP ÷	RPV :
K103N	0	60	0	60	0

Drug resistance interpretation: IN

HIVDB 9.5.1 (2023-11-05)

HIVDB 9.5.1 (2023-11-05)

INSTI Major Mutations: None INSTI Accessory Mutations: None

IN Other Mutations:

Integrase Strand Transfer Inhibitors

bictegravir (BIC) Susceptible Susceptible cabotegravir (CAB) dolutegravir (DTG) Susceptible elvitegravir (EVG) Susceptible raltegravir (RAL) Susceptible

HIVDB 9.5.1 (2023-11-05)

Mutation scoring: IN