

Drug resistance interpretation: PR

HIVDB 9.5.1 (2023-11-05)

PI Major Mutations:

None

PI Accessory Mutations:

None

PR Other Mutations:

L10V100%
seen(3,302) • I13N110%
seen(3,759) • V49L100%
seen(3,302) • E35D100%
seen(3,302) • P39PS914%
seen(3,302) • R41K99%
seen(3,338) • D60E99%
seen(3,380) • L63P99%
seen(3,271) • I64V100%
seen(3,271)

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

- L10I/V are polymorphic, PI-selected accessory mutations that increase the replication of viruses with other PI-resistance mutations.

Mutation scoring: PR

HIVDB 9.5.1 (2023-11-05)

No drug resistance mutations were found for PI.

Drug resistance interpretation: RT

HIVDB 9.5.1 (2023-11-05)

NRTI Mutations:

None

NNRTI Mutations:

V179VD114%
seen(4,552)

RT Other Mutations:

V35T100%
seen(3,333) • K49R99%
seen(3,302) • I50V114%
seen(2,217) • V60I100%
seen(3,332) • K103KR914%
seen(2,307) • K122E99%
seen(3,335) • D123N99%
seen(3,335) • E169D99%
seen(4,367) • D177E100%
seen(4,332) • I178IL1114%
seen(4,303) • V189I99%
seen(4,112) • T200TI114%
seen(3,333) • I202V100%
seen(3,381) • Q207E100%
seen(3,311) • R211K100%
seen(3,314) • F214L99%
seen(3,371) • V245Q99%
seen(3,382)

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)

Susceptible

efavirenz (EFV)

Intermediate Resistance

etravirine (ETR)

Potential Low-Level Resistance

nevirapine (NVP)

Intermediate Resistance

rilpivirine (RPV)

Low-Level Resistance

RT comments

NNRTI

- V179D/E are somewhat polymorphic accessory NNRTI-selected mutation. In combination with other NNRTI DRMs, they appear to contribute low-levels of reduced susceptibility to each of the NNRTIs. In particular, the combinations of K103R/V179D and V106I/V179D act synergistically to reduce NVP and EFV susceptibility.

Other

- K103R is a polymorphic mutation that alone has no effect on NNRTI susceptibility. However, in combination with V179D, it reduces NVP and EFV susceptibility about 15-fold.
- 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)

No drug resistance mutations were found for NRTI.

Drug resistance mutation scores of NNRTI:

Download CSV

| Rule | DOR ⚖ | EFV ⚖ | ETR ⚖ | NVP ⚖ | RPV ⚖ |
|-----------------|-------|-------|-------|-------|-------|
| K103KR + V179VD | 0 | 20 | 0 | 20 | 15 |
| V179VD | 0 | 10 | 10 | 10 | 10 |
| Total | 0 | 30 | 10 | 30 | 25 |