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

PI Major Mutations: None

None PI Accessory Mutations:

113V 88% - K14R 83% - M36I 93% - R41K 99% - L63Q 54% - I64ISV 159%, 5: 24%, 1: 15% - H69HK H: 85%, K: 11% cov-86.821 504 cov-21.500 PR Other Mutations:

Protease Inhibitors

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

Mutation scoring: PR

No drug resistance mutations were found for Pl.

Drug resistance interpretation: RT

None

None NNRTI Mutations:

RT Other Mutations:

A272P 0476 - K277R 0576 - L282C 0776 - P294T 0776 - V317A 0776 - G335D 0576 - P345Q 0576 - F346Y 0576 - F346Y

Non-nucleoside Reverse Transcriptase Inhibitors

Nucleoside Reverse Transcriptase Inhibitors

zidovudine (AZT) emtricitabine (FTC) Susceptible susceptible susceptible susceptible susceptible susceptible nevirapine (NVP)		•	
emtricitabine (FTC) Susceptible etravirine (ETR) nevirapine (NVP)	abacavir (ABC)	Susceptible	doravirine (DOR)
lamivudine (3TC) Susceptible nevirapine (NVP)	zidovudine (AZT)	Susceptible	efavirenz (EFV)
	emtricitabine (FTC)	Susceptible	etravirine (ETR)
tenofovir (TDF) Susceptible rilpivirine (RPV)	lamivudine (3TC)	Susceptible	nevirapine (NVP)
	tenofovir (TDF)	Susceptible	rilpivirine (RPV)

RT comments

Mutation scoring: RT

NRTI Mutations:

Other

L100I is a non-polymorphic mutation that usually occurs in combination with K103N. In this setting it confers high-level resistance to ETR and DOR. L100V is a rare mutations that likely has effects similar to L100I. L100S is a highly unusual mutation at this position.

No drug resistance mutations were found for NRTI.

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No drug resistance mutations were found for NNRTI.