

PI Major Mutations:None

PI Accessory Mutations:None

PR Other Mutations:T12P100%1000/1,280 • I13V10%1000/1,280 • K14R10%1000/1,280 • Q18H10%1000/1,280 • K20R10%1000/1,280 • E33D10%1000/1,280 • M36I10%1000/1,280 • N37D10%1000/1,280 • R41K100%1000/1,280 • K45R10%1000/1,280 • R57K10%1000/1,280 • L63P100%1000/1,280 • I64L10%1000/1,280 • H69K10%1000/1,280 • L89M100%1000/1,280

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

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

No drug resistance mutations were found for PI.

NRTI Mutations:[T69TD](#)100%1000/1,280 T: 40%, D: 17% • [V75VM](#)100%1000/1,280 N: 40%, V: 10% • [M184V](#)100%1000/1,280

NNRTI Mutations:None

RT Other Mutations:V35T100%1000/1,280 • T39A100%1000/1,280 • S48Q10%1000/1,280 • V60I100%1000/1,280 • K122E100%1000/1,280 • D123NS0-100%1000/1,280 N: 100%, S: 10% • I135T100%1000/1,280 • S162A100%1000/1,280 • K173S10%1000/1,280 • Q174K40%1000/1,280 • V179I10%1000/1,280 • T200A100%1000/1,280 • Q207A100%1000/1,280 • R211S10%1000/1,280 • V243KQ0-100%1000/1,280 K: 40%, Q: 17% • D250E100%1000/1,280 • S519N100%1000/1,280 • Q520H100%1000/1,280 • Q524EK0-100%1000/1,280 N: 100%, K: 10% • E529D100%1000/1,280 • K530R100%1000/1,280 • A534S100%1000/1,280 • V548V0-100%1000/1,280 V: 40%, L: 10% • A554S100%1000/1,280

Nucleoside Reverse Transcriptase Inhibitors		Non-nucleoside Reverse Transcriptase Inhibitors	
abacavir (ABC)	Low-Level Resistance	doravirine (DOR)	Susceptible
zidovudine (AZT)	Susceptible	efavirenz (EFV)	Susceptible
stavudine (D4T)	Intermediate Resistance	etravirine (ETR)	Susceptible
didanosine (DDI)	Intermediate Resistance	nevirapine (NVP)	Susceptible
emtricitabine (FTC)	High-Level Resistance	rilpivirine (RPV)	Susceptible
lamivudine (3TC)	High-Level Resistance		
tenofovir (TDF)	Susceptible		

RT comments

NRTI

- T69D is a nonpolymorphic mutation selected by early NRTIs that does not appear to reduce AZT, ABC, or TDF susceptibility.
- V75T(M/A/S are nonpolymorphic accessory NRTI-selected mutations. They appear to have minimal phenotypic effects on AZT, ABC, and TDF.
- M184V(I cause high-level in vitro resistance to 3TC and FTC and low/intermediate resistance to ABC (3-fold reduced susceptibility). M184V(I are not contraindications to continued treatment with 3TC or FTC because they increase susceptibility to AZT and TDF and are associated with clinically significant reductions in HIV-1 replication.

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.

Drug resistance mutation scores of NRTI:

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Rule	ABC ÷	AZT ÷	D4T ÷	DDI ÷	FTC ÷	3TC ÷	TDF ÷
<a href="#">M184V</a>	15	-10	-10	10	60	60	-10
<a href="#">V75VM</a>	0	10	30	15	0	0	0
<a href="#">T69TD</a>	0	0	10	30	0	0	0
Total	15	0	30	55	60	60	-10

No drug resistance mutations were found for NNRTI.

INSTI Major Mutations: None

INSTI Accessory Mutations: None

IN Other Mutations: K7R 100%  
seen:147 • K14KR 0.40%  
seen:240 • S17N 100%  
seen:201 • L28I 100%  
seen:170 • P30A 100%  
seen:196 • I60V 100%  
seen:82 • I72V 100%  
seen:117 • T112I 1.14%  
seen:170 • I113V 100%  
seen:170 • T124A 100%  
seen:114 • T125A 100%  
seen:114 • K136Q 100%  
seen:90 • D167E 99%  
seen:109 • V201I 99%  
seen:124 • T218T 1.14%  
seen:97 • L234I 100%  
seen:106 • M275V 99%  
seen:126 • S283G 100%  
seen:171 • D286N 100%  
seen:80

Integrase Strand Transfer Inhibitors

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

No drug resistance mutations were found for INSTI.