

PI Major Mutations:None

PI Accessory Mutations:[L33LF](#) (L: 4.7%
H: 0.0%)

PR Other Mutations:[I13V](#) (I: 0.0%
H: 12.1%) • [K20KER](#) (K: 1.0%
H: 1.0%) • [E35M](#) (E: 0.0%
H: 0.0%) • [M36I](#) (M: 0.0%
H: 0.0%) • [N37D](#) (N: 0.0%
H: 0.0%) • [R41RK](#) (R: 0.0%
H: 0.0%) • [D60E](#) (D: 0.0%
H: 0.0%) • [H69K](#) (H: 0.0%
H: 0.0%) • [I72M](#) (I: 0.0%
H: 0.0%) • [L89IM](#) (L: 0.0%
H: 0.0%)

Protease Inhibitors	
atazanavir/r (ATV/r)	Susceptible
darunavir/r (DRV/r)	Susceptible
fosamprenavir/r (FPV/r)	Potential Low-Level Resistance
indinavir/r (IDV/r)	Susceptible
lopinavir/r (LPV/r)	Susceptible
nelfinavir (NFV)	Potential Low-Level Resistance
saquinavir/r (SQV/r)	Susceptible
tipranavir/r (TPV/r)	Potential Low-Level Resistance

PR comments

Accessory

- [L33F](#) is a relatively non-polymorphic accessory mutation selected by each of the PIs. In combination with other PI-resistance mutations, it is associated with reduced susceptibility to LPV, ATV, and DRV.

Other

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

Drug resistance mutation scores of PI:

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Rule	ATV/r	DRV/r	FPV/r	IDV/r	LPV/r	NFV	SQV/r	TPV/r
L33LF	5	5	10	5	5	10	5	10

NRTI Mutations:[D67N](#) (D: 0.0%
H: 0.0%) • [K70R](#) (K: 0.0%
H: 0.0%) • [M184V](#) (M: 0.0%
H: 0.0%) • [K219Q](#) (K: 0.0%
H: 0.0%)

NNRTI Mutations:None

RT Other Mutations:[ISIV](#) (I: 0.0%
H: 0.0%) • [V35T](#) (V: 0.0%
H: 0.0%) • [T39L](#) (T: 0.0%
H: 0.0%) • [I50V](#) (I: 0.0%
H: 0.0%) • [V60I](#) (V: 0.0%
H: 0.0%) • [T69N](#) (T: 0.0%
H: 0.0%) • [K122E](#) (K: 0.0%
H: 0.0%) • [D123S](#) (D: 0.0%
H: 0.0%) • [I135T](#) (I: 0.0%
H: 0.0%) • [K173S](#) (K: 0.0%
H: 0.0%) • [Q174N](#) (Q: 0.0%
H: 0.0%) • [D177E](#) (D: 0.0%
H: 0.0%) • [I178IM](#) (I: 0.0%
H: 0.0%) • [I195IL](#) (I: 0.0%
H: 0.0%) • [G196E](#) (G: 0.0%
H: 0.0%) • [Q207A](#) (Q: 0.0%
H: 0.0%) • [R211A](#) (R: 0.0%
H: 0.0%) • [F214L](#) (F: 0.0%
H: 0.0%) • [Q209L](#) (Q: 0.0%
H: 0.0%) • [D911S](#) (D: 0.0%
H: 0.0%) • [K512S](#) (K: 0.0%
H: 0.0%) • [L517I](#) (L: 0.0%
H: 0.0%) • [S519N](#) (S: 0.0%
H: 0.0%) • [Q524K](#) (Q: 0.0%
H: 0.0%) • [I526IT](#) (I: 0.0%
H: 0.0%) • [K527EG](#) (K: 0.0%
H: 0.0%) • [E529D](#) (E: 0.0%
H: 0.0%) • [A554S](#) (A: 0.0%
H: 0.0%)

Nucleoside Reverse Transcriptase Inhibitors		Non-nucleoside Reverse Transcriptase Inhibitors	
abacavir (ABC)	Intermediate Resistance	doravirine (DOR)	Susceptible
zidovudine (AZT)	High-Level Resistance	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)	Low-Level Resistance		

RT comments

NRTI

- [D67N](#) is a non-polymorphic TAM associated with low-level resistance to AZT.
- [K70R](#) is a TAM that confers intermediate resistance to AZT and contributes to reduced ABC and TDF susceptibility in combination with other TAMs.
- [M184V](#)/I cause high-level in vitro resistance to 3TC 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.
- [K219E](#)/[Q](#)/[N](#)/[R](#) are accessory TAMs that usually occur in combination with multiple other TAMs.

Other

- [T69N](#)([S](#)/[A](#)/[I](#))/[E](#) are relatively non-polymorphic mutations weakly selected in persons receiving NRTIs. They may minimally contribute reduced AZT susceptibility.

Drug resistance mutation scores of NRTI:

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Rule	ABC	AZT	D4T	DDI	FTC	3TC	TDF
D67N	5	15	15	5	0	0	5
D67N + K70R + M184V + K219Q	10	0	0	0	0	0	0
D67N + K70R + K219Q	10	15	10	10	10	10	10
K70R	5	30	15	10	0	0	5
M184V	15	-10	-10	10	60	60	-10
K219Q	5	10	10	5	0	0	5
Total	50	60	40	40	70	70	15

No drug resistance mutations were found for NNRTI.

INSTI Major Mutations:

None

INSTI Accessory Mutations:

None

IN Other Mutations:

E11D 100%
seen:207 • K14R 99%
seen:224 • N27NS N: 40%, S: 10%
seen:275 • V31I 99%
seen:280 • L101I 100%
seen:282 • T112V 100%
seen:285 • I113V 99%
seen:285 • T124A 100%
seen:284 • T125A 100%
seen:284 • G134D 99%
seen:283 • K136Q 99%
seen:283 • G163V 100%
seen:282 • V201I 100%
seen:279 • L234I 99%
seen:287 • **D256DN** D: 99%, N: 21%
seen:273 • S283G 100%
seen:284 • R284G 99%
seen:284 • D286N 99%
seen:278

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.