

NRTI Mutations:

NNRTI Mutations:

RT Other Mutations:

K65R

S68G

V75A

L100I

K103N

F227L

P1K

S3R

P4R

I5V

K22N

Q23P

P25S

V35T

T39K

K43E

K49R

P55F

N57Y

R78T

Q85R

E89Q

G112S

D121Y

K122E

D123E

T128S

N137Y

T165L

P170H

D177E

I178M

M184E

G190R

T200R

K201N

I202T

E203Q

E204Q

Q207E

L210

R211T

F214L

T216S

P217S

K219

E224E_T

P226I

L228V

W229R

K238Q

V241A

Q242P

P243L

I244S

V245

K249

D250G

S251R

W252L

T253D

V254C

N255

D256L

I257N

Q258I

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

RT comments

NRTI

- K65R confers intermediate reductions in susceptibility to TDF, ABC, and 3TC/FTC. It increases AZT susceptibility. In NRTI-experienced, INSTI-naïve patients with K65R, TDF+3TC+DTG is usually highly effective and more effective than AZT/3TC/DTG. However, in patients receiving TDF+3TC+DTG, there is a risk of emergent DTG resistance that does not arise in NRTI-naïve patients receiving TDF+3TC+DTG.
- S68G is a polymorphic mutation that is often selected in combination with K65R. It partially restores the replication defect associated with K65R.
- V75T/M/A/S are nonpolymorphic accessory NRTI-selected mutations. They appear to have minimal phenotypic effects on AZT, ABC, and TDF.

NNRTI

- L100I is a non-polymorphic mutation that usually occurs in combination with K103N. In this setting it confers high-level resistance to NVP, EFV, and RPV and intermediate resistance to ETR and DOR.
- K103N is a non-polymorphic mutation that confers high-level reductions in NVP and EFV susceptibility. It is the most commonly transmitted DRM.
- F227L is a non-polymorphic mutation that usually occurs in combination with V106A. It is selected in vivo and in vitro with both NVP and DOR. In this context it is associated with high-level reductions in NVP and DOR susceptibility and intermediate reductions in EFV susceptibility. F227I/V are extremely rare mutations that have been selected in vitro by DOR.

Other

- 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. M184E is a highly unusual mutation at this position.
- K238T/N are uncommon non-polymorphic mutations selected in persons receiving NVP and EFV usually in combination with K103N. Alone, K238T/N appear to have minimal effects on NNRTI susceptibility. K238Q is a highly unusual mutation at this position.

Drug resistance mutation scores of NRTI:

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Rule	ABC ⇅	AZT ⇅	D4T ⇅	DDI ⇅	FTC ⇅	3TC ⇅	TDF ⇅
K65R	45	-10	60	60	30	30	50
V75A	0	10	30	15	0	0	0
K65R + S68G	0	0	0	0	0	0	5
Total	45	0	90	75	30	30	55

Drug resistance mutation scores of NNRTI:

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Rule	DOR ⇅	EFV ⇅	ETR ⇅	NVP ⇅	RPV ⇅
L100I	15	60	30	60	60
L100I + K103N	15	0	0	0	0
F227L	60	15	0	30	0
K103N	0	60	0	60	0
Total	90	135	30	150	60