fosamprenavir/r (FPV/r)	Susceptible
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	
K43T is a nonpolymorphic	accessory mutation selected by ATV and LPV. Its phenotypic effect on currently used F
Other	
. Want is the consensus amis	an acid in subtana C and CDC03. AC In subtanas B and C M304 is a Blasclasted mutation

Download CSV

None

K43T

V11M • T12L • I13A • K14R • I15V • G16V • K20I • E21* • T26P • T31P • L33* • E35D • M36I • N37K • R41K • V56E • L63S • I64M • H69K • I72E • N83S • L89M

Drug resistance interpretation: PR

Protease Inhibitors

Susceptible

Susceptible

Rule $|ATV|r \Leftrightarrow |DRV|r \Leftrightarrow |FPV|r \Leftrightarrow |IDV|r \Leftrightarrow |LPV|r \Leftrightarrow |NFV \Leftrightarrow |SQV|r \Leftrightarrow |TPV|r \Leftrightarrow |ATV|r \Leftrightarrow |ATV|r$

PI Major Mutations: PLAccessory Mutations: PR Other Mutations:

atazanavir/r (ATV/r)

darunavir/r (DRV/r)

Mutation scoring: PR

Drug resistance mutation scores of PI:

PIs is uncertain. K20I is the consensus amino acid in subtype G and CRF02_AG. In subtypes B and C, K20I is a PI-selected mutation of uncertain effects on currently used PIs.

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