PatientID: HIVDR-789-23

Sebuttemba 27, 2023

Color Code

HR: High-Level Resistance
LR: Low-Level Resistance
IR: Intermediate Resistance

S: Susceptible

DRUG.CLASS	DRUG	RESISTANCE.PROFILE	DRMS.above.20.percent.prevalence	
PI	ATV	HR		
	DRV	PLR		
	FPV	$^{ m HR}$	M46I;I54V;V82A;T74P;L10F;K43T	
	IDV	$^{ m HR}$		
	LPV	$^{ m HR}$		
	NFV	$^{ m HR}$		
	SQV	$^{ m HR}$		
	TPV	HR		
NRTI	ABC	HR		
	AZT	$^{ m HR}$		
	D4T	$^{ m HR}$		
	DDI	$^{ m HR}$	M41L;D67N;M184V;L210W;T215Y;K219N	
	FTC	$^{ m HR}$		
	LMV	$^{ m HR}$		
	TDF	$^{ m HR}$		
	DOR	IR		
NNRTI	EFV	$^{ m HR}$		
	ETR	IR	H221HY;A98G;V108I;K101H;G190A	
	NVP	$_{ m HR}$		
	RPV	IR		

Appendix

Drug abbreviations in full

DRUG.CLASS	ABBREVIATION	DRUG.NAME
	ATV	Atazanavir
	DRV	Darunavir
	FPV	Fosamprenavir
PI	IDV	Indinavir
11	LPV	Lopinavir
	NFV	Nelfinavir
	SQV	Saquinavir
	TPV	Tipranavir
	ABC	Abacavir
	AZT	Azidothymidine
	DFT	Stavudine
NRTI	DDI	Didanosine
	FTC	Emtricitabine
	LMV	Lamivudine
	TDF	Tenofovir
	DOR	Doravirine
	EFV	Efavirenz
NNRTI	ETR	Etravirine
	NVP	Nevirapine
	RPV	Rilpivirine
	BIC	Bictegravir
	CAB	Cabotegravir
INSTI	DTG	Dolutegravir
	EVG	Elvitegravir
	RAL	Raltegravir

Comments

DRUG.CLASS	COMMENTS		
	I54V is a non-polymorphic PI-selected mutation that contributes reduced susceptibility to		
	each of the PIs except DRV.		
	K43T is a nonpolymorphic accessory mutation selected by ATV and LPV. Its phenotypic		
	effect on currently used PIs is uncertain.		
	L10F is a common non-polymorphic, PI-selected accessory mutation associated with		
	reduced in vitro susceptibility to LPV and DRV.		
	M46I/L are relatively non-polymorphic PI-selected mutations. In combination with other		
	PI-resistance mutations, they are associated with reduced susceptibility to each of the PIs		
	except DRV.		
PI	T74P is a nonpolymorphic PI-selected accessory mutation that occurs primarily in viruses		
	from persons who have received multiple PIs. In combination with other mutations, It is		
	associated with reduced susceptibility to ATV and DRV.		
	V82A is a non-polymorphic mutation selected primarily by IDV and LPV. It is associated		
	with reduced susceptibility to LPV and to a lesser extent ATV. It increases DRV		
	susceptibility.		
	D67N is a non-polymorphic TAM associated with low-level resistance to AZT.		

	K219E/Q/N/R are accessory TAMS that usually occur in combination with multiple other TAMs.
	L210W is a TAM that usually occurs in combination with M41L and T215Y. The
	combination of M41, L210W and T215Y causes high-level resistance to AZT and
	intermediate resistance to 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.
NRTI	M41L is a TAM that usually occurs with T215Y. In combination, M41L plus T215Y confer
111111	intermediate / high-level resistance to AZT and d4T and contribute to reduced ddI, ABC
	and TDF susceptibility.
	T215Y/F are TAMs that causes intermediate/high-level resistance to AZT and potentially
	low-level resistance to ABC and TDF.
	A98G is a non-polymorphic accessory mutation associated with low-level reduced
	susceptibility to each of the NNRTIs.
	G190A is a non-polymorphic mutation that causes high-level resistance to NVP and
	intermediate resistance to EFV. It does not significantly reduce susceptibility to RPV,
	ETR, or DOR.
	H221Y is a non-polymorphic accessory mutation selected primarily by NVP, RPV, and
	DOR. It frequently occurs in combination with Y181C.
	K101H is a non-polymorphic accessory mutation selected by NVP, EFV and ETR. When
NNRTI	present with other NNRTI-resistance mutations, it contributes reduces susceptibility to
	these NNRTIs.
	V108I is a relatively non-polymorphic accessory mutation selected in vitro and/or in vivo
	with each of the NNRTIs. It appears to contribute to reduced susceptibility to most
	NNRTIs only in combination with other NNRTI-resistance mutations.
INSTI	