PatientID: HDR45

Okitobba 06, 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	\mathbf{S}	
	DRV	\mathbf{S}	
	FPV	\mathbf{S}	
	IDV	\mathbf{S}	
	LPV	\mathbf{S}	
	NFV	\mathbf{S}	
	SQV	\mathbf{S}	
	TPV	${f S}$	
	ABC	LR	
	AZT	${f S}$	
	D4T	${f S}$	
NRTI	DDI	PLR	M184V
	FTC	$_{ m HR}$	
	LMV	$_{ m HR}$	
	TDF	\mathbf{S}	
NNRTI	DOR	$_{ m HR}$	
	EFV	$_{ m HR}$	
	ETR	LR	F227I;K101H;G190A
	NVP	$_{ m HR}$	
	RPV	LR	
INSTI	BIC	${f S}$	
	CAB	${f S}$	
	DTG	${f S}$	
	EVG	${f S}$	
	RAL	${f S}$	

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		
PI			
NRTI	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.		
	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.		
	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,		
NNRTI	ETR, or DOR.		
	K101H is a non-polymorphic accessory mutation selected by NVP, EFV and ETR. When		
	present with other NNRTI-resistance mutations, it contributes reduces susceptibility to		
	these NNRTIs.		
INSTI			