

Samuel H. Pepkowitz, MD, Medical Director
 345 Oyster Point Blvd
 South San Francisco, CA 94080 - Tel: (800) 777-0177

Patient Name:	DOB	Patient ID/Medical Record #	Gender	Monogram Accession #
Date Collected	Date Received	Date Reported	Mode	Report Status
			Reference Lab ID/Order #	
Comments			HIV-1 Subtype: B	

	DRUG		PHENOSENSE™ SUSCEPTIBILITY				Evidence of Susceptibility		Net Assessment	
	Generic Name	Brand Name	Cutoffs (Lower - Upper)	Fold Change	Drug Susceptibility		Pheno Sense	Gene Seq		
NRTI	Abacavir	Ziagen	(4.5 - 6.5)	3.98			Y	N	Sensitive	16
	Didanosine	Videx	(1.3 - 2.2)	1.99			P	N	Partially Sensitive	
	Emtricitabine	Emtriva	(3.5)	>MAX			N	N	Resistant	
	Lamivudine	Epivir	(3.5)	>MAX			N	N	Resistant	
	Stavudine	Zerit	(1.7)	1.51			Y	N	Sensitive	3
	Zidovudine	Retrovir	(1.9)	7.91			N	N	Resistant	3
	Tenofovir	Viread	(1.4 - 4)	1.16			Y	N	Sensitive	3
	NRTI Mutations		M41L, M184V, T215Y							

NNRTI	Delavirdine	Rescriptor	(6.2)	3.91			Y	N	Sensitive	1
	Efavirenz	Sustiva	(3)	30			N	N	Resistant	
	Etravirine	Intelence	(2.9 - 10)	0.56			Y	N	Sensitive	1
	Nevirapine	Viramune	(4.5)	>MAX			N	N	Resistant	
	Rilpivirine	Edurant	(2)	1.29			Y	N	Resistant	1
NNRTI Mutations		Y188Y/F/L, H221H/Y								

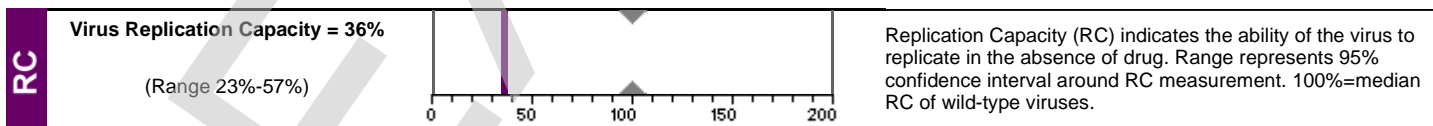
PI	Atazanavir	Reyataz	(2.2)	4.96			N	N	Resistant	
		Reyataz / r†	(5.2)	4.96			Y	N	Sensitive	16
	Darunavir	Prezista / r†	(10 - 90)	1.34			Y	Y	Sensitive	
	Fosamprenavir	Lexiva / r†	(4 - 11)	4.00			Y	Y	Sensitive	
	Indinavir	Crixivan / r†	(10)	5.81			Y	Y	Sensitive	
	Lopinavir	Kaletra†	(9 - 55)	1.69			Y	Y	Sensitive	
	Nelfinavir	Viracept	(3.6)	17			N	N	Resistant	
	Ritonavir	Norvir	(2.5)	4.30			N	N	Resistant	
	Saquinavir	Invirase / r†	(2.3 - 12)	3.88			P	N	Partially Sensitive	
	Tipranavir	Aptivus / r†	(2 - 8)	2.87			P	N	Partially Sensitive	
PI Mutations		L10V, I13V, K20T, E35G, M36I, I62V, L63T, T74S, L90M								

Lower Clinical Cutoff (in bold)
 Upper Clinical Cutoff (in bold)
 Biological Cutoff
 Hypersusceptibility
 Cutoff
 Sensitive
 Partially Sensitive
 Resistant
 Y Evidence of Drug Sensitivity
 P Evidence of Partial Drug Sensitivity
 N Evidence of Drug Resistance

Samuel H. Pepkowitz, MD, Medical Director
345 Oyster Point Blvd
South San Francisco, CA 94080 - Tel: (800) 777-0177

Patient Name:	Date Collected:	Monogram Acc#:	Status:
---------------	-----------------	----------------	---------

Combination Phenotype/Genotype Net Assessment			
	SENSITIVE	PARTIALLY SENSITIVE	RESISTANT
NRTI	Abacavir Stavudine Tenofovir	Didanosine	Emtricitabine Lamivudine Zidovudine
NNRTI	Delavirdine Etravirine		Efavirenz Nevirapine Rilpivirine
PI	Atazanavir / r Darunavir / r Fosamprenavir / r Indinavir / r Lopinavir / r	Saquinavir / r Tipranavir / r	Atazanavir Nelfinavir Ritonavir



Phenotype / Genotype Comments (clinical significance may vary)

- 1 - **Mixture**: Mixtures detected at resistance-associated position(s); minor populations with decreased susceptibility may be present and may increase in the presence of drug pressure.
- 3 - **IC50 reduced**: Phenotypic measurement reflects possible enhanced susceptibility due to M184I or V.
- 16 - **Unexplained discordance**: Genotypic correlates of susceptibility not accounted for by current rules.

Samuel H. Pepkowitz, MD, Medical Director
 345 Oyster Point Blvd
 South San Francisco, CA 94080 - Tel: (800) 777-0177

Patient Name:	Date Collected:	Monogram Acc#:	Status:
---------------	-----------------	----------------	---------

Complete List of Mutations Detected

RT: P19P/L, V21V/I, V35V/I, M41L, V60I, Q102K, K122K/E, I135T, C162S, M184V, Y188Y/F/L, G196E, Q207E, T215Y, H221H/Y, A272A/S, R277R/K, L283I, P294T, E297E/K

PR: L10V, I13V, K14K/R, I15V, K20T, E35G, M36I, N37D/E, I62V, L63T, I64V, E65D, I72V, T74S, V77I, L90M, I93L

Important Definitions

IC50: Concentration of drug required to inhibit viral replication by 50%.

$$\text{Fold Change} = \frac{\text{IC50 patient}}{\text{IC50 reference}}$$

Clinical Cutoffs: *Lower clinical cutoff* denotes the fold change which was the best discriminator of reduced clinical response using drug-specific clinical outcome data. Reduced response was defined by the clinical endpoint for the specific clinical cohort analyzed for each cutoff value. *Upper clinical cutoff* denotes the fold change above which a clinical response is unlikely (<.5 log reduction in HIV RNA) and which was determined using the same drug-specific clinical cohort data as for the lower clinical cutoff. Biological cutoffs are used for specific antiretrovirals (ZDV, the NNRTIs and specific protease inhibitors when not pharmacokinetically enhanced with ritonavir). These values are defined as the fold change value below which reside 99% of tested wild-type isolates, i.e., those without known drug resistance mutations. Fold Change <0.4 indicates enhanced susceptibility.

Mixtures are indicated by amino acids separated by a slash. Deletions in the amino acid sequence are indicated by a ^ symbol.

* **Boosted PIs:** Clinical cutoff and genotypic interpretation algorithms for ritonavir-boosted protease inhibitors derived from individual studies using the following dosages: AMP/r 600mg/100mg BID; ATV/r 300mg/100mg QD; DRV/r 600mg/100mg BID; IDV/r 800mg/200mg BID; LPV/r 400mg/100mg BID; SQV/r 1000mg/100mg BID; and TPV/r 500mg/200mg BID.

Assessment of drug susceptibility is based upon detected mutations and interpreted using an advanced proprietary algorithm (version 14)

Patient-Specific Results

Drugs	ABC	ddl	FTC	3TC	d4T	ZDV	TFV	DLV	EFV	ETR	NVP	RPV	ATV	DRV	AMP	IDV	LPV	NFV	RTV	SQV	TPV
IC50 (µM)	6.6	11.56	>100	>300	1.4	0.349	1.274	0.118	0.1157	0.001315	>20	0.001275	0.009990	0.001211	0.0567	0.0408	0.0121	0.2611	0.163	0.0197	0.3945
Fold Change	3.98	1.99	>MAX	>MAX	1.51	7.91	1.16	3.91	30	0.56	>MAX	1.29	4.96	1.34	4.00	5.81	1.69	17	4.30	3.88	2.87

For more information on interpreting this report, please visit www.MonogramHIV.com or call Customer Service at 800-777-0177 between the hours of 6:30am to 5:00pm PT Monday through Friday.

PhenoSense GT is a proprietary assay that combines the technology of PhenoSense HIV and GeneSeq HIV with expert interpretation. This assay meets the standards for performance characteristics and all other quality control and assurance requirements established by the Clinical Laboratory Improvement Amendments. This test is validated for testing specimens with HIV-1 viral loads equal to or above 500 copies/mL and should be interpreted only on such specimens. The results should not be used as the sole criteria for patient management. The results have been disclosed to you from confidential records protected by law and are not to be disclosed to unauthorized persons. Further disclosure of these results is prohibited without specific consent of the persons to whom it pertains, or as permitted by law.