

# HIV Drug Resistance Testing Basics

**Brian R. Wood, MD**  
**Medical Director, MWAETC Project ECHO**  
**Associate Professor of Medicine, University of Washington**

Last Updated: April 20, 2023

# Disclosures

---

No conflicts of interest or relationships to disclose.

# Disclaimer

Funding for this presentation was made possible by U1OHA29296 from the Human Resources and Services Administration HIV/AIDS Bureau. The views expressed do not necessarily reflect the official policies of the Department of Health and Human Services nor does mention of trade names, commercial practices, or organizations imply endorsement by the U.S. Government. *Any trade/brand names for products mentioned during this presentation are for training and identification purposes only.*

# Learning Objectives

- Understand the process of genotype versus phenotype resistance testing and why one is preferred over the other
- Describe the indications for a traditional genotype (RT/PR), an integrase (IN) genotype, and a phenotype
- Know the resources for help with interpretation of resistance-associated mutations

# Poll

You are seeing a patient recently diagnosed with HIV for their first clinic visit. They have no history of PrEP use. Which of the following is recommended as part of the baseline laboratory evaluation?

- A) Genotype resistance assay (integrase resistance testing not necessary)
- B) Genotype resistance assay with integrase resistance testing
- C) Phenotype resistance assay with integrase resistance testing
- D) Genotype and phenotype resistance assays (integrase not necessary)

# Resistance Test Comparison

Genotype	Phenotype
Sequence reverse transcriptase (RT) and protease (PR) genes, +/- integrase (IN) gene (or, rarely, envelope gene)	Grow virus in culture, add ARV drugs in various amounts, compare IC <sub>50</sub> to IC <sub>50</sub> of wild type virus (“fold change”)
Quicker, lower cost, more sensitive	Takes longer, more expensive
Interpretation challenging if numerous mutations	Helpful if complex resistance history, especially to protease inhibitors (PI)

\*Both types: need circulating RNA; resistance only detected if >10-20% of virus population

# Example genotype report

## HIV-1 Genotyping

See Note

### NRTI DRUGS

EPIVIR, (lamivudine, 3TC)	None
EMTRIVA, (emtricitabine, FTC)	None
RETROVIR, (zidovudine, AZT)	None
VIDEX, (didanosine, ddI)	None
ZERIT, (stavudine, d4T)	None
ZIAGEN, (abacavir, ABC)	None
VIREAD, (tenofovir, TDF)	None

NRTI associated resistance mutations found: None

### NNRTI DRUGS

RESCRIPTOR, (delavirdine, DLV)	Resistance
SUSTIVA, (efavirenz, EFV)	Resistance
VIRAMUNE, (nevirapine, NVP)	Resistance
INTELENCE, (etravirine, ETR)	None

NNRTI associated resistance mutations found: K103N

### Protease inhibitors

AGENERASE, (amprenavir, APV)	None
LEXIVA, (fosamprenavir, FOS)	None
CRIXIVAN, (indinavir, IDV)	None
FORTOVASE / INVIRASE, (saquinavir, SQV)	None
KALETRA, (lopinavir + ritonavir, LPV)	None
PREZISTA, (darunavir, DRV)	None
VIRACEPT, (nelfinavir, NFV)	None
REYATAZ, (atazanavir, ATV)	None
APTIVUS, (tipranavir, TPV)	None

# Example phenotype report

	DRUG		PHENOTYPE™ SUSCEPTIBILITY			ASSESSMENT		
	Generic Name	Brand Name	Cutoffs (Lower - Upper)	Fold Change	Increasing Drug Susceptibility	Decreasing	Drug	
NRTI	Abacavir	Ziagen	(4.5 - 6.5)	1.20			ABC	Sensitive
	Didanosine	Videx	(1.3 - 2.2)	1.38			ddl	Partially Sensitive
	Emtricitabine	Emtriva	(3.5)	1.20			FTC	Sensitive
	Lamivudine	EpiVir	(3.5)	1.27			3TC	Sensitive
	Stavudine	Zerit	(1.7)	1.20			d4T	Sensitive
	Tenofovir	Viread	(1.4 - 4)	1.16			TFV	Sensitive
	Zidovudine	Retrovir	(1.9)	1.28			ZDV	Sensitive

NNRTI	Delavirdine	Rescriptor	(6.2)	3.10			DLV	Sensitive
	Efavirenz	Sustiva	(3)	1.18			EFV	Sensitive
	Etravirine	Intelence	(2.9 - 10)	1.28			ETR	Sensitive
	Nevirapine	Viramune	(4.5)	1.39			NVP	Sensitive
	Rilpivirine	Edurant	(2)	1.29			RPV	Sensitive

PI	Atazanavir	Reyataz	(2.2)	3.07			ATV	Resistant
		Reyataz / r*	(5.2)	3.07			ATV/r	Sensitive
	Darunavir	Prezista / r*	(10 - 90)	4.13			DRV/r	Sensitive
	Fosamprenavir	Lexiva / r*	(4 - 11)	3.92			AMP/r	Sensitive
	Indinavir	Crixivan / r*	(10)	1.07			IDV/r	Sensitive
	Lopinavir	Kaletra*	(9 - 55)	2.50			LPV/r	Sensitive
	Nelfinavir	Viracept	(3.6)	1.28			NFV	Sensitive
	Ritonavir	Norvir	(2.5)	5.04			RTV	Resistant
	Saquinavir	Invirase / r*	(2.3 - 12)	2.05			SQV/r	Sensitive
	Tipranavir	Aplivus / r*	(2 - 8)	3.07			TPV/r	Partially Sensitive

Lower Clinical Cutoff (in bold)  
 Upper Clinical Cutoff (in bold)  
 Biological Cutoff

Hypersusceptibility  
 Cutoff

Sensitive  
 Partial Sensitivity  
 Resistance

# Example phenotype report

DRUG		PHENOSENSE™ SUSCEPTIBILITY			Evidence of Susceptibility		Net Assessment	
Generic Name	Brand Name	Cutoffs (Lower - Upper)	Fold Change	Increasing Drug Susceptibility	Decreasing	Pheno Sense		Gene Seq
Abacavir	Ziagen	(4.5 - 6.5)	>MAX			N	N	Resistant
Didanosine	Videx	(1.3 - 2.2)	20			N	N	Resistant
Emtricitabine	Emtriva	(3.5)	>MAX			N	N	Resistant
Lamivudine	Eplivir	(3.5)	>MAX			N	N	Resistant
Stavudine	Zerit	(1.7)	7.87			N	N	Resistant
Zidovudine	Retrovir	(1.9)	282			N	N	Resistant
Tenofovir	Viread	(1.4 - 4)	1.71			P	N	Partially Sensitive
NRTI Mutations		A62V, T69I/V, V75I, F77L, Y115F, F116Y, Q151M, M184V, K219K/N						
Delavirdine	Rescriptor	(5.2)	>MAX			N	N	Resistant
Efavirenz	Sustiva	(3)	24			N	N	Resistant
Etravirine	Intelence	(2.9 - 10)	106			N	N	Resistant
Nevirapine	Viramune	(4.5)	>MAX			N	N	Resistant
NNRTI Mutations		V179V/I, Y181I, V189V/I, G190A						
Atazanavir	Reyataz	(2.2)	>MAX			N	N	Resistant
	Reyataz / r <sup>2</sup>	(5.2)	>MAX			N	N	Resistant
Darunavir	Prezista / r <sup>2</sup>	(10 - 90)	>MAX			N	N	Resistant
Fosamprenavir	Lexiva / r <sup>2</sup>	(4 - 11)	>MAX			N	N	Resistant
Indinavir	Crixivan / r <sup>2</sup>	(10)	30			N	N	Resistant
Lopinavir	Kaletra	(9 - 55)	>MAX			N	N	Resistant
Nelfinavir	Viracapt	(3.6)	38			N	N	Resistant
Ritonavir	Norvir	(2.5)	>MAX			N	N	Resistant
Saquinavir	Inlvase / r <sup>2</sup>	(2.3 - 12)	19			N	N	Resistant
Tipranavir	Aptivus / r <sup>2</sup>	(2 - 8)	27			N	N	Resistant
PI Mutations		L10V, V11I, I13V, K20T, V32I, L33F, E35D, M36I, M46L, I54L, D60E, A71V, G73T, V82V/I, I84V						

# Indications for Genotype Resistance Testing

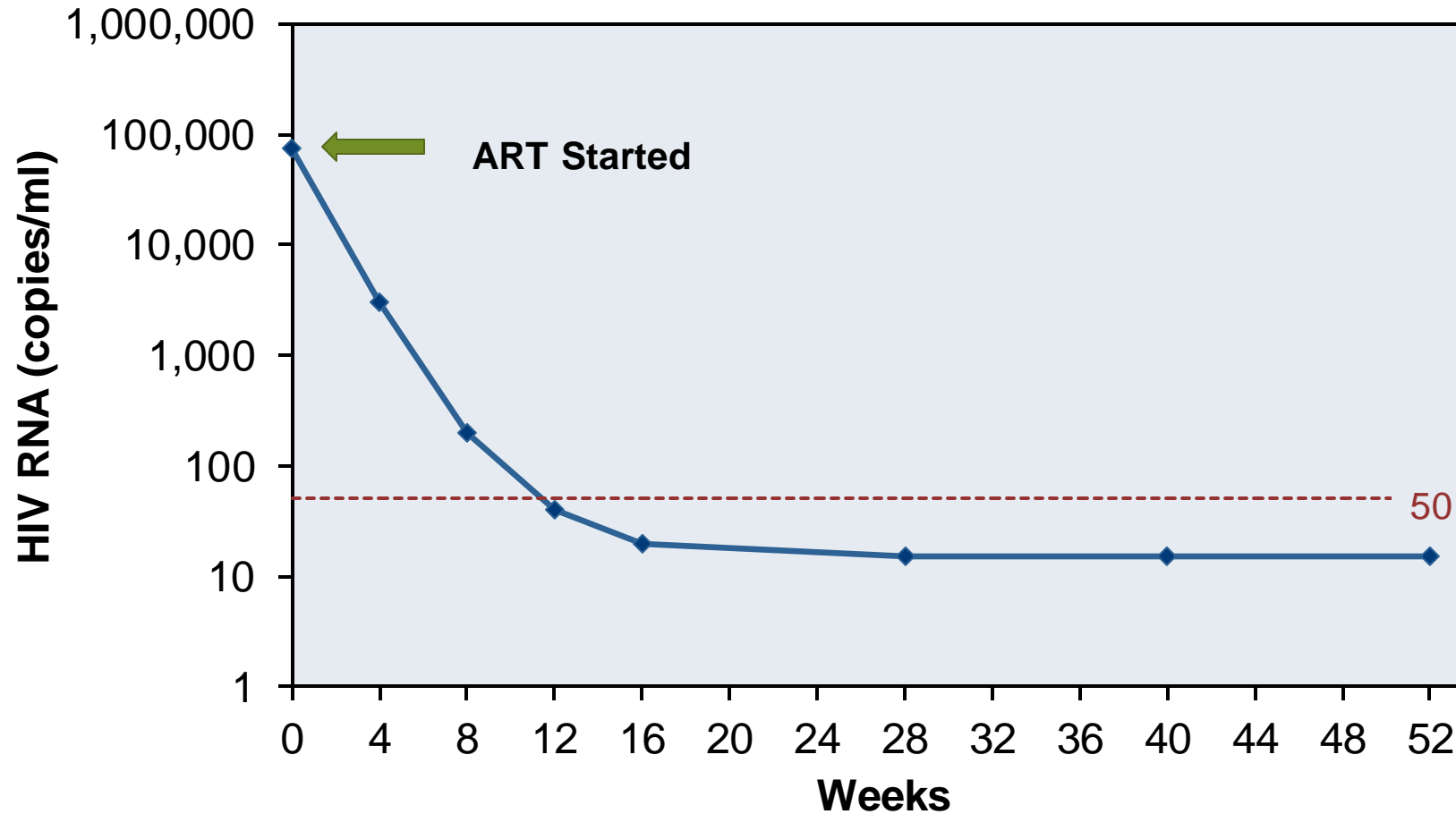
- **Indication #1: all treatment-naïve patients at entry into care**
  - Frequency of transmitted mutations: 5-15% (mostly NNRTI)
  - Check even if deferring ART
  - Ok to start ART before results return
  - Integrase resistance testing not routinely indicated at baseline

# Indications for Genotype Resistance Testing

- **Indication #2: Virologic failure or suboptimal virologic suppression**
  - Virologic failure: HIV RNA rebound to >200 copies/mL (genotype may be unsuccessful if RNA 200-500 copies/mL, but should be considered)
  - For non-long-acting ART, ideally perform genotype within 4 weeks of stopping ART (not always realistic)

# Virologic Responses on Antiretroviral Therapy

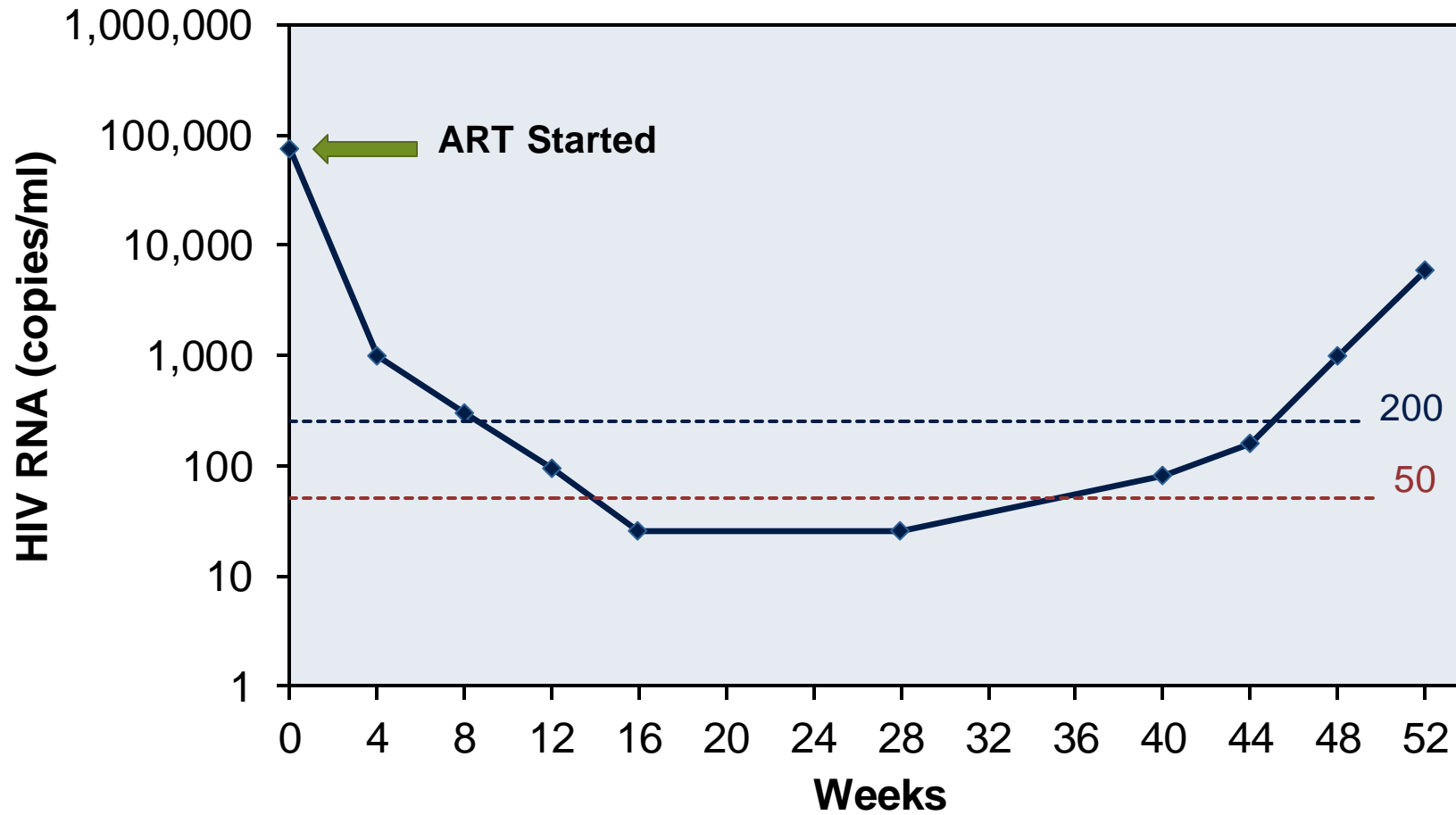
## Virologic Suppression



A confirmed HIV RNA level below the lower limit of assay detection.

# Virologic Responses on Antiretroviral Therapy

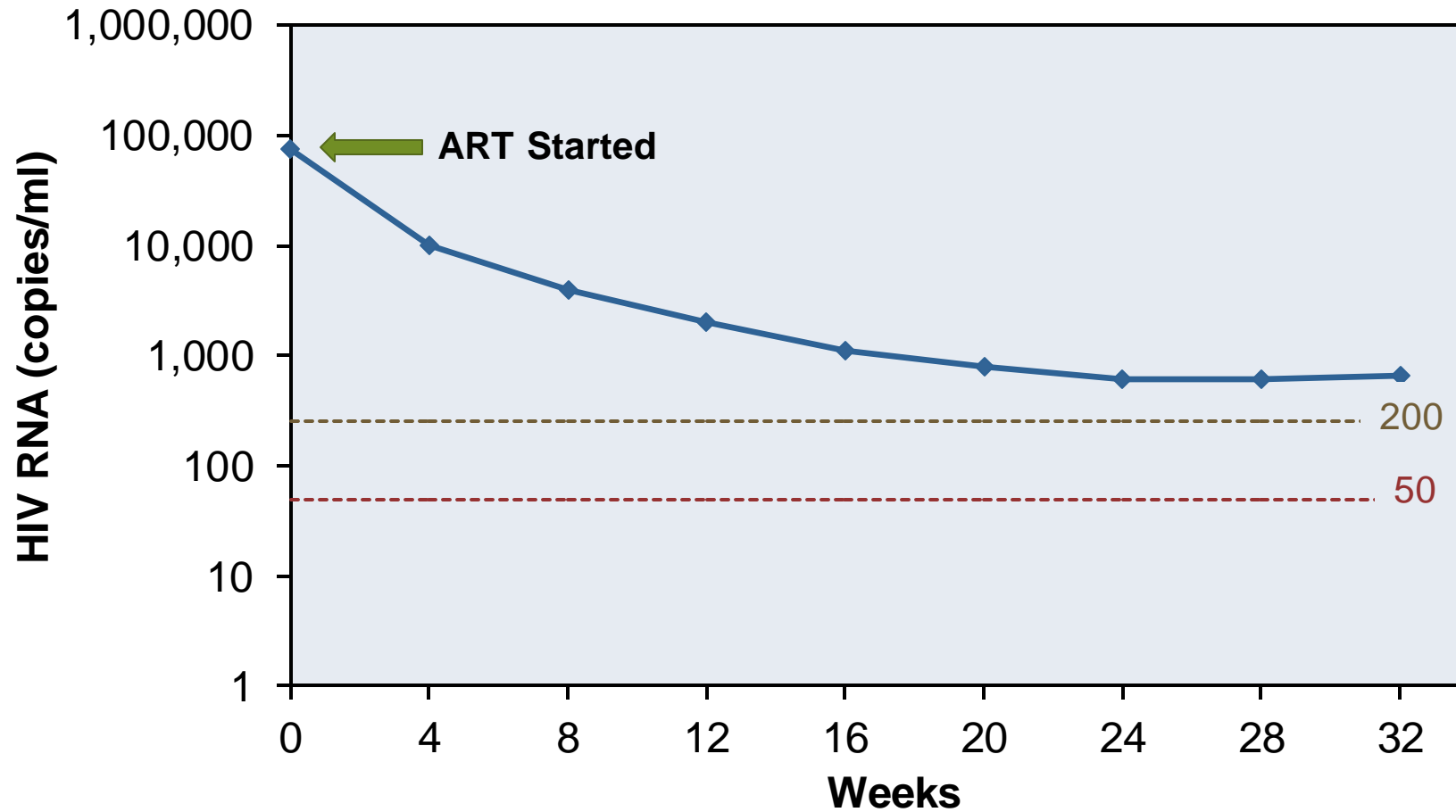
## Virologic Failure



Confirmed HIV RNA  $\geq 200$  copies/mL after virologic suppression

# Virologic Responses on Antiretroviral Therapy

## Incomplete or Suboptimal Virologic Response



Two consecutive plasma HIV RNA levels >200 copies/mL after 24 weeks on an ARV regimen.  
Baseline HIV RNA may affect the time course of response, and some regimens will take longer than others to suppress HIV RNA levels.

# Indications for Integrase (IN) Genotype

- **Indication #1:** virologic failure while taking an integrase inhibitor
- **Indication #2:** add to RT/PR genotype at baseline if past integrase inhibitor exposure (prior cabotegravir for PrEP) or integrase inhibitor resistance exposure

\*Remember, integrase resistance testing may require a separate order!

# Another Genotype Option: PBMC DNA Resistance Testing (also called: archive, DNA, proviral, or PBMC genotype)

- What is it? Sequence mutations in **proviral DNA, instead of plasma RNA**
- Advantage: available at any RNA level, including undetectable
- Disadvantage: less sensitive than cumulative RNA genotypes
  - Why? Takes weeks to months for mutations to accumulate in PBMCs, especially if low HIV RNA levels or periods of virologic failure brief
- **Indication:** taking salvage ART, need resistance data in order to change or simplify regimen, and cannot obtain past RNA genotype results

1. Delaugerre C et al. HIV Medicine. 2012;13:517–525.

2. Chu C, et al. Clin Microbiol Rev. 2022 Dec 21;35(4):e0005222.

# Indications for Phenotype Resistance Testing

- Per guidelines: add to genotype if known or suspected complex mutation pattern
- In practice: almost never

# Case

- A 55-year-old patient, who was prescribed rilpivirine/tenofovir alafenamide/emtricitabine, presents after an absence from care. They report missed doses of ART over the prior 3 months. The prior HIV RNA levels were suppressed but a repeat level returns at just over 1,000 copies/mL and an RT/PR genotype resistance assay shows the RT mutations K103N and M184V.
- How can you obtain help interpreting the effects of these mutations?



Stanford University

# HIV DRUG RESISTANCE DATABASE

*A curated public database to represent, store and analyze HIV drug resistance data.*

HOME

GENOTYPE-RX

GENOTYPE-PHENO

GENOTYPE-CLINICAL

HIVDB PROGRAM

ABOUT HIVDB

SUPPORT HIVDB!



**Sierra 3.4.2**  
[release notes / web service](#)  
 Dec 14, 2022

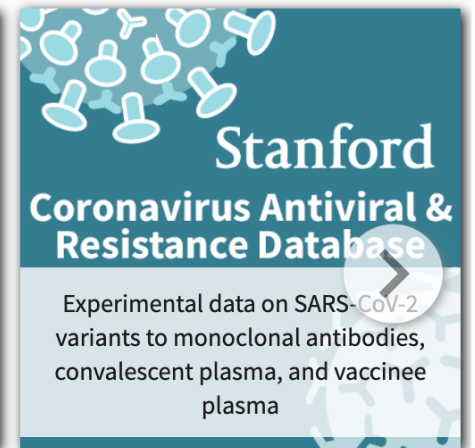


**HIVDB Algorithm Version 9.4**  
 Dec 7, 2022

**HIV in vitro selection**  
 HIV in vitro selected PR, RT, IN and CA mutations  
 Mar 13, 2023



**<ASIEditor />**  
 JavaScript-based Algorithm Specification Interface (ASI) editor  
 Aug 18, 2022



**Stanford Coronavirus Antiviral & Resistance Database**  
 Experimental data on SARS-CoV-2 variants to monoclonal antibodies, convalescent plasma, and vaccinee plasma



**CPR** Calibrated Population Resistance

**HIVDB released on January 10, 2023**

**Query / Download**

**HIVdb Program**

## Reverse Transcriptase

Input mutation(s)

Select mutations:

40	41	44	62
---	---	---	---
65	67	68	69
---	---	---	---
70	74	75	77
---	---	---	---
90	98	100	101
---	---	---	---
103	106	108	115
---	---	---	---
116	118	138	151
---	---	---	---
179	181	184	188
---	---	---	---
190	210	215	219
---	---	---	---
221	225	227	230
---	---	---	---
234	236	238	318
---	---	---	---

## Protease

Input mutation(s)

Select mutations:

10	11	13	20
---	---	---	---
23	24	30	32
---	---	---	---
33	35	36	43
---	---	---	---
46	47	48	50
---	---	---	---
53	54	58	63
---	---	---	---
71	73	74	76
---	---	---	---
77	82	83	84
---	---	---	---
85	88	89	90
---	---	---	---
93			
---			

## Integrase

Input mutation(s)

Select mutations:

51	66	74	92
---	---	---	---
95	97	114	118
---	---	---	---
121	128	138	140
---	---	---	---
143	145	146	147
---	---	---	---
148	151	153	155
---	---	---	---
157	163	230	263
---	---	---	---

### Reverse Transcriptase

Input mutation(s)

### Protease

Input mutation(s)

### Integrase

Input mutation(s)

Select mutations:

40	41	44	62
---	---	---	---
65	67	68	69
---	---	---	---
70	74	75	77
---	---	---	---
90	98	100	101
---	---	---	---
103	106	108	115
---	---	---	---
116	118	138	151
---	---	---	---
179	181	184	188
---	---	---	---
190	210	I	219
---	---	V	---
221	225	*	230
---	---	---	---
234	236	238	318
---	---	---	---
348			
---			

Select mutations:

10	11	13	20
---	---	---	---
23	24	30	32
---	---	---	---
33	35	36	43
---	---	---	---
46	47	48	50
---	---	---	---
53	54	58	63
---	---	---	---
71	73	74	76
---	---	---	---
77	82	83	84
---	---	---	---
85	88	89	90
---	---	---	---
93			
---			

Select mutations:

51	66	74	92
---	---	---	---
95	97	114	118
---	---	---	---
121	128	138	140
---	---	---	---
143	145	146	147
---	---	---	---
148	151	153	155
---	---	---	---
157	163	230	263
---	---	---	---

Select mutations:

40	41	44	62
---	---	---	---
65	67	68	69
---	---	---	---
70	74	75	77
---	---	---	---
90	98	100	101
---	---	---	---
103	106	108	115
---	---	---	---
E	118	138	151
H	---	---	---
N	181	184	188
Q	---	---	---
R	210	215	219
S	---	---	---
T	225	227	230
*	---	---	---
...	236	238	318
---	---	---	---

Select mutations:

10	11	13	20
---	---	---	---
23	24	30	32
---	---	---	---
33	35	36	43
---	---	---	---
46	47	48	50
---	---	---	---
53	54	58	63
---	---	---	---
71	73	74	76
---	---	---	---
77	82	83	84
---	---	---	---
85	88	89	90
---	---	---	---
93			
---			

Select mutations:

51	66	74	92
---	---	---	---
95	97	114	118
---	---	---	---
121	128	138	140
---	---	---	---
143	145	146	147
---	---	---	---
148	151	153	155
---	---	---	---
157	163	230	263
---	---	---	---

K103N x M184V x Input mutation(s)

Input mutation(s)

Input mutation(s)

Select mutations:

40	41	44	62
---	---	---	---
65	67	68	69
---	---	---	---
70	74	75	77
---	---	---	---
90	98	100	101
---	---	---	---
103	106	108	115
*	---	---	---
116	118	138	151
---	---	---	---
179	181	184	188
---	---	---	---
190	210	215	219
---	---	---	---
221	225	227	230
---	---	---	---
234	236	238	318
---	---	---	---
348			
---			

Select mutations:

10	11	13	20
---	---	---	---
23	24	30	32
---	---	---	---
33	35	36	43
---	---	---	---
46	47	48	50
---	---	---	---
53	54	58	63
---	---	---	---
71	73	74	76
---	---	---	---
77	82	83	84
---	---	---	---
85	88	89	90
---	---	---	---
93			
---			

Select mutations:

51	66	74	92
---	---	---	---
95	97	114	118
---	---	---	---
121	128	138	140
---	---	---	---
143	145	146	147
---	---	---	---
148	151	153	155
---	---	---	---
157	163	230	263
---	---	---	---

Keep input mutations when browsing back

Reset

Analyze



## **Nucleoside Reverse Transcriptase Inhibitors**

---

<b>abacavir (ABC)</b>	Low-Level Resistance
<b>zidovudine (AZT)</b>	Susceptible
<b>emtricitabine (FTC)</b>	High-Level Resistance
<b>lamivudine (3TC)</b>	High-Level Resistance
<b>tenofovir (TDF)</b>	Susceptible

## **Non-nucleoside Reverse Transcriptase Inhibitors**

---

<b>doravirine (DOR)</b>	Susceptible
<b>efavirenz (EFV)</b>	High-Level Resistance
<b>etravirine (ETR)</b>	Susceptible
<b>nevirapine (NVP)</b>	High-Level Resistance
<b>rilpivirine (RPV)</b>	Susceptible

## RT comments

### NRTI

- **M184V/I** cause high-level in vitro resistance to 3TC and FTC and low-level resistance to ddI and ABC. However, **M184V/I** are not contraindications to continued treatment with 3TC or FTC because they increase susceptibility to AZT, TDF and d4T and are associated with clinically significant reductions in HIV-1 replication.

### NNRTI

- **K103N** is a non-polymorphic mutation that causes high-level reductions in NVP and EFV susceptibility.

## Mutation scoring: RT

*Drug resistance mutation scores of NRTI:*

Copy to clipboard



Rule	ABC ⚡	AZT ⚡	FTC ⚡	3TC ⚡	TDF ⚡
<u>M184V</u>	15	-10	60	60	-10

*Drug resistance mutation scores of NNRTI:*

Copy to clipboard



Rule	DOR ⚡	EFV ⚡	ETR ⚡	NVP ⚡	RPV ⚡
<u>K103N</u>	0	60	0	60	0

\*Scores <10 indicate susceptible; scores 10-14 indicate potential low-level resistance; scores 15-29 indicate low-level resistance; scores 30-59 indicate intermediate resistance; scores 60 or higher indicate high-level resistance.

# Case

- A 31-year-old patient, who was prescribed elvitegravir/cobicistat/tenofovir alafenamide/emtricitabine, presents for a first visit after transferring care. They report that due to a move and other factors they were taking their medication every other day for about 3 months and then were completely out for about 3 months. Lab testing is performed, including an RT/PR genotype and IN genotype. These demonstrate the RT mutations M184V, K65R, and L74V, plus IN mutations Q148H/K and G140A/S.
- How would you interpret these resistance mutations?

## Reverse Transcriptase

K65R x L74V x M184V x

Input mutation(s)

## Protease

Input mutation(s)

## Integrase

G140AS x Q148HK x

Input mutation(s)

## Drug resistance interpretation: RT

NRTI Resistance Mutations:	<b>K65R, L74V, M184V</b>
NNRTI Resistance Mutations:	None
Other Mutations:	None

### **Nucleoside Reverse Transcriptase Inhibitors**

---

<b>abacavir (ABC)</b>	High-Level Resistance
<b>zidovudine (AZT)</b>	Susceptible
<b>emtricitabine (FTC)</b>	High-Level Resistance
<b>lamivudine (3TC)</b>	High-Level Resistance
<b>tenofovir (TDF)</b>	Intermediate Resistance

### **Non-nucleoside Reverse Transcriptase Inhibitors**

---

<b>doravirine (DOR)</b>	Susceptible
<b>efavirenz (EFV)</b>	Susceptible
<b>etravirine (ETR)</b>	Susceptible
<b>nevirapine (NVP)</b>	Susceptible
<b>rilpivirine (RPV)</b>	Susceptible

## RT comments

### NRTI

- **K65R** causes intermediate/high-level resistance to TDF, ddI, ABC and d4T and low/intermediate resistance to 3TC and FTC. **K65R** increases susceptibility to AZT.
- **L74V/I** cause high-level resistance to ddI and intermediate resistance to ABC.
- **M184V/I** cause high-level in vitro resistance to 3TC and FTC and low-level resistance to ddI and ABC. However, **M184V/I** are not contraindications to continued treatment with 3TC or FTC because they increase susceptibility to AZT, TDF and d4T and are associated with clinically significant reductions in HIV-1 replication.

## Mutation scoring: RT

*Drug resistance mutation scores of NRTI:*

Copy to clipboard



Rule	ABC ↕	AZT ↕	FTC ↕	3TC ↕	TDF ↕
<u>K65R</u>	45	-10	30	30	50
<u>L74V</u>	30	0	0	0	0
<u>L74V + M184V</u>	15	0	0	0	0
<u>M184V</u>	15	-10	60	60	-10
<b>Total</b>	105	-20	90	90	40

## Drug resistance interpretation: IN

IN Major Resistance Mutations:	<b>G140AS, Q148HK</b>
IN Accessory Resistance Mutations:	None
Other Mutations:	None

### **Integrase Strand Transfer Inhibitors**

---

<b>bictegravir (BIC)</b>	Intermediate Resistance
<b>dolutegravir (DTG)</b>	Intermediate Resistance
<b>elvitegravir (EVG)</b>	High-Level Resistance
<b>raltegravir (RAL)</b>	High-Level Resistance

## IN comments

### IN Major

- **G140S/A/C** are non-polymorphic mutations that usually occur with Q148 mutations. Alone, they have minimal effects on INSTI susceptibility. However, in combination with Q148 mutations they are associated with high-level resistance to RAL and EVG and intermediate reductions in DTG and BIC susceptibility.
- **Q148H/K/R** are non-polymorphic mutations selected by RAL, EVG, and rarely DTG. **Q148H/R/K** are associated with high-level reductions in RAL and EVG susceptibility particularly when they occur in combination with E138 or G140 mutations. Alone, **Q148H/K/R** have minimal effects on DTG and BIC susceptibility. But in combination with E138 and G140 mutations they cause moderate and occasionally high-level reductions in DTG and BIC susceptibility.

### Dosage Considerations

- There is evidence for intermediate **DTG** resistance. If **DTG** is used, it should be administered twice daily.

## Mutation scoring: IN

*Drug resistance mutation scores of INSTI:*

Copy to clipboard



Rule	BIC ⚡	CAB ⚡	DTG ⚡	EVG ⚡	RAL ⚡
<u>G140AS</u>	10	10	10	30	30
<u>G140AS + Q148HK</u>	10	20	10	0	0
<u>Q148HK</u>	30	50	30	60	60
Total	50	80	50	90	90

# Take-Home Points

- Genotype is the principal resistance test used in clinical care
  - Indicated for all at baseline (integrase testing not routinely indicated)
  - Also indicated for virologic failure or incomplete virologic response
  - If virologic failure occurs while taking integrase inhibitor, add integrase testing
  - Genotype of proviral DNA in PBMC (aka, archive genotype) rarely indicated
- Stanford Database (db) is a powerful tool for interpreting & learning mutations
  - Remember to enter all resistance mutations from all past genotype tests!

# Other Resources for Learning Key Mutations

- National HIV Curriculum module Evaluating and Managing Virologic Failure:  
<https://www.hiv.uw.edu>
- Project ECHO video archive:  
<https://www.youtube.com/@MWAETCProjectECHO/videos>
- Prior relevant ECHO talks:
  - Introduction to HIV Resistance Testing (Spach)
  - NNRTI Resistance (Wood)
  - NNRTI Resistance 2015 (Spach)
  - NRTI Resistance (Wood)
  - Resistance to Integrase Strand Transfer Inhibitors (Spach)
  - Recent Trials of Second-Line ART (Wood)
  - Management of NRTI Resistance (Spach)

# Acknowledgment

This Mountain West AIDS Education and Training (MWAETC) program is supported by the Health Resources and Services Administration (HRSA) of the U.S. Department of Health and Human Services (HHS) as part of an award totaling \$3,098,654 with 0% financed with non-governmental sources.

The content in this presentation are those of the author(s) and do not necessarily represent the official views of, nor an endorsement by, HRSA, HHS, or the U.S. Government.

