



Lessons Learned from a Novice HIV Pharmacist

HIV Pharmacy Education Day 2025
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- **About me:**
 - Community Pharmacist – Prime Care Pharmacy
 - HIV Pharmacy Consulting – Hive Health Services
- **Goal:**
 - Share a real-world case to build foundational skills in HIV care
 - Designed for those newer to HIV practice
- **Focus:**
 - Where to start, what resources to use, how to find information



Pharmacy Consult:

Can you assess for
regimen simplification?

Patient VS

- 53 year old, cisgender female
- HIV History → HIV+ 1996 (life insurance required test)
- Allergies → Nevirapine (Stevens-Johnson Syndrome)
- Drug Coverage → private plan





Medications

- progesterone 100mg daily
- estradiol 1mg daily
- Genvoya (FTC/TAF/EVG/cobi)
one tablet daily
- darunavir 800mg daily
- recently stopped rosuvastatin
10mg daily



The Power of a Good ARV History

Viral load
undetectable since
March 2007

DATE	REGIMEN	COMMENTS
1996 - 1998	AZT + ddi	
1998	d4T + 3TC + S	Rosenthal Syndrome secondary to NVP
1998-2003?	d4T, 3TC, S	second son type
2003-2006?	drug holiday	otype
Dec 2006 - Nov 2007	TDF + 3TC + S	
Nov 2007 - Jul 2014	TDF/FTC + S	
Jul 2014 - Jun 2016	DTG + ABC/3TC + DRV 800mg + RTV	
Jun 2016 - Oct 2019	ABC/3TC/DTG + DRV 800mg + RTV	
Oct 2019 - Present	FTC/TAF/EVG/c + DRV 800mg	

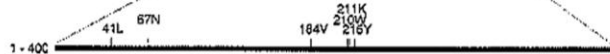


Lesson 2

Understanding Genotypes & Resistance

- Genotypic testing **detects resistance mutations** in relevant viral genes
- Requires viral load ≥ 250 copies/mL
- Mutation shorthand:
 - Example → **M184V**: methionine (M) → valine (V) at position 184
- Mutations help identify reduced ARV susceptibility
- Viral load sample → Public Health Ontario → BCCE for genotyping → Interpretation
- Watch the three part **CHAP 101 Learning Series: Tackling ART Resistance** by Linda Robinson

1-99

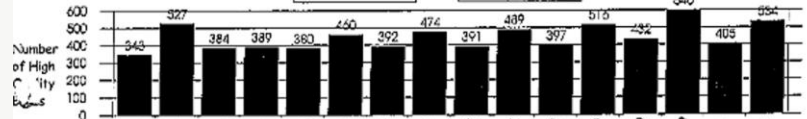
analysis ¹

change



K: NOTATIONS

T39A/T
M41L
D67N
E122K
Q123E
S162C
I178L
M184V
T200A
Q207E
L210W
R211K
L214F
T215V
P272A
R277K
I293V
Z329L
T376A
K395R
E399D



NRTI - 41L, 67N, 184V, 211K, 210W, 215Y

SUMMARY REPORT

DRUGS		FOLD CHANGE ¹	CUT-OFF ²		RESISTANCE ANALYSIS ³	CLINICAL NOTES <small>(see p2 for details)</small>
NRTI / NtRTI mutations: 211K						
NRTI/NRTI	Retrovir®	Zidovudine	1.0	1.9	14.4	MAXIMAL RESPONSE
	Epivir®	Lamivudine	0.9	1.1	3.7	MAXIMAL RESPONSE
	Videx®	Didanosine	0.8	1.3	3.0	MAXIMAL RESPONSE
	Hivid®	Zalcitabine	0.9		3.0	SUSCEPTIBLE
	Zenit®	Stavudine	0.8	1.1	2.2	MAXIMAL RESPONSE
	Ziagen®	Abacavir	0.7		2.1	SUSCEPTIBLE
	Emtriva®	Emtricitabine	0.8		3.7	SUSCEPTIBLE
	Viread®	Tenofovir DF	0.8	1.0	2.0	MAXIMAL RESPONSE

NNRTI mutations: 283wt/I						
NNRTI	Viramune®	Nevirapine	1.2		5.2	SUSCEPTIBLE
	Rescriptor®	Delavirdine	1.6		7.7	SUSCEPTIBLE
	Sustiva®, Stocrin®	Efavirenz	1.0		3.4	SUSCEPTIBLE

PI mutations: 77wt/I						
PI	Crixivan®	Indinavir	0.7	0.8	2.2	MAXIMAL RESPONSE
	Crixivan®, boosted	Indinavir/r	0.7	4.1	21.2	MAXIMAL RESPONSE
	Norvir®	Ritonavir	0.7		2.4	SUSCEPTIBLE
	Viracept®	Nelfinavir	0.9	1.0	1.5	MAXIMAL RESPONSE
	Invirase®	Saquinavir	0.6	0.7	1.0	MAXIMAL RESPONSE
	Invirase®, boosted	Saquinavir/r	0.6	1.1	12.0	MAXIMAL RESPONSE
	Agenerase®	Amprenavir	0.6	0.7	1.4	MAXIMAL RESPONSE
	Agenerase®, boosted	Amprenavir/r	0.6	0.9	6.5	MAXIMAL RESPONSE
	Lexiva®, Prezista®	Fosamprenavir	0.6		1.8	SUSCEPTIBLE
	Kaletra®	Lopinavir/r	0.6	1.0	0.9	MAXIMAL RESPONSE
	Royataz®	Atazanavir	0.7		2.0	SUSCEPTIBLE
	Aptivus®	Tipranavir	0.7		1.6	SUSCEPTIBLE

1. Predicted fold change in 50% inhibitory concentration (IC₅₀) relative to susceptible reference virus. 2. Cut-off values for maximal and minimal clinical response (Clinical Cut-Off) as for normal resistance testing. 3. Resistance Analysis based on the magnitude of the Fold Change relative to the Clinical or the Biological Cut-Offs. See page 3 for definitions.

September 27, 2006

NRTI - 211K

NNRTI - 283wt/I

PI - 77wt/I

Stanford Drug Resistance Database



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

VISTAS PROGRAM

ABOUT HIVDB

SUPPORT HIVDB!



**HIVDB Algorithm
Version 9.8**

Jan 05, 2025



Sierra 3.5.3

[release notes](#) / [web service](#)

Jan 05, 2025

HIV Drug Resistance Tutorials

[NRTI](#) / [NNRTI](#) / [PI](#) / [INSTI](#) / [HIVDR Interpretation Program](#)

Jun 02, 2024

**HIVDB Viral Sequence and
Treatment Submission
(VISTAS) Program**


HIV, HBV, HCV Genbank submission tool

Jun 28, 2024

GenBank2PubMed

Connect GenBank virus sequences to publications

May 07, 2025



**Calibrated
Population
Resistance**



INTERACTIVE MAP

HIVDB released on Sep 26, 2025

Query / Download



Genotype-treatment

[ARV selection data](#) comprising 236,806 protease, 251,203 RT, 40,313 integrase and 25,362 capsid HIV-1 virus sequences from 279,615 persons; 1,091 protease, 898 RT and 358 integrase HIV-2 virus sequences from 1,153 persons. [In vitro selection data](#) includes 1,111 HIV-1 in vitro selection data of PR, RT and IN.



Genotype-phenotype

[Drug susceptibility data](#) comprising 30,676 PI, 24,762 NRTI, 15,180 NNRTI and 5,644 INI susceptibility results from HIV-1 virus isolates



Genotype-clinical

[Clinical outcome data](#) comprising genotype, treatments, plasma HIV-1 RNA levels and CD4 counts from

HIVdb Program

**Drug Resistance Summaries
(Download PDF)**

[PIs](#) [NRTIs](#) [NNRTIs](#) [INSTIs](#) [CAIs](#)

HIV Drug Resistance Tutorials

[NRTIs](#) [NNRTIs](#) [PIs](#) [INSTIs](#)

[HIVDR Interpretation Program](#)

Questions or suggestion are welcomed:
hivdbteam@lists.stanford.edu

HIVdb Program: Mutations Analysis

HIVdb accepts user-submitted protease, RT, and integrase sequences or mutations and returns inferred levels of resistance to the most commonly used protease, nucleoside, non-nucleoside, and integrase inhibitors. Its purpose is educational and as such it provides extensive comments and a highly transparent scoring system that is hyperlinked to data in the HIV Drug Resistance Database. A detailed description of the program as well as all updates is in the [Release Notes](#). A [web service](#) has been created to allow users to access HIVdb programmatically.

New: this program is now available for analyzing SARS-CoV-2 mutations, FASTA, and FASTQ (NGS) sequences.

Protease, RT, and integrase mutations can be entered using either the text box or auto-suggestion boxes. To use the text box, type each mutation separated by one or more spaces. The consensus wildtype and separating commas are optional. If there is a mixture of more than one amino acid at a position, write both amino acids (an intervening slash is optional). Insertions should be indicated by "Insertion" and deletions by "Deletion".

Drug display options

By default, results will be shown for checked ARVs. Use checkboxes for additional ARVs. ([select all](#))

NRTI: ☒ ABC ☒ AZT ☒ FTC ☒ 3TC ☒ TDF ☐ D4T ☐ DDV

INSTI: ☒ BIC ☒ CAB ☒ DTG ☒ EVG ☒ RAL

NNRTI: ☒ DOR ☒ EFV ☒ ETR ☒ NVP ☒ RPV ☐ DPV

PI: ☒ ATV/r ☒ DRV/r ☒ LPV/r ☐ FPV/r ☐ IDV/r ☐ NFV ☐ SQV/r ☐ TPV/r

Input mutations Input sequences Input sequence reads

Reverse Transcriptase

M41L x D67N x M184V x R211K x L210W x T215Y x L283I x Enter/paste 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							
...							

Protease

V77I x Enter/paste 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

Enter/paste mutations

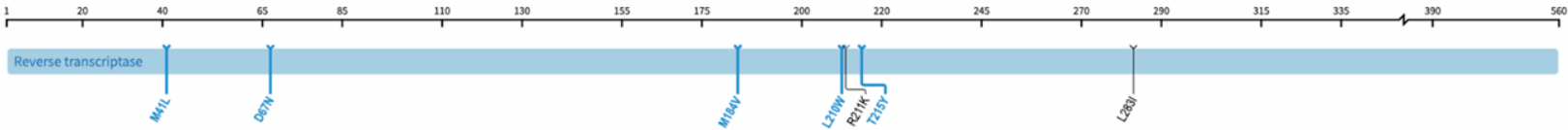
51	66	74	92	95	97	114	118
...
121	128	138	140	143	145	146	147
...
148	149	151	153	155		163	230
...
232	263						
...	...						

☒ Save input mutations in my browser for future use

Reset

Analyze

Reverse transcriptase (RT)



There are no known mutation quality issues.

Drug resistance interpretation: PR

HIVDB 9.8 (2025-01-05)

PI Major Mutations: None
PI Accessory Mutations: None
PR Other Mutations: V77I

Protease Inhibitors

atazanavir/r (ATV/r) Susceptible
darunavir/r (DRV/r) Susceptible
lopinavir/r (LPV/r) Susceptible

Mutation scoring: PR

HIVDB 9.8 (2025-01-05)

No drug resistance mutations were found for PI.

Drug resistance interpretation: RT

HIVDB 9.8 (2025-01-05)

NRTI Mutations: **M41L • D67N • M184V • L210W • T215Y**
NNRTI Mutations: None
RT Other Mutations: R211K • L283I

Nucleoside Reverse Transcriptase Inhibitors

abacavir (ABC) High-Level Resistance
zidovudine (AZT) High-Level Resistance
emtricitabine (FTC) High-Level Resistance
lamivudine (3TC) High-Level Resistance
tenofovir (TDF) High-Level Resistance

Non-nucleoside Reverse Transcriptase Inhibitors

doravirine (DOR) Susceptible
efavirenz (EFV) Susceptible
etravirine (ETR) Susceptible
nevirapine (NVP) Susceptible
rilpivirine (RPV) Susceptible

RT comments

NRTI

- **M41L** is a TAM that usually occurs with T215Y. In combination, **M41L** plus T215Y confer intermediate / high-level resistance to AZT and d4T and contribute to reduced ddI, ABC and TDF susceptibility.
- **D67N** is a non-polymorphic TAM associated with low-level resistance to AZT.
- **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.
- **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.
- **T215Y/F** are TAMs that causes intermediate/high-level resistance to AZT and potentially low-level resistance to ABC and TDF.

Assessment

- Stanford resistance profile shows **high-level resistance** to the **entire NRTI** class
- **Severe allergy to nevirapine** (SJS) → concern for cross-sensitivity with NNRTIs
- No evidence of NNRTI re-exposure since the reaction → avoid unless no alternatives
- Can we simplify??

Lesson 3

Use Your Community

- CHAP google group → poll the country!
- CHAP observership opportunities
- Expert and experienced colleagues
- Medical liaisons



Alternatives

2

B/F/TAF alone

- Use with TAF mutations is off-label
- Canadian study switching to B/F/TAF in the presence of 1-8 NRTI mutations → 98% maintained suppression
 - 2 had same 5 NRTI RAMs; 9 were on same baseline regimen
- Trial with close VL monitoring
- If virologic rebound occurs → add DRV/c

1

B/F/TAF + DRV/c

- Remains PI-based, two-tablet regimen
- Modernizes integrase inhibitor from EVG → BIC
- Simple, safe, and effective switch

3

DTG/3TC + DRV/c

- PI-based, two-tablet regimen
- Modernizes integrase inhibitor from EVG → DTG
- HBV immunity confirmed

Outcome

- VS met with the physician to discuss simplifying her regimen to B/F/TAF alone
- **VS declined any change!** She reported feeling great with her current regimen and prefers to remain on current therapy due to concerns about tolerating switches
- Valuable learning gained through the case despite no change





Thank you!

Do you have
any comments
or questions?