

Virologic Failure: Resistance is Not Futile!

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- I have no financial disclosures

Objectives

- Identify factors that influence drug resistance.
- Understand the methods available for measuring HIV drug resistance.
- Identify strategies for responding to detection of moderate to high levels of drug resistant HIV.

So You've Started Someone on Antiretroviral Therapy

- Can be intimidating
- Is it working? How do you know?
- If it is not working, what can you do about it?



What is antiretroviral therapy (ART) resistance?

- Reduced susceptibility of Human Immunodeficiency Virus (HIV) to a specific antiviral drug
 - Drug less effective in replication blockade
 - > 200 copies/mL of blood (DHHS guidelines, 2016)
- Caused by mutations in viral genome
- May be:
 - Agent specific
 - Cross-resistance (multiple drugs)
- Magnitude of resistance variable by mutation and number

Approach to Potential Resistance

- Is it really resistance?
- Selecting a resistance test
- Considering the entire library of resistance
- Address root cause
- Use resources:
 - Colleagues
 - Resistance Databases (Stanford)
 - National Clinical Consultation Center



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



Calibrated Population Resistance

INTERACTIVE MAP

Surveillance Mutations

Point-of-Care / Essential Mutations

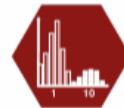
Query Pages



Genotype-treatment

Retrieve sequences (and/or mutations) from persons receiving selected HIV drugs

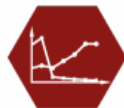
Retrieve sequences and treatments from viruses with specific mutations



Genotype-phenotype

Retrieve drug susceptibility data for isolates with selected mutations

Download genotype-phenotype research datasets



Genotype-clinical

Summaries of genotype-clinical outcome studies

Genotype-clinical outcome datasets (download)



References

Published drug resistance studies in HIVDB

Published studies by Stanford database group

New Submissions

[View All](#)

HIVdb Program

Drug Resistance Summaries (Download PDF)

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

HIVseq Program

HIValg Program

HIV-1 Genetic Variability for Drug Resistance

<https://hivdb.stanford.edu/>

Making a “Master” Genotype

Drug display options

By default, results will be shown for checked ARVs. Use checkboxes for additional ARVs. (select all ARVs, revert to default)

NRTI: ABC AZT FTC 3TC TDF D4T DDI NNRTI: EFV ETR NVP RPV

INSTI: DTG EVG RAL PI: ATV/r DRV/r LPV/r FPV/r IDV/r NFV SQV/r TPV/r

Input mutations Input sequences

Reverse Transcriptase **Protease** **Integrase**

Input mutation(s)

Select mutations:

40	41	44	62
---	---	---	---
65	67	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	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	121
---	---	---	---
128	138	140	143
---	---	---	---
145	146	147	148
---	---	---	---
151	153	155	157
---	---	---	---
163	230	263	
---	---	---	

Reset Analyze

34-yo-male with HIV initiated on antiretroviral therapy 3 years ago with consistently suppressed virus presents for follow up. Previous CD4 1100 cells/mm³ 6 months ago. Labs today: CD4 1264 cells/mm³ and viral load 65 copies/mL.

What is going on here?

What is our next step?

Just A Little Bit of Virus?

The Blip

- An episode of low-level viremia that is preceded and followed by suppression below the quantification limit of an assay WITHOUT a change in therapy
- Less than 200 copies/ML
- Often unclear cause
- Repeat viral load

Persistent Low Level Viremia (PLLV)

- A viral load <200 copies/mL but not “undetectable”
- Action controversial
- Follow closely
- Consider Archive Genotype*

* We'll come back to this

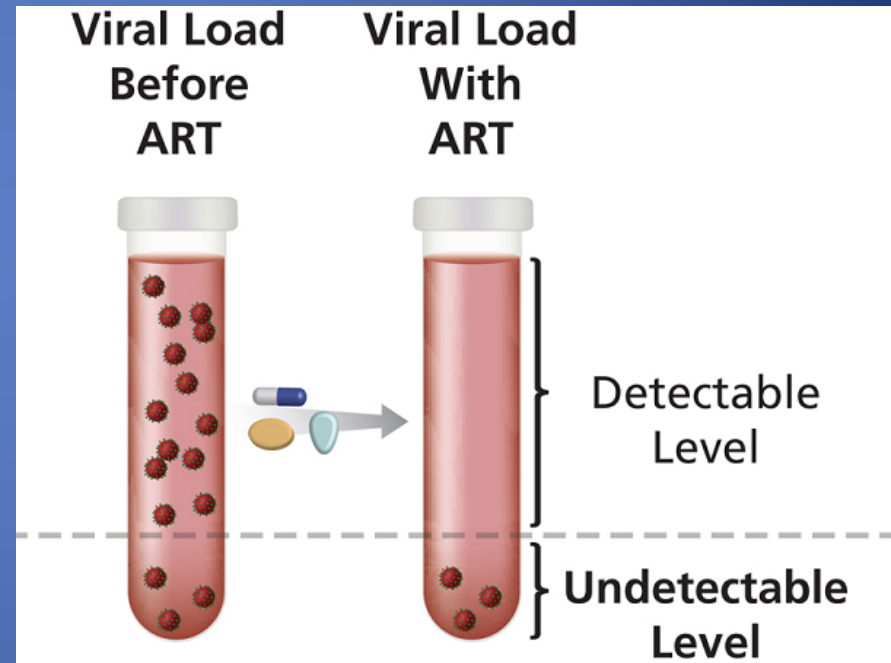
56-yo-male with HIV diagnosed 7 years ago. He has been on antiretroviral therapy for >5 years consistently with a viral load < 20 copies/mL. He has recently signed up for your health system patient access portal. He calls you to discuss the following result:

```
SPECIMEN DESCRIPTION:          (PURPLE TOP) EDTA WHOLE BLOOD/PLASMA
HIV QUANTITATIVE VIRAL LOAD RESULT:  HIV-1 RNA Detected
Log base ten value = <1.3
Copies RNA per mL = <20
```

How should we address his concerns?

“Detectable but not Quantifiable”

- Usually viral load of <math><20-50</math> copies/mL
- Moving target
- Most would not change therapy
- Future standards?



Resistance or Not?

- Baseline resistance testing
- Repeat an abnormal result (ie. viral load) prior to making therapy changes
- If established that resistance present, try to determine cause:
 - Adherence
 - Absorption
 - Drug interactions

Testing for Resistance

- When
 - Entrance to care
 - Initiation of therapy
 - In virologic failure:
 - Failure with HIV RNA >1000 copies/mL
 - Failure with HIV RNA 500 -1000 copies/mL
 - Testing should be attempted but may not be successful
 - Suboptimal viral load reduction
 - On therapy if able

DHHS Guidelines for Use of Antiretroviral Agents in HIV-1 Infected Adults and Adolescents, 2016

Resistance Testing: Genotype vs. Phenotype

Genotype

- Advantages
 - Availability
 - Less technically demanding
 - Mutations likely precede phenotypic resistance
 - Less expensive

Phenotype

- Advantages
 - Direct measure of susceptibility
 - More familiar reporting results



Genotype vs. Phenotype

Limitations

Genotype

- Limitations
 - Indirect measure
 - May not correlate to phenotype
 - Expert interpretation required
 - Insensitive for minor mutations
 - Relies upon known mutations
 - May need separate test for integrase inhibitor panel

Phenotype

- Limitations
 - Limited availability
 - Technically demanding
 - Insensitive for detecting minor mutations
 - Clinically significant breakpoints not defined
 - Costly

Archive Genotype

- Introduced 2014
- Designed to measure resistance in virologically suppressed patients
- Facilitate therapy switch in otherwise suppressed patients.

Archived vs Plasma Virus

“The PBMC-derived HIV-1 DNA and circulating HIV-1 RNA represent two different viral compartments in the same individual.”*

HIV-1 DNA (“Archived”)

- Virus transmitted and archived at the time of acute infection
- Mutations acquired during the course of the patient’s ARV treatment

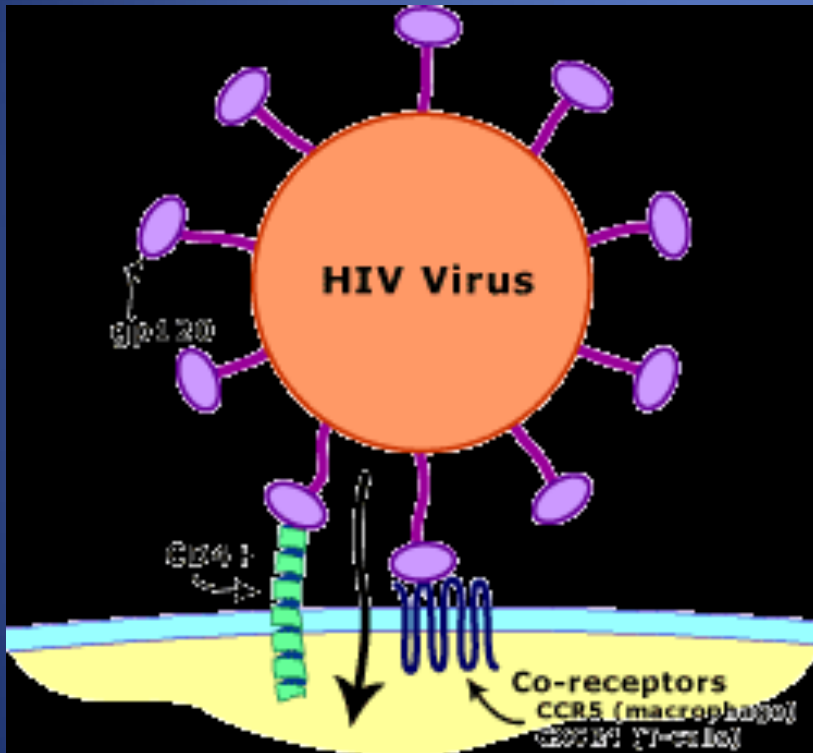
HIV-1 RNA (“Plasma”)

- The actively replicating virus from productively infected cells.
- The most current form of the virus.

* VanDamme AM, et al. *AIDS Rev.* 2011; 13:77-108.

Co-Receptor Tropism Testing

- Perform when:
 - Considering use of CCR5 co-receptor antagonist
 - Virologic failure on CCR5 antagonist



<http://genetics.thetech.org>



aidsinfo.nih.gov

Mutation Reporting Convention

Wild-type Amino Acid Designation

Substituted Amino Acid Designation



K103N



Position of Codon in Genome

Lysine (K) replaced by asparagine (N)

Mutations Behave Like:

Light Switch



westsidewholesale.com

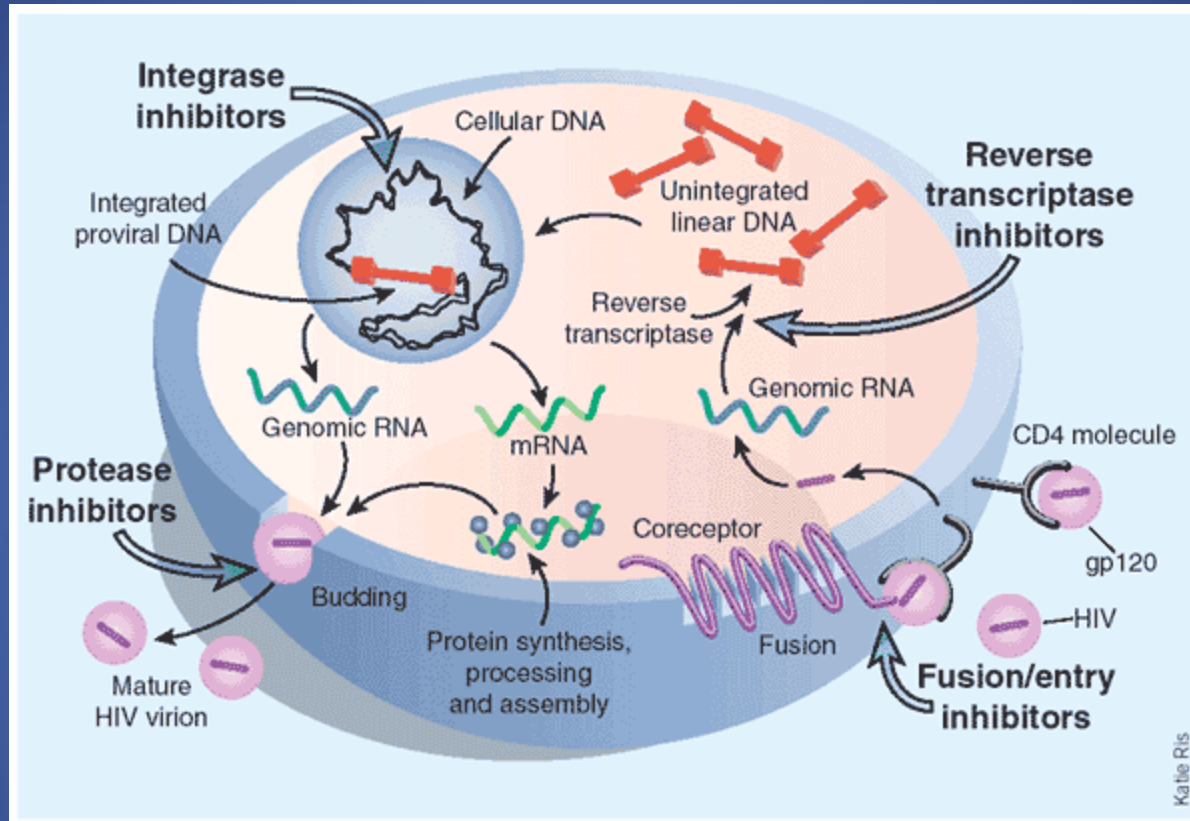
Dimmer Switch



Adapted from David Spach, MD

Drug Categories

NRTI	NNRTI	Protease Inh	Integrase Inh	Entry Inh
Tenofovir (TDF/TAF)	Efavirenz (EFV)	Darunavir (DRV)	Raltegravir (RAL)	Maraviroc (MVC)
Emtricitabine (FTC)	Rilpivirine (RPV)	Atazanavir (ATV)	Elvitegravir (EVG)	Enfuvirtide (ENF)
Abacavir (ABC)	Etravirine (ETR)	Ritonavir (/r)	Dolutegravir (DTG)	
Lamivudine (3TC)	Nevirapine (NVP)	Lopinavir (LPV)		
Zidovudine (AZT)	Delavirdine (DLV)	Fosamprenavir (FPV)		
Stavudine (D4T)		Tipranavir (TPV)		
Didanosine (DDI)		Saquinavir (SQV)		
		Nelfinavir (NFV)		
		Indinavir (IDV)		



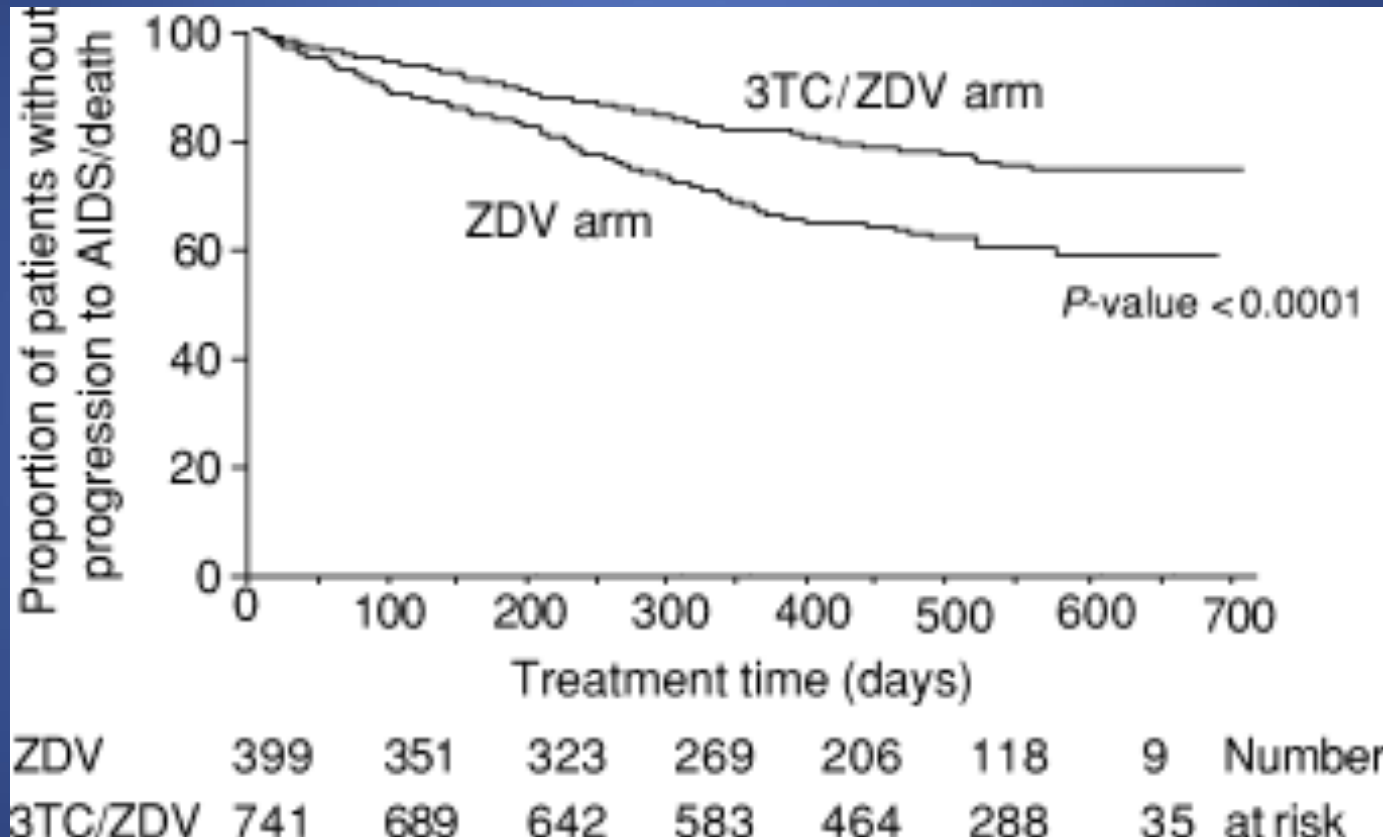
35-yo-female with HIV/AIDS that returns to clinic for follow up. She misses 1 or 2 out of every 3 appointments scheduled. Currently on TDF/FTC/DRV/r. Viral load today is 10,000. HLAB5701 is negative. A genotype reveals the following mutation: M184I

Assuming good adherence to regimen, what changes would you suggest?

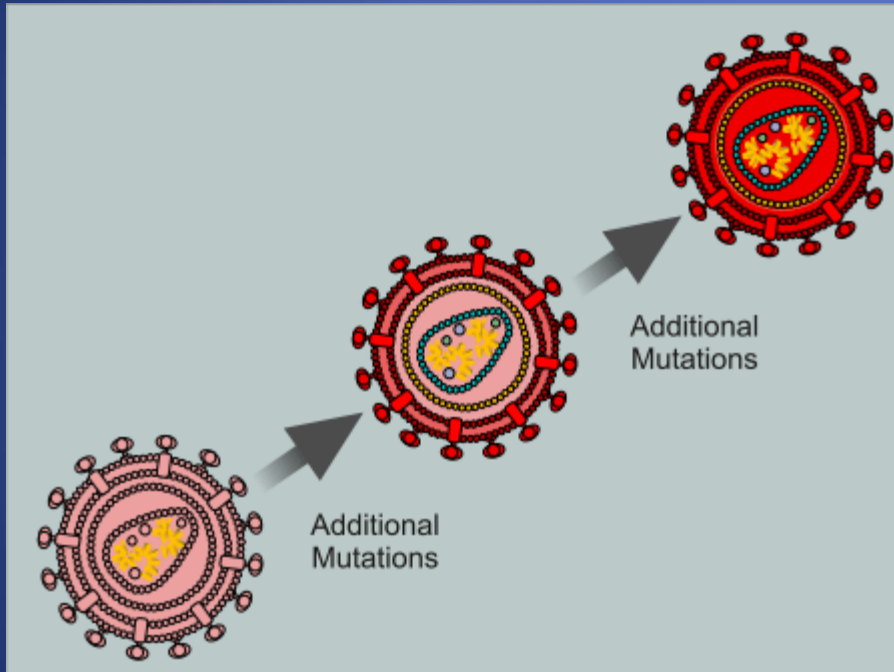
184V

- M184V most common nucleoside reverse transcriptase inhibitor (NRTI) mutation
- Selected for by 3TC, FTC
 - Reduces susceptibility to these drugs x100 fold
 - Slight decrease in ABC activity
 - Increased activity: AZT, TDF (probably TAF), d4T
- Decreased viral fitness
 - virus “expends more energy to maintain”
 - Reversion to wild type without “pressure”

The Impact of the M184V Substitution in HIV-1 Reverse Transcriptase on Treatment Response



M184I



fhmb.tqn.com

- Precursor to 184V
- Same impact
 - Resistance profile
 - Decreased fitness
- 184I +138K-> increased RPV resistance

35-yo-female with HIV/AIDS that returns to clinic for follow up. She misses 1 or 2 out of every 3 appointments scheduled. Currently on TDF/FTC/DRV/r. Viral load today is 10,000. HLAB5701 is negative. A genotype reveals the following mutation: M184I

What drugs are active? Which are inactive?

What changes should we make?

K65R

- Selected for by: TDF, ABC, d4T, ddI, rarely 3TC
- Reduces susceptibility:
 - 2 fold: TDF, ABC, ddI
 - 5-10 fold: 3TC, FTC
 - Can co-exist with 184V/I
- Increases susceptibility:
 - AZT (except in presence of 151M)

A 57-yr-old male with HIV/AIDS for 10 years presents to clinic to establish care. He has had a long history of antiretroviral treatment in the past and cannot recall all of his previous medications. He is not currently on antiretroviral therapy. A HIV genotype reveals the following mutations: 41L, 103N, 181C, 215Y, 184V

What therapy should we consider?

Thymidine Analog Mutations (TAMS)

- M41L
- D67N
- K70R
- L210W
- T215Y
- Selected by: AZT, d4T
- “Dimmer switch”
- Cumulative effect
- If multiple TAMS
assume M184V!
- Reduced susceptibility:
AZT, d4T, ABC, ddI, TDF
- Not all are created
equal

A 57-yr-old male with HIV/AIDS for 10 years presents to clinic to establish care. He has had a long history of antiretroviral treatment in the past and cannot recall all of his previous medications. He is not currently on antiretroviral therapy. A HIV genotype reveals the following mutations: 41L, 103N, 181C, 215Y, 184V

What therapy should we consider?

NRTI Resistance to Know

- M184V/I
 - Common
 - May “hide” in a wild type
- TAMSA
 - Cumulative
 - “Dimmer switch”
- K65R

50-yo-male recently diagnosed with HIV here to establish care. No prior history of antiretroviral therapy in the past. He is interested in starting medications. On review of genotype you note the following mutations: K103N. Initial viral load 8,000 copies/mL. CD4 >500 cells/mm³

What drug(s) should we avoid?

What other mutations might be present?

Transmitted Resistance

- “Resistance testing at entry to care regardless of whether antiretroviral therapy (ART) will be initiated immediately.”
 - DHHS Treatment Guidelines (2016)
- Frequency of ≥ 1 major resistance mutations in treatment naïve patients (North America): 10-24%
- Durable
- Measure of the dominant species present

K103N

- Most common non-nucleoside reverse transcriptase inhibitor (NNRTI) mutation
- “Light Switch”
- Selected by:
 - NVP, EFV
- Reduces susceptibility:
 - NVP (50X), EFV (20x)
- When coexists with M184V:
 - 2/3 patients
 - If keep of EFV can lead to more mutations
 - No set guidelines for regimen in setting; would favor integrase inhibitor

Your new patient sees a drug chart on the wall and is intrigued by the small pill size of TAF/FTC/RPV. After discussing potential side effects, drug-drug interactions, dosing strategy, you both agree to start this medication. He is lost to follow up. He returns to care 9 months later with reported “good” adherence to medications. Labs as follows:

HIV viral RNA: 4500 copies/mL

Genotype: K103N, E138K, M184I

What now?

E138K

- NNRTI mutation selected by RVP
 - Decreased activity of RVP
 - Decreased activity of ETV compared with K103N
- In combination with M184I
 - Can cause failure on RPV-containing regimen
 - Significant activity decrease in ETV

Etravirine (ETV)

- Consider in salvage
- “Dimmer drug”
- Multiple mutations to completely eliminate activity; determined by scoring system
- Use mutation calculator to assist in assessing activity

Adapted from David Spach, MD

Hivdb.stanford.edu;

NNRTI Resistance to Know

- K103N
 - Common
 - May have masked M184V
- E138K
 - RVP and ETV to some extent
 - Enhancement of M184I
- Etravirine scoring

Protease Inhibitor Resistance

- Class typically with high ceiling for resistance
- “Dimmer Switch” mutations
- Complex
- Use resistance data base



45-yo-female with HIV on TDF/FTC, RAL presents for follow up. Previously well-controlled, but admits to only taking 3-4 doses/week since last visit 4 months ago. Current viral load 1644 copies/mL.

Baseline genotype: no mutations.

What has happened here?

Integrase Inhibitor Resistance

- Newest class; evolving
- Often not evaluated on standard genotype
- Variable ceiling for resistance



Raltegravir



Elvitegravir



Dolutegravir

Integrase Inhibitor Mutations

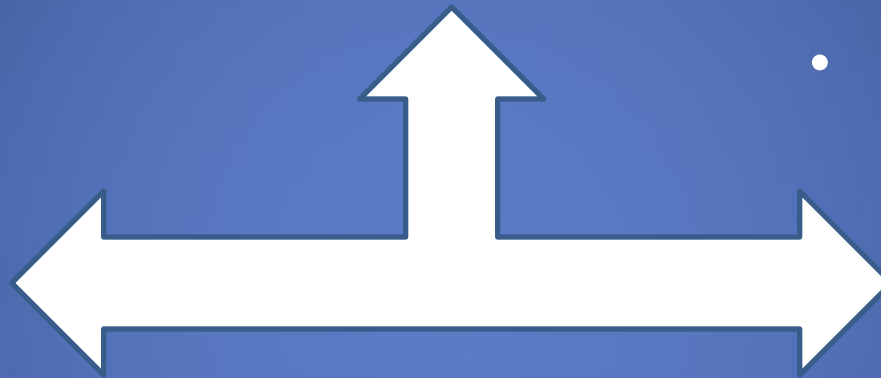
Cross Resistance

Raltegravir

- Low barrier
 - Y143
 - Q148
 - N155

Elvitegravir

- Medium Barrier
 - E92
 - Q148
 - N155



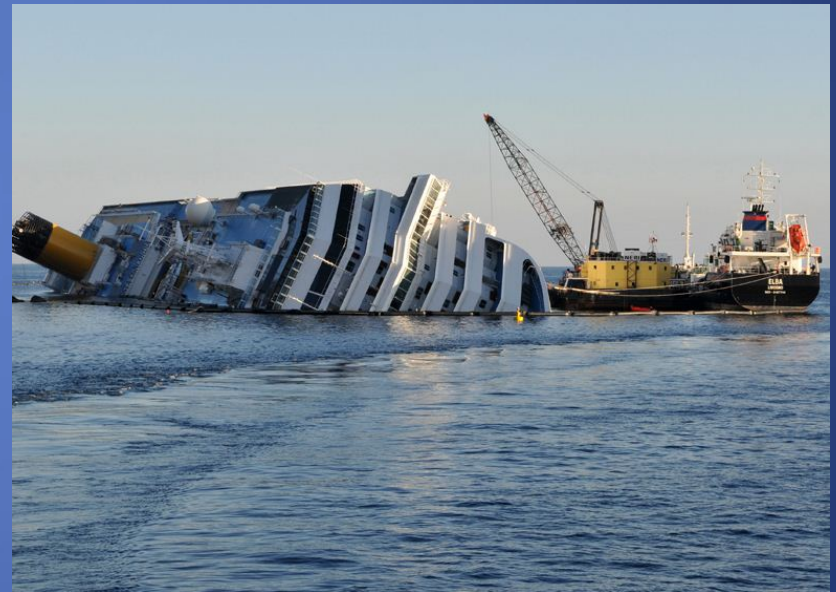
- G140+ G148 = high degree of resistance
- Q148 pathway can lead to DOL resistance
- Avoid prolonged failure!

Dolutegravir

- No clear reports of dolutegravir resistance in setting of no prior integrase resistance
- “Dimmer” drug
- Once daily vs. Twice daily dosing
 - If 148 mutation present would use twice daily dosing—not clear in package insert (Viking-3 Study)

Deep Salvage

- Deep salvage = Deep breaths
- Genotypes are additive
- Try to add two potentially active drugs
- Think outside the box!
 - Atypical backbone
 - Experimental drugs
 - Use resources!



Resources

- National Clinical Consultation Center/Clinicians' Warmline (CDC)
 - M-F 9AM-8PM EST
 - 1-800-933-3413
- HIV Drug Resistance Database
 - Hivdb.stanford.edu