

# ARV Resistance and why it still matters



## Dr. Edmund Wilkins

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North Manchester General Hospital

# Resistance...



# Outline

- Theory
  - Relevance
  - How to manage without a resistance test
- 
- Please interact....



# What is resistance..



What happens..



# How does it occur?

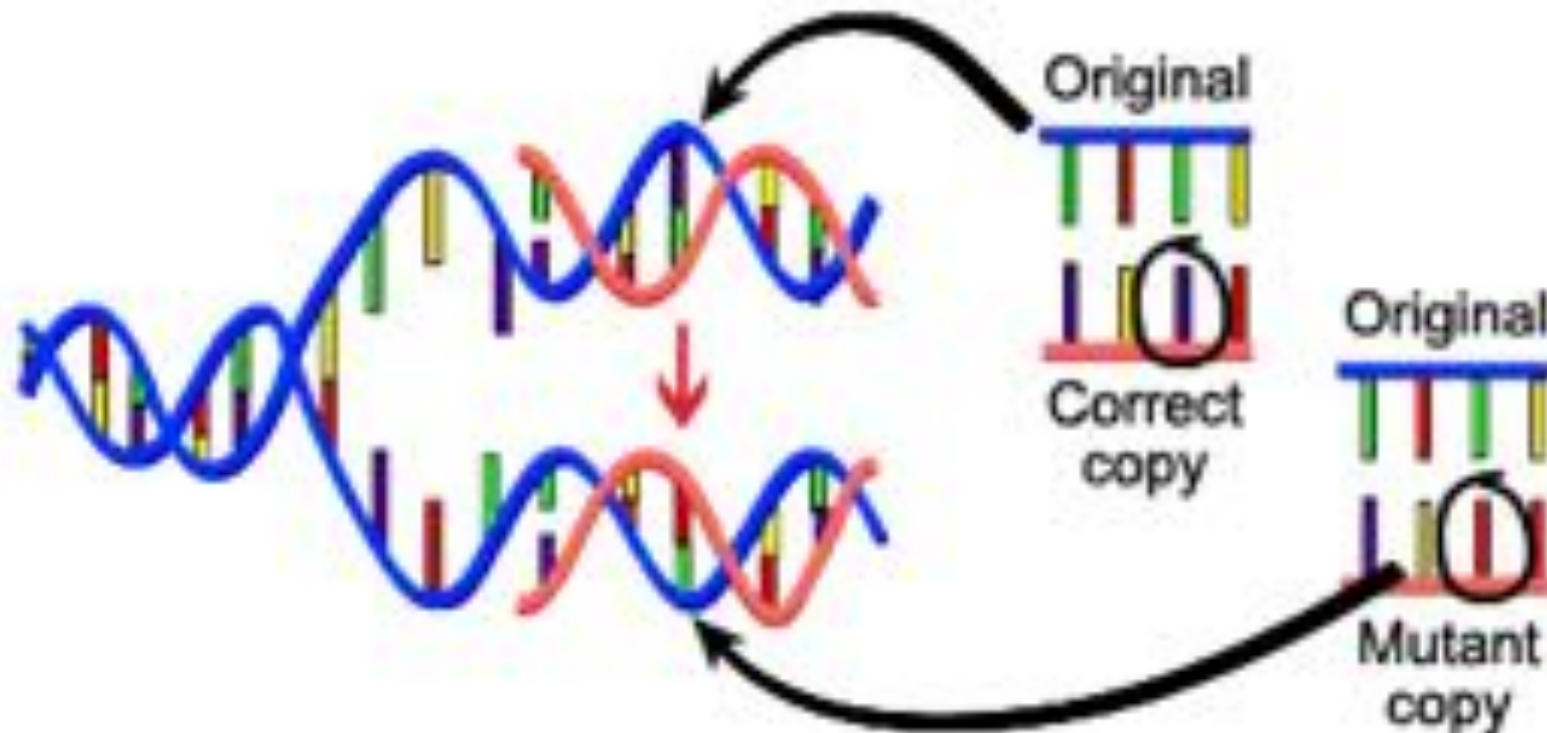
- **A mutation of the *viral genetic material***



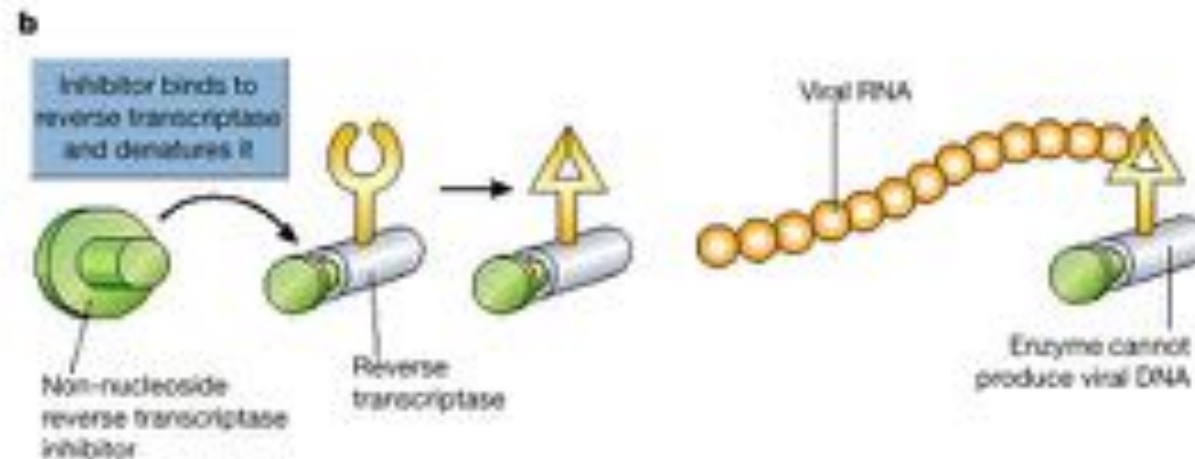
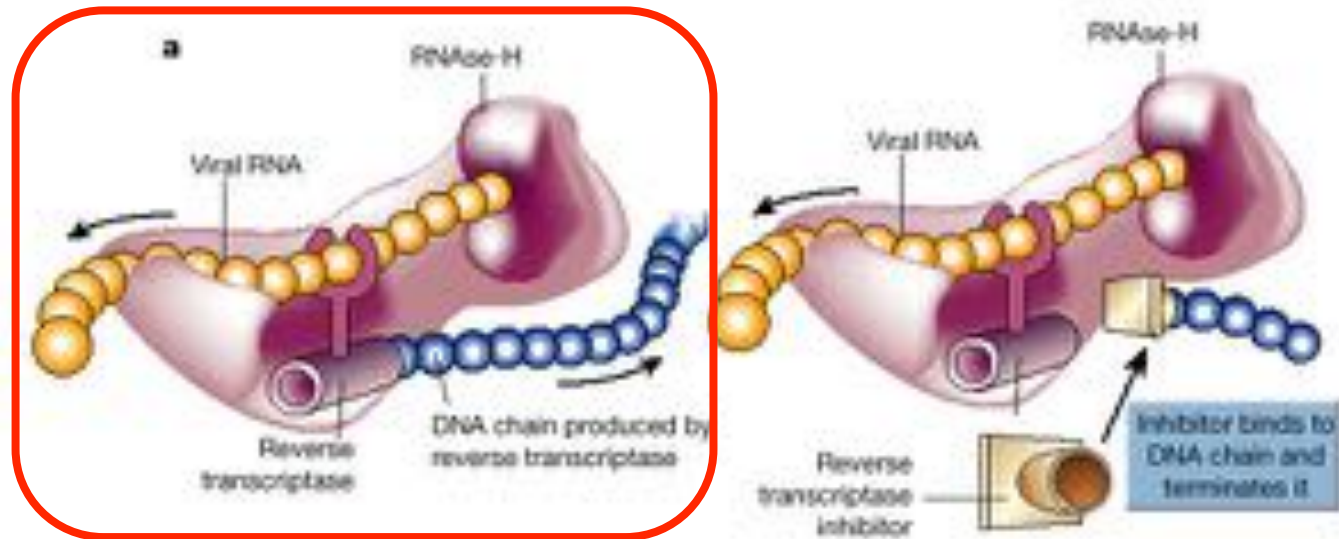


# And in HIV....

- **A mutation of the *viral genetic material*** that results in the drug no longer being able to block viral replication

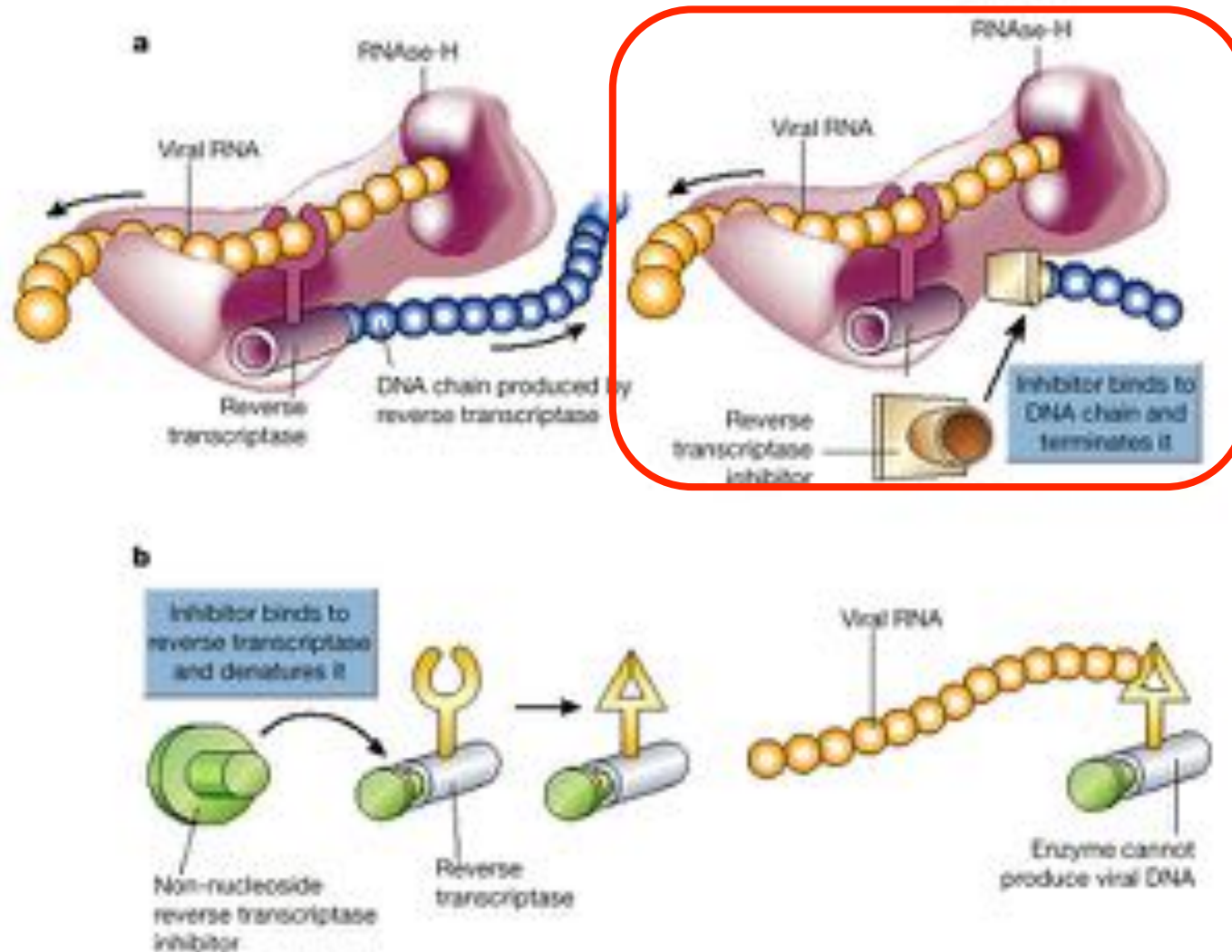


# What happens...

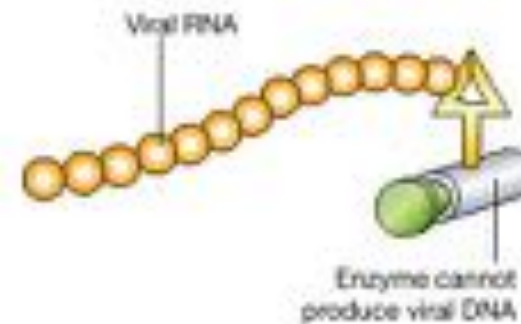
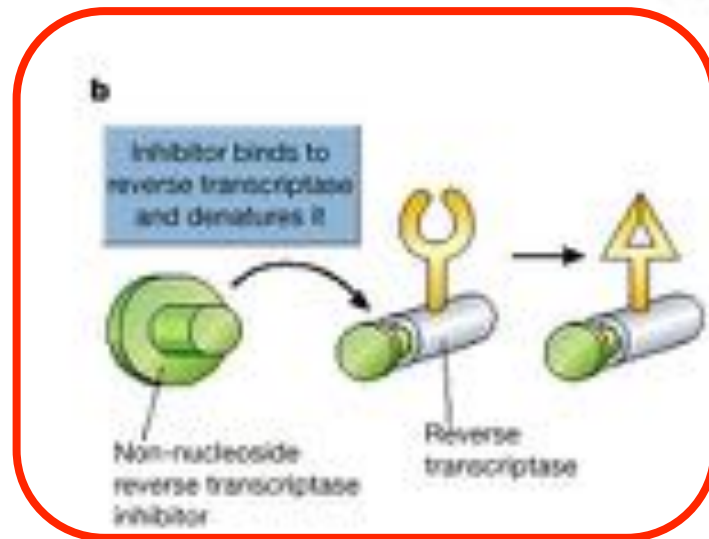
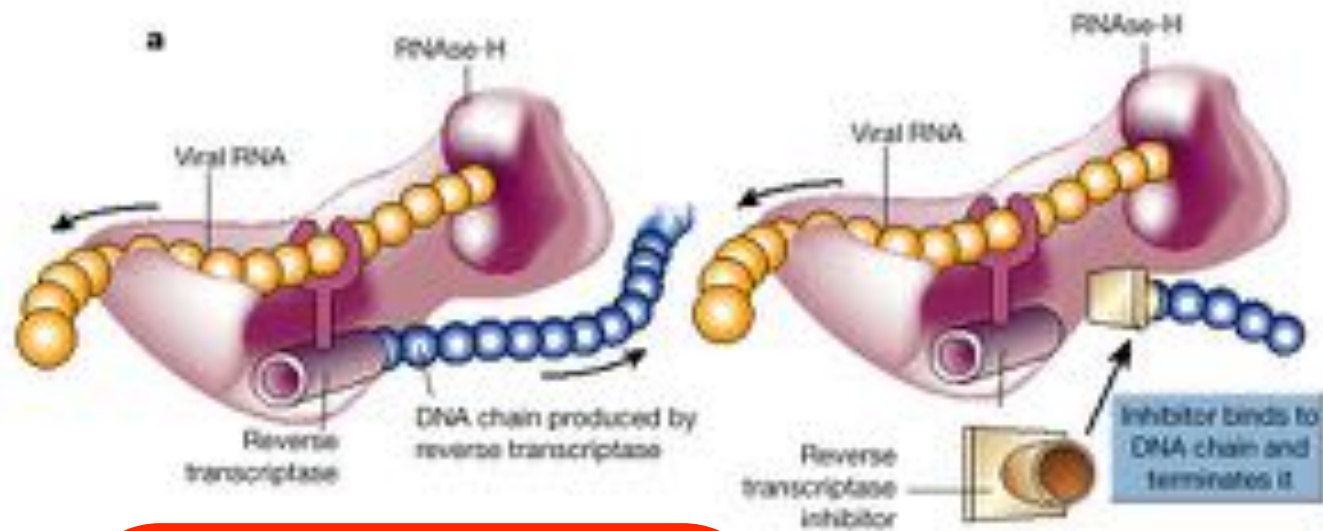




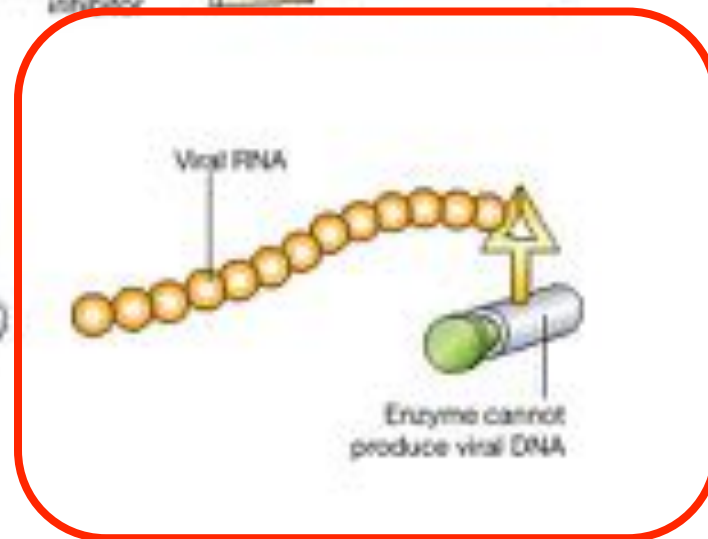
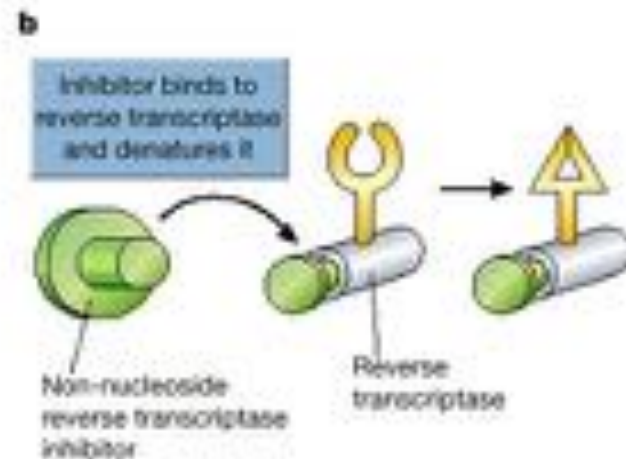
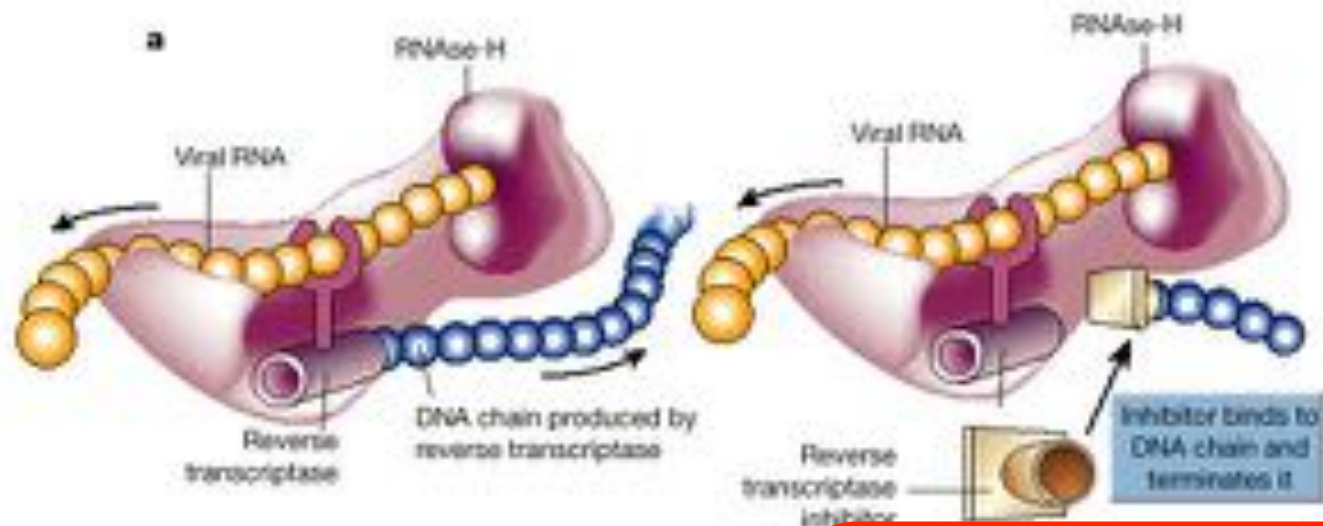
# What happens...



# What happens...



# What happens...



# Why does it occur?

- High mutation rate
  - HIV is 'poor' at replicating itself accurately
    - Many mistakes occur
  - Therefore lots of potential to develop resistance
    - In untreated patient: >1 billion viral particles made/day, with at least one mutation per 1000 viruses → 1-10 million mutations/day.
    - In a patient with a moderate viral load, every single mutation is possible in the HIV genome, every single day...
- Low barrier to resistance
  - It doesn't take many resistance mutations to knock out a drug
  - These mutations not 'lethal' or significantly hampering for the virus



# Why does it occur?

- Viral replication in presence of detectable drug(s)
  - Inadequate combination of drugs
    - **Not potent enough**

Resistance will develop with suboptimal treatment



# Inevitable consequence of pre-ART therapy

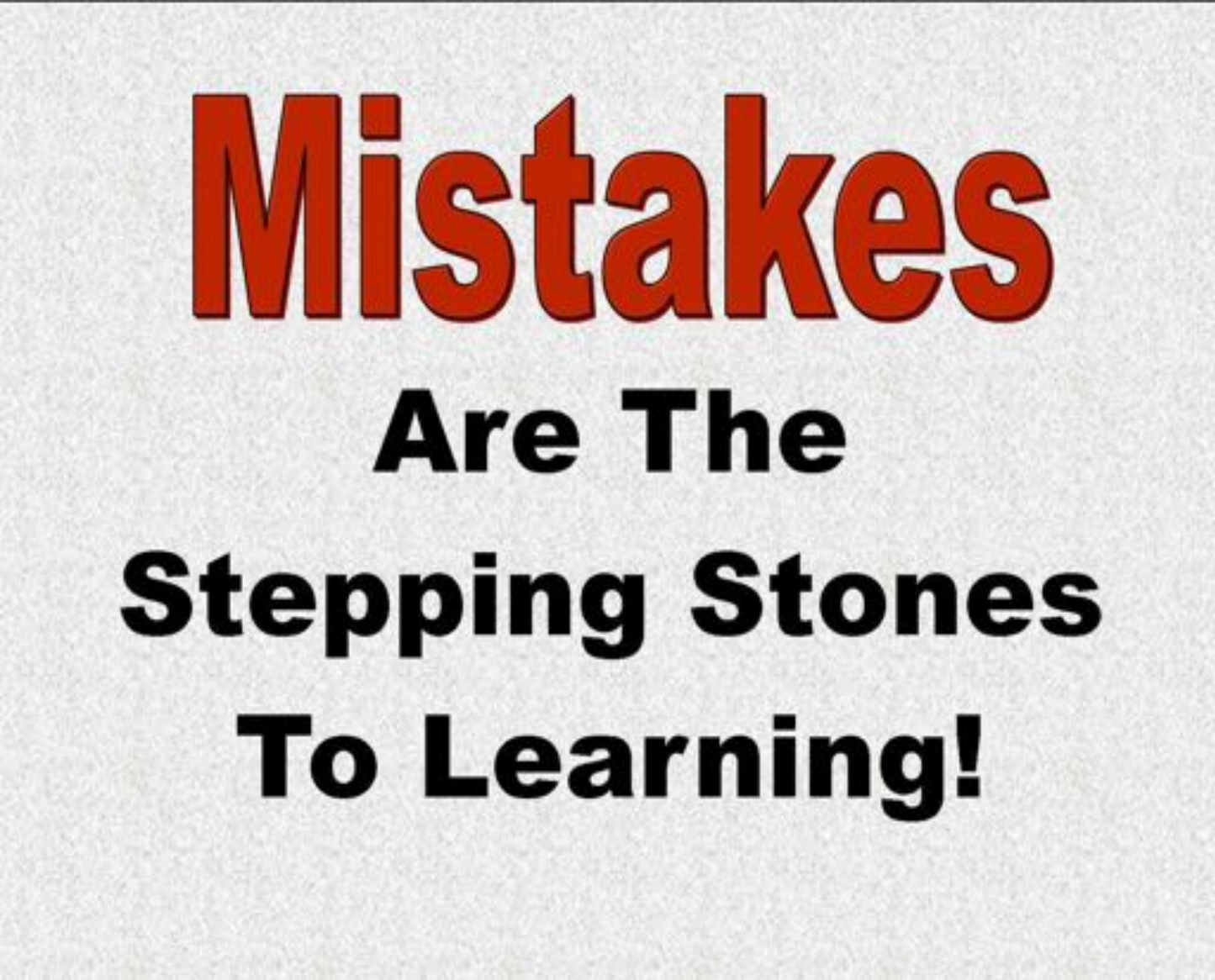


At a time when treatment was  
for survival





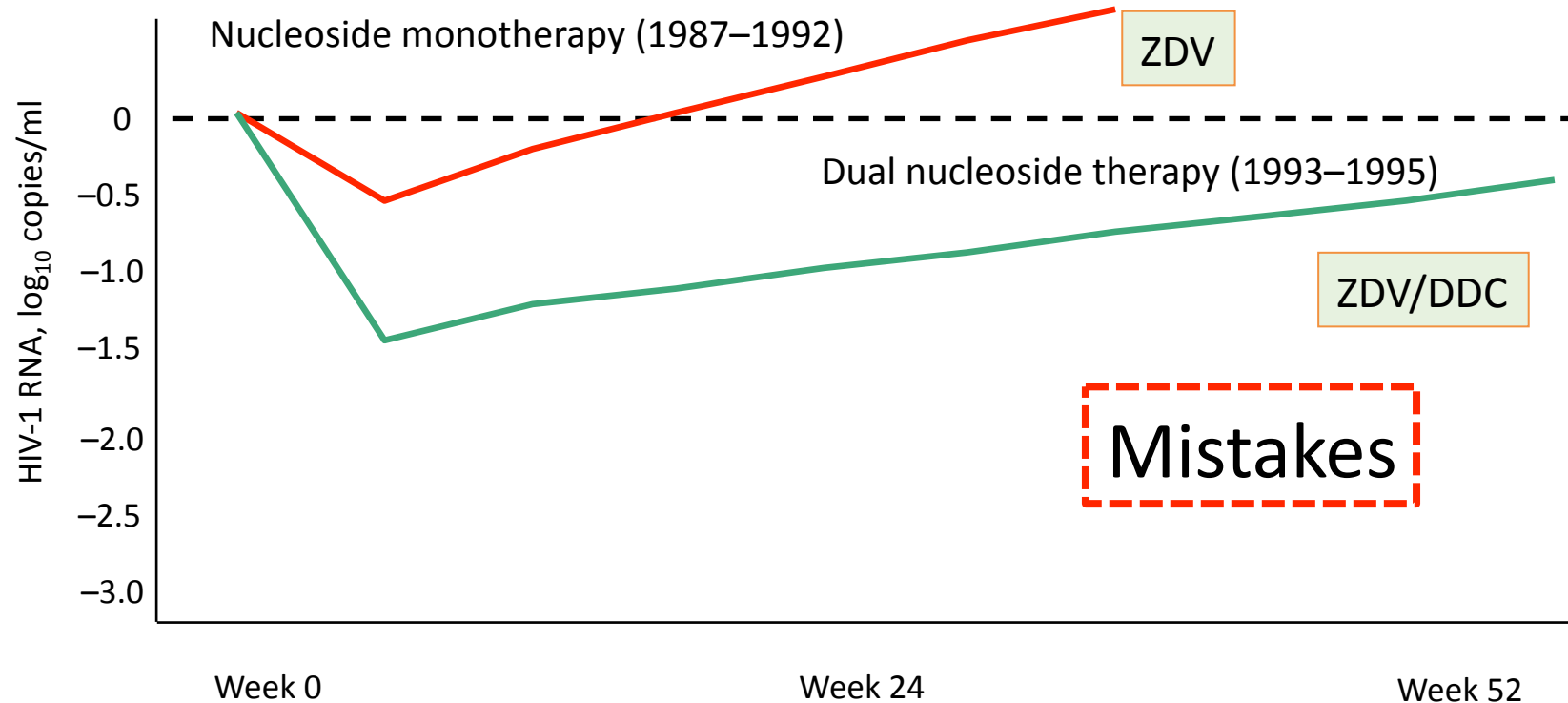
And mistakes were being  
made



**Mistakes**  
**Are The**  
**Stepping Stones**  
**To Learning!**

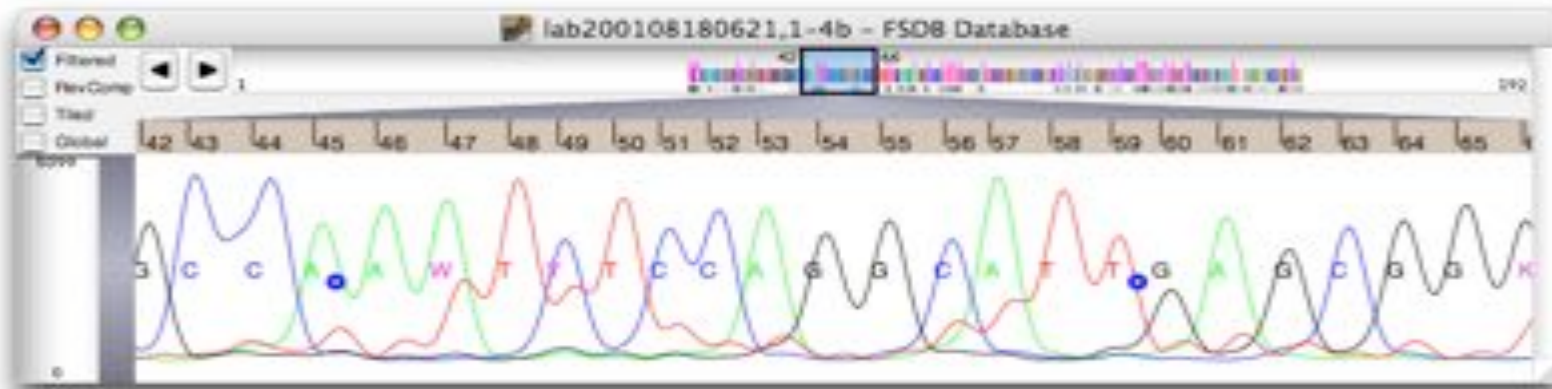
# 2NRTI therapy failed

Relative viral load suppression with mono- and combination therapies



# Resistance testing became available...

Nucleotide sequencing



**TRUGENE<sup>®</sup> HIV-1**  
**GuideLines<sup>™</sup> Rules 13.0**  
**RESISTANCE REPORT**

Sample ID: 180C060104  
 Patient ID: 1013dew  
 Patient Name: John Doe  
 Date Drawn: 20030606  
 Physician: Dr. Jane Doe  
 Institution: City Hospital  
 Report Date: 2007/11/12

State Central Labs  
 22 State St. Shelton, CT 06789

Bayer Reference Testing Laboratory  
 Example Report  
 725 Potter Street (APC3)  
 Berkeley, CA 94710  
 Tel: 800-434-2447  
 Fax: 510-705-5902

Resistance associated RT Mutations: L100I, K103N, T215S/Y

Nucleoside and Nucleotide RT Inhibitors	Resistance Interpretation
abacavir (ABC)	No Evidence of Resistance
didanosine (ddI)	No Evidence of Resistance
lamivudine (3TC)/emtricitabine (FTC)	No Evidence of Resistance
stavudine (d4T)	Resistance
tenofovir (TDF)	No Evidence of Resistance
zidovudine (AZT)	Resistance

NonNucleoside RT Inhibitors	Resistance Interpretation
efavirenz (EFV)	Resistance
nevirapine (NVP)	Resistance

Resistance associated PR Mutations: L19I, M46L\*, L63P, A71T

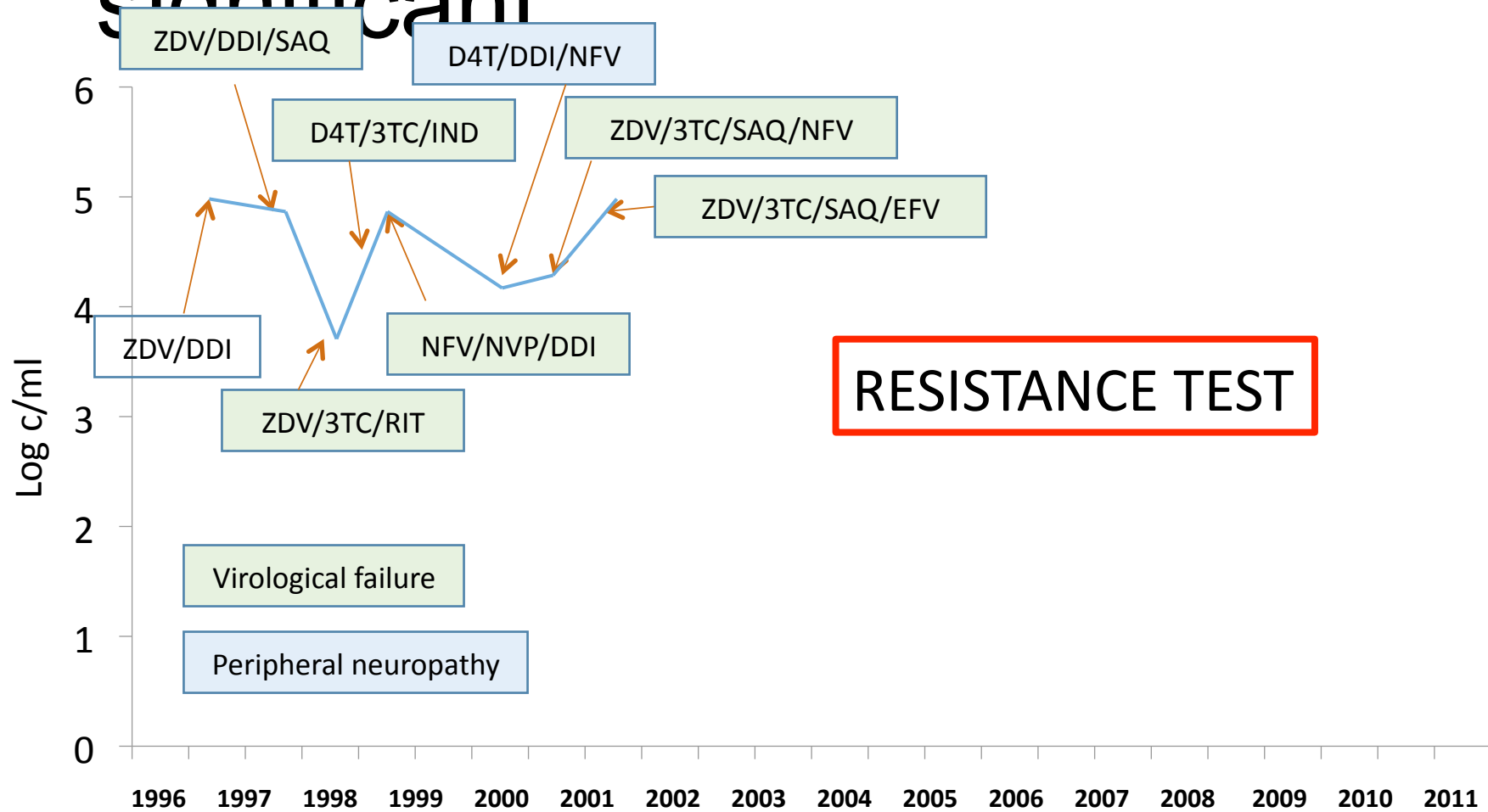
Protease Inhibitors	Resistance Interpretation
amprenavir (APV)/fosamprenavir (FPV)	Resistance
APV or FPV**	Resistance
atazanavir (ATV)	No Evidence of Resistance
ATV/r**	No Evidence of Resistance
darunavir + ritonavir (DRV/r)	No Evidence of Resistance
indinavir (IDV)	Resistance
IDV/r**	Possible Resistance
lopinavir + ritonavir (LPV/r)	No Evidence of Resistance
nelfinavir (NFV)	Possible Resistance
saqvinavir + ritonavir (SQV/r)	No Evidence of Resistance
tipranavir + ritonavir (TPV/r)	No Evidence of Resistance

\*\* Protease Inhibitors administered with low-dose ritonavir for pharmacological boosting.

Resistance interpretation is based upon interpretation by an international expert panel (the International Panel of HIV-1 and HIV-2) using phenotypic and genotypic response data available as of June 2007 for combination of Protease and RT sequences to determine drug resistance. These include primary and secondary resistance.

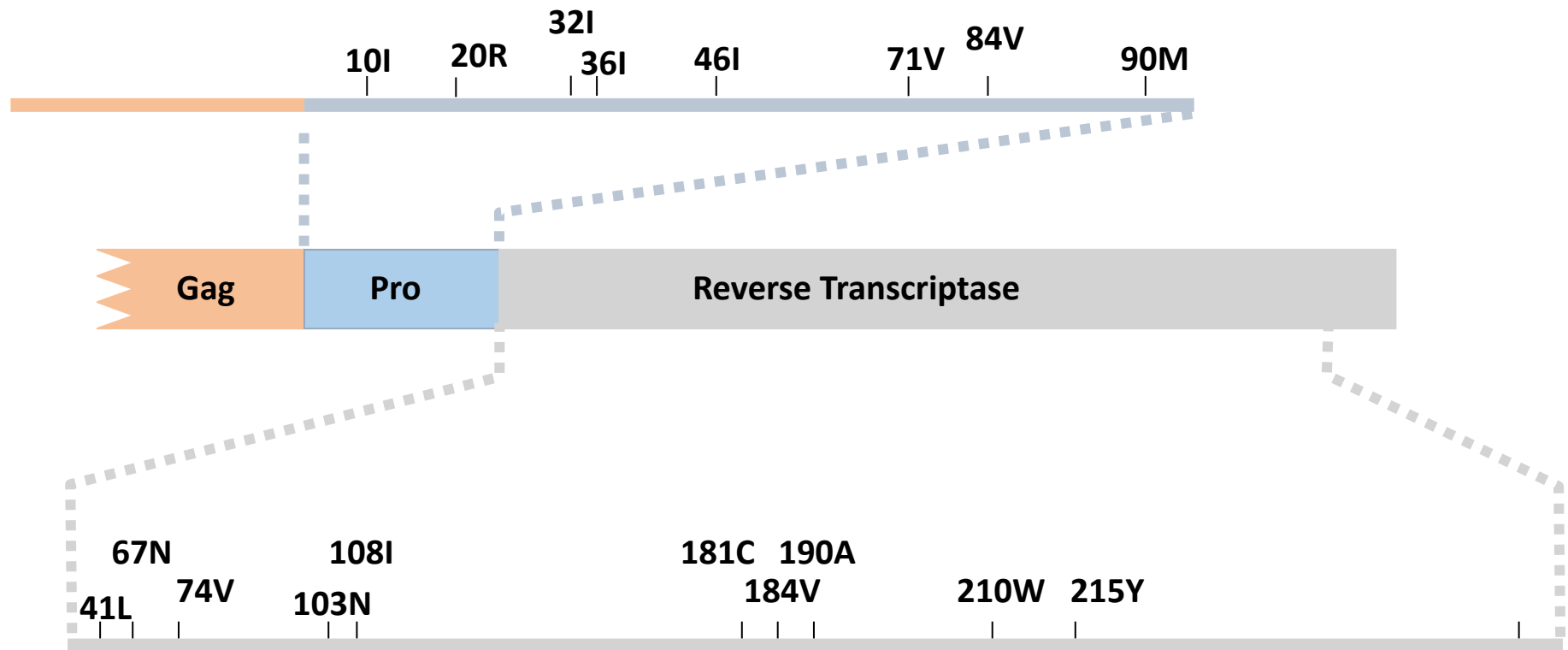


# Consequences were significant





# GT resistance test



# Why does it occur?

- Viral replication in presence of detectable drug(s)
  - Inadequate combination of drugs
    - Not potent enough
    - **Pre-existing resistance**

# Case 1

- 33 year old heterosexual male
- Presents with oropharyngeal and oesophageal candidiasis
- CD4 142 cells/mL
- Viral load 37,567 copies/ml
- Started on Septrin

# So what are you going to choose..

1. AZT and 3TC
2. TDF and FTC
3. TDF and 3TC
4. TDF and AZT
5. Other

Audience  
vote



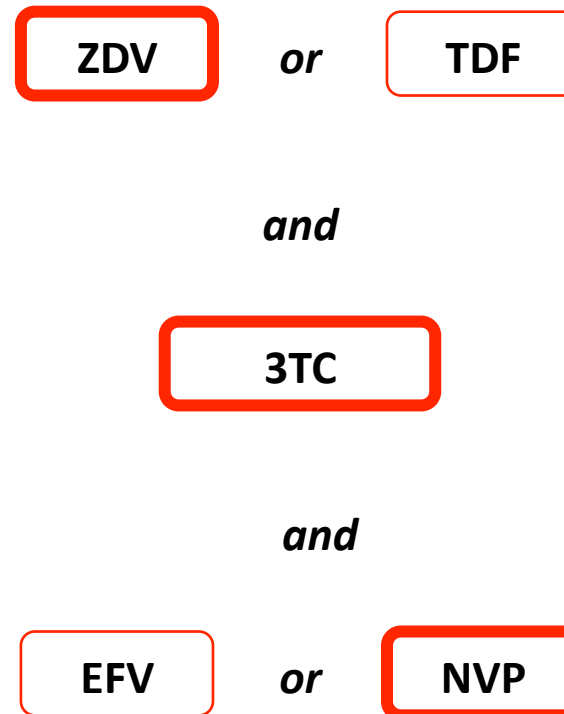
# So what are you going to choose..

1. NVP
2. EFV
3. ATAZ/r
4. LOP/r
5. Other

Audience  
vote

# Case 1

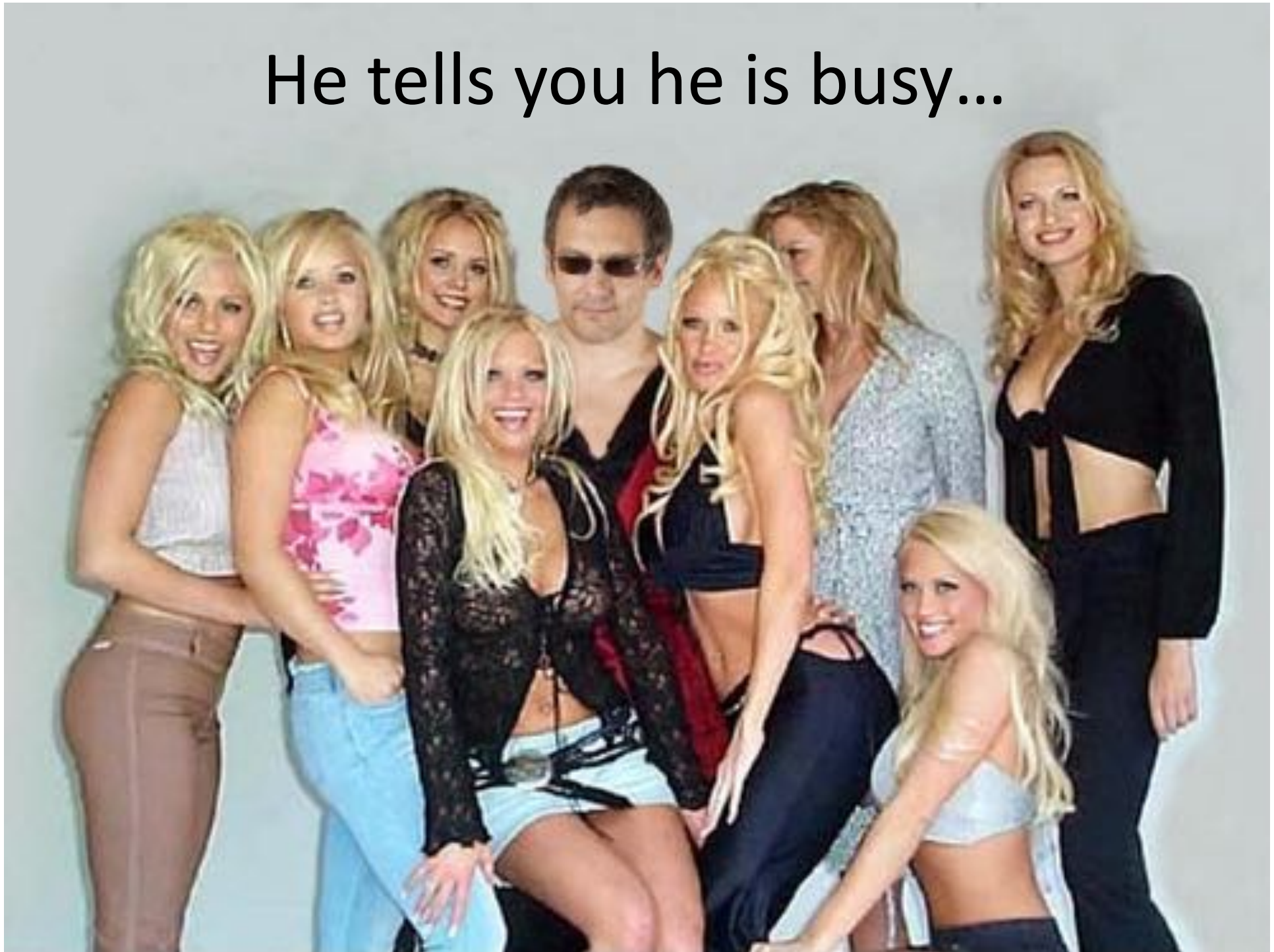
- 33 year old heterosexual male
- Presents with oropharyngeal and oesophageal candidiasis
- CD4 142 cells/mL
- Viral load 37,567 copies/ml
- Started on Septrin



Ask professor for help in getting/  
interpreting a resistance test...



He tells you he is busy...





And he is!





# His football club need him

## Man Utd 2 Chelsea 1

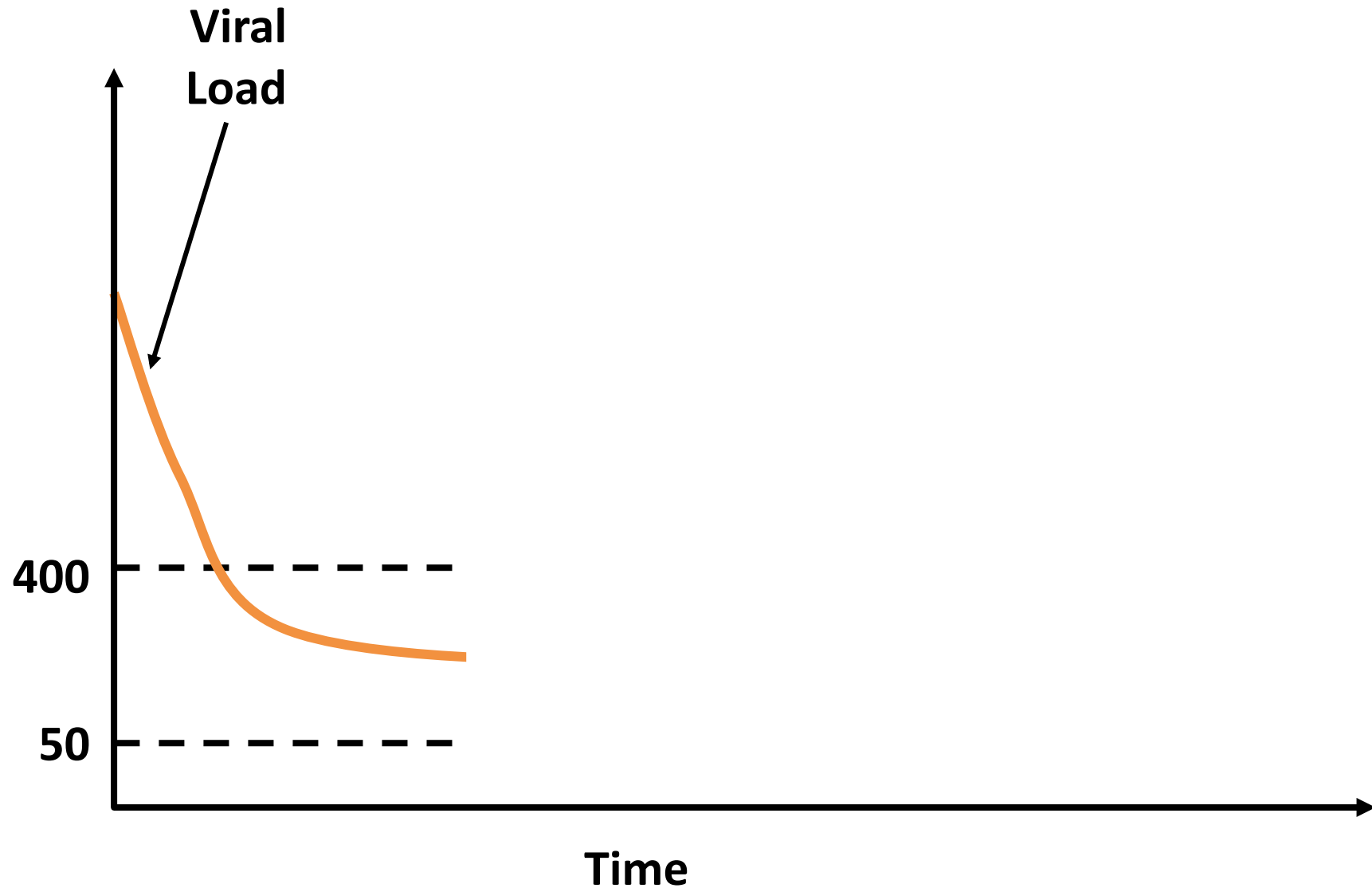


Barclays Premier League table

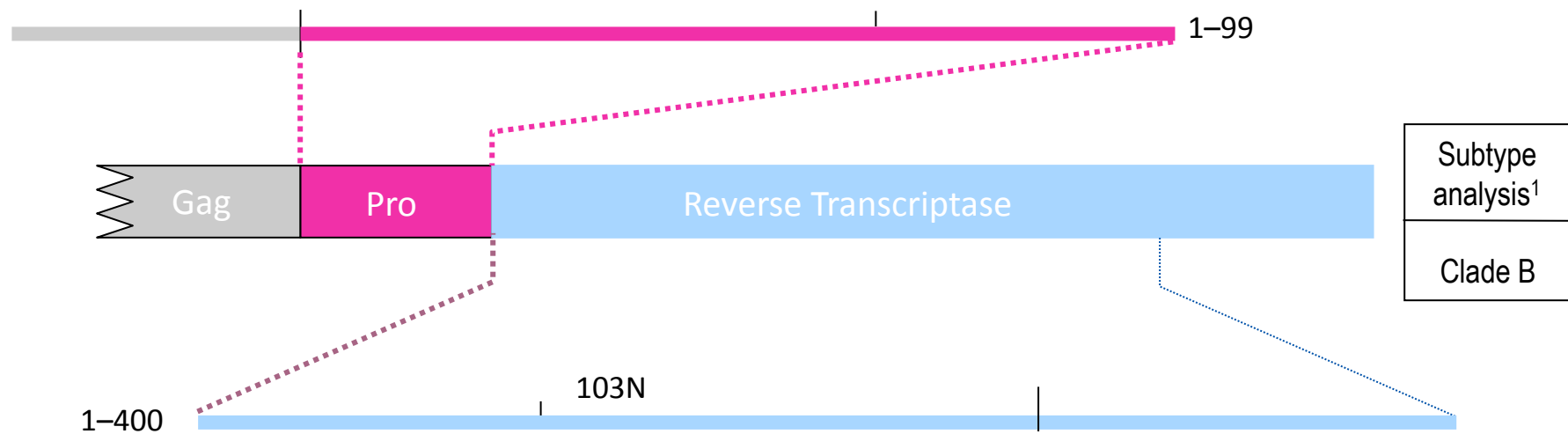
Season: 11/12 (30/03/12)

Team	P	W	PTS
1. Man Utd	38	26	78
2. Chelsea	38	21	70
3. Arsenal	38	20	67
4. Man City	38	21	62
5. Tottenham	38	17	54
6. Liverpool	38	18	53
7. Everton	38	9	31
8. Stoke	38	7	26
9. Bolton	38	7	26
10. Fulham	38	7	25
11. Newcastle	38	7	24
12. Sunderland	38	7	24
13. West Brom	38	11	23
14. Aston Villa	38	7	23
15. Blackburn	38	7	20
16. Birmingham	38	7	19
17. Wolves	38	5	17
18. Blackpool	38	5	16
19. Wigan	38	5	16
20. Reading	38	5	16

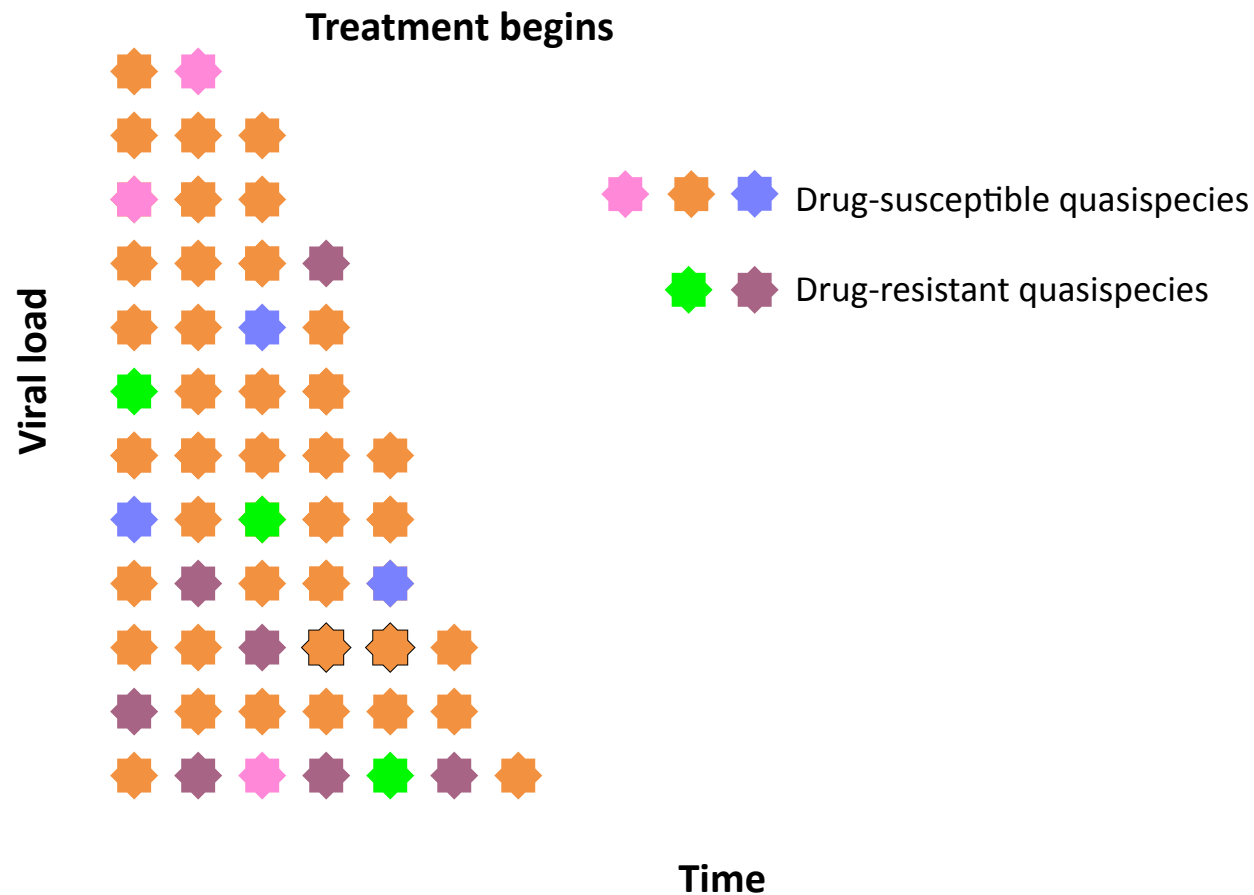
# Response to therapy – case 1



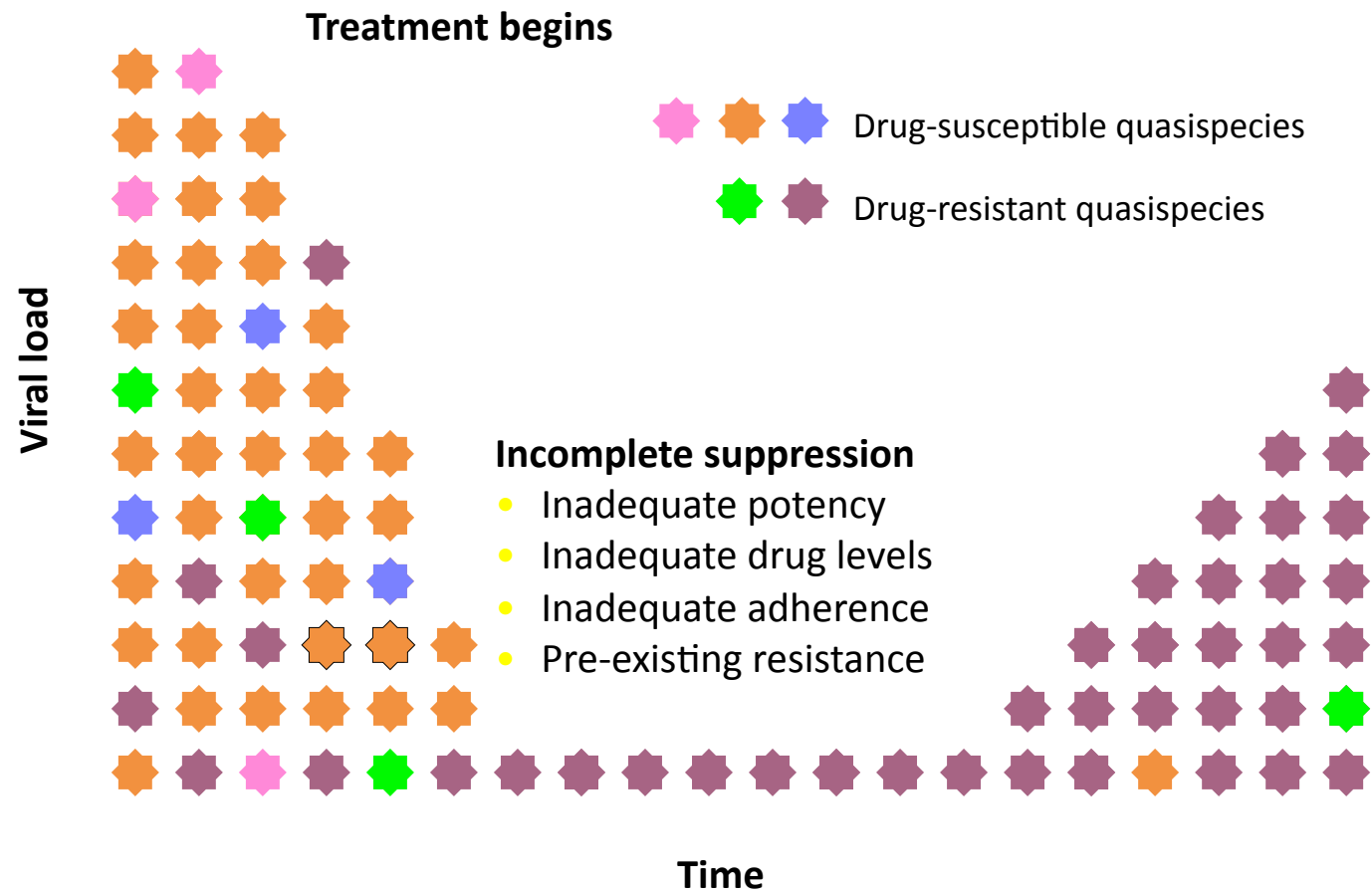
# Resistance-associated mutation identified on baseline test



# So what happens when resistance is present? Selective Pressure of Therapy



# Selective Pressure of Therapy



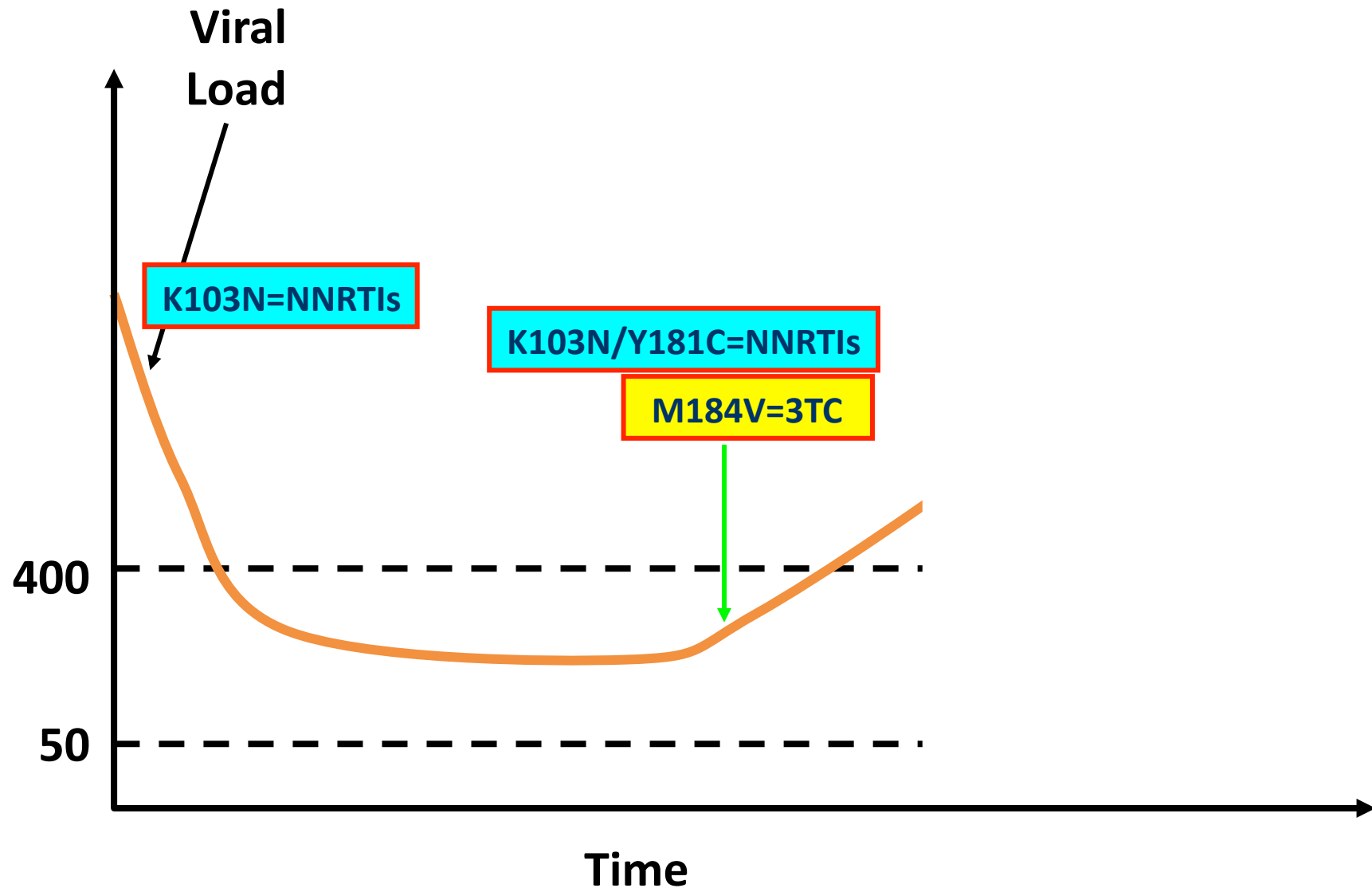


# Patient – CD4 falls to 230 from 501 – what would you do?

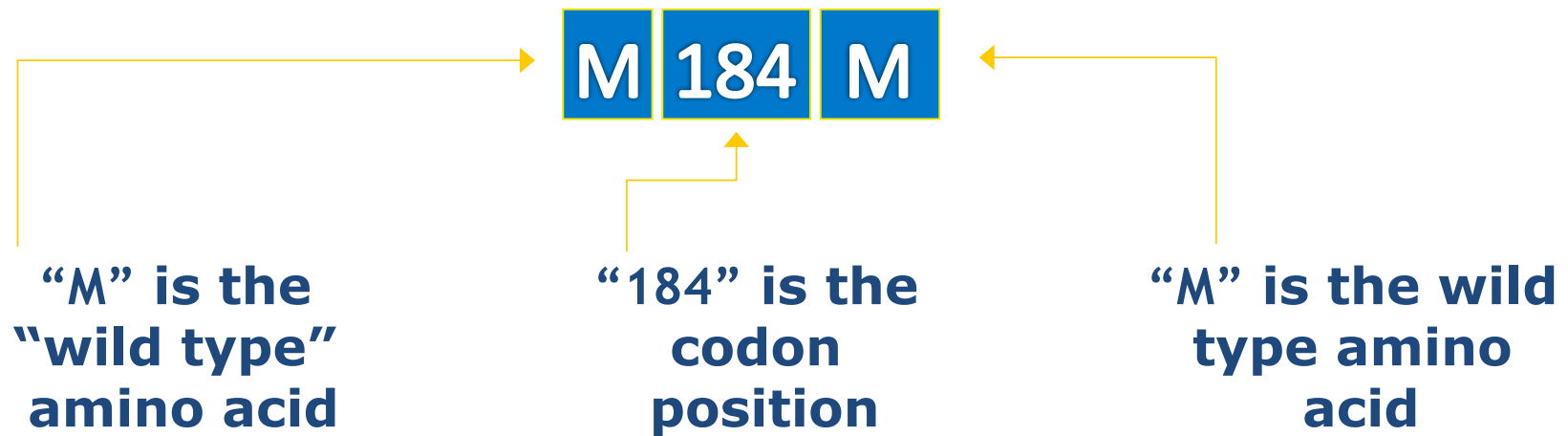
1. Carry on with NVP/AZT/3TC
2. Switch to EFV/TDF/FTC
3. Switch to ATAZ/r/TDF/FTC
4. Persuade Dr Nelson to organise another resistance test
5. Repeat CD4 and wait till fall further
6. Other

Audience  
vote

# Response to therapy – case 1

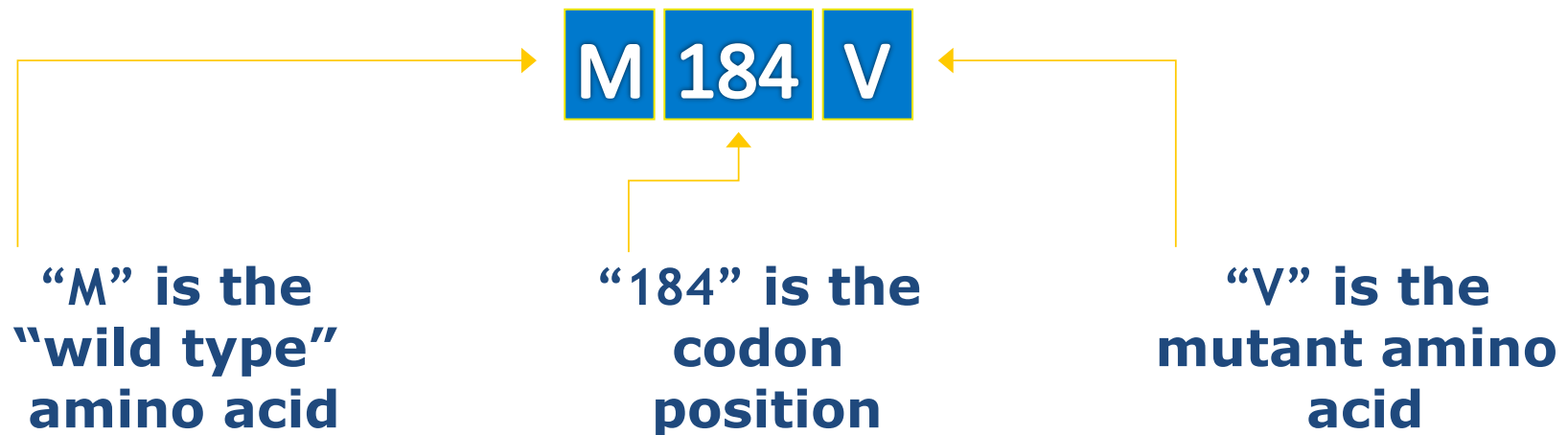


# What do these mean..... Before 3TC

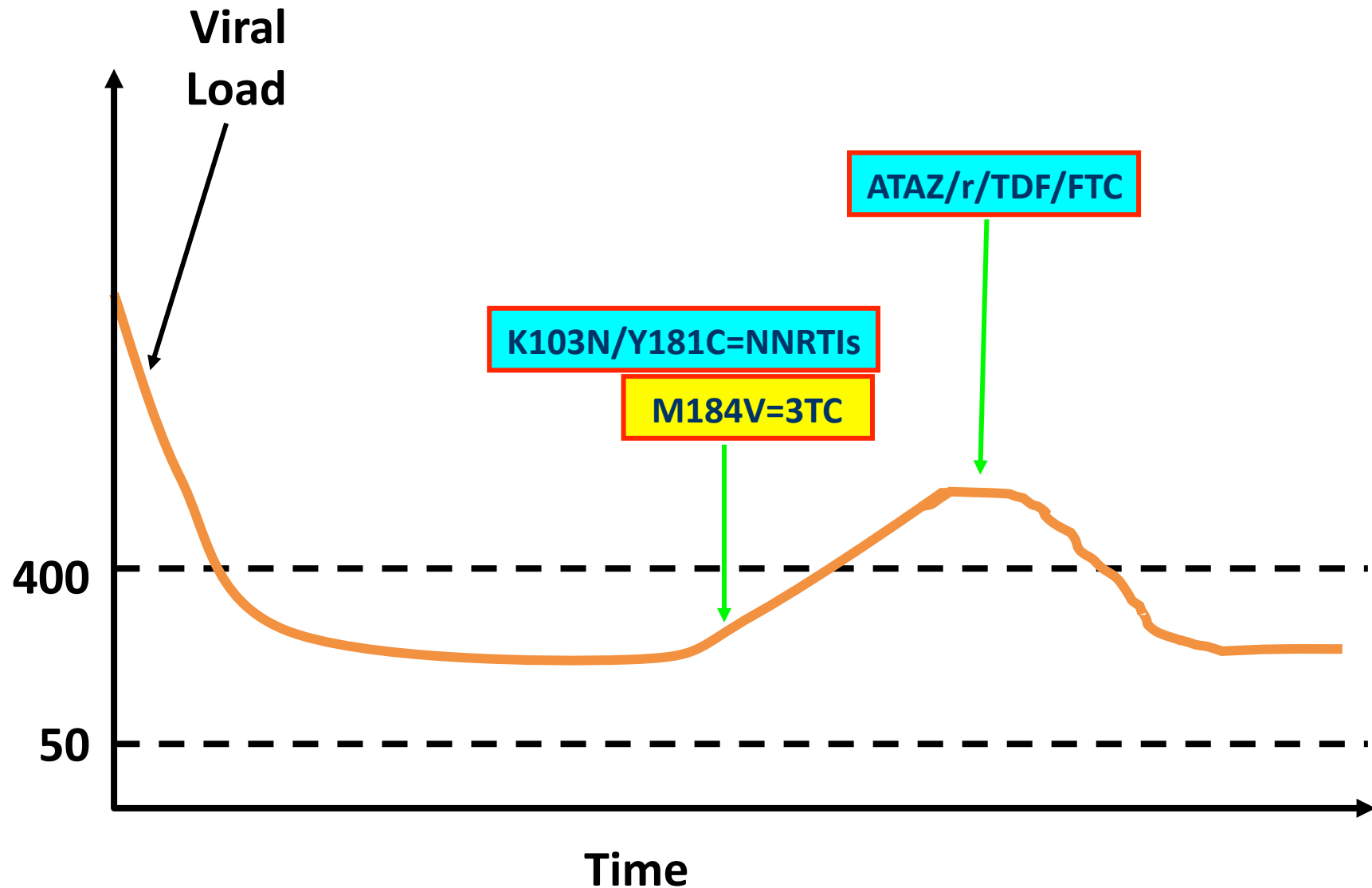


# After 3TC and resistance...

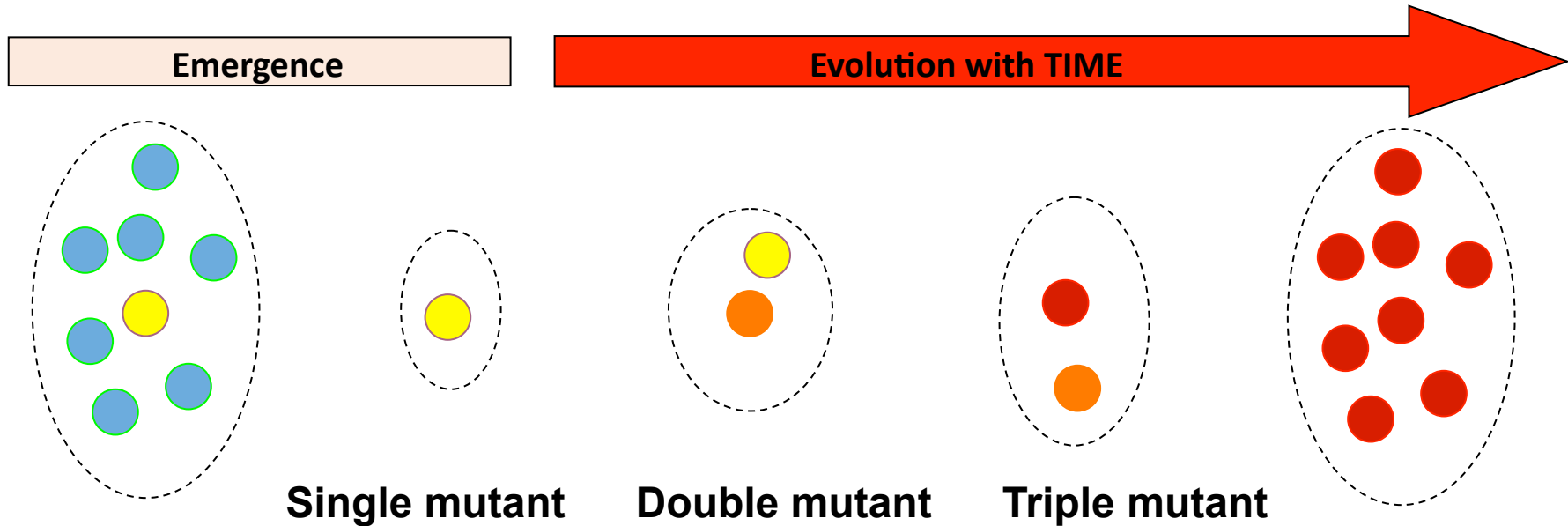
- How do we identify a resistance mutation?



# Response to therapy – case 1



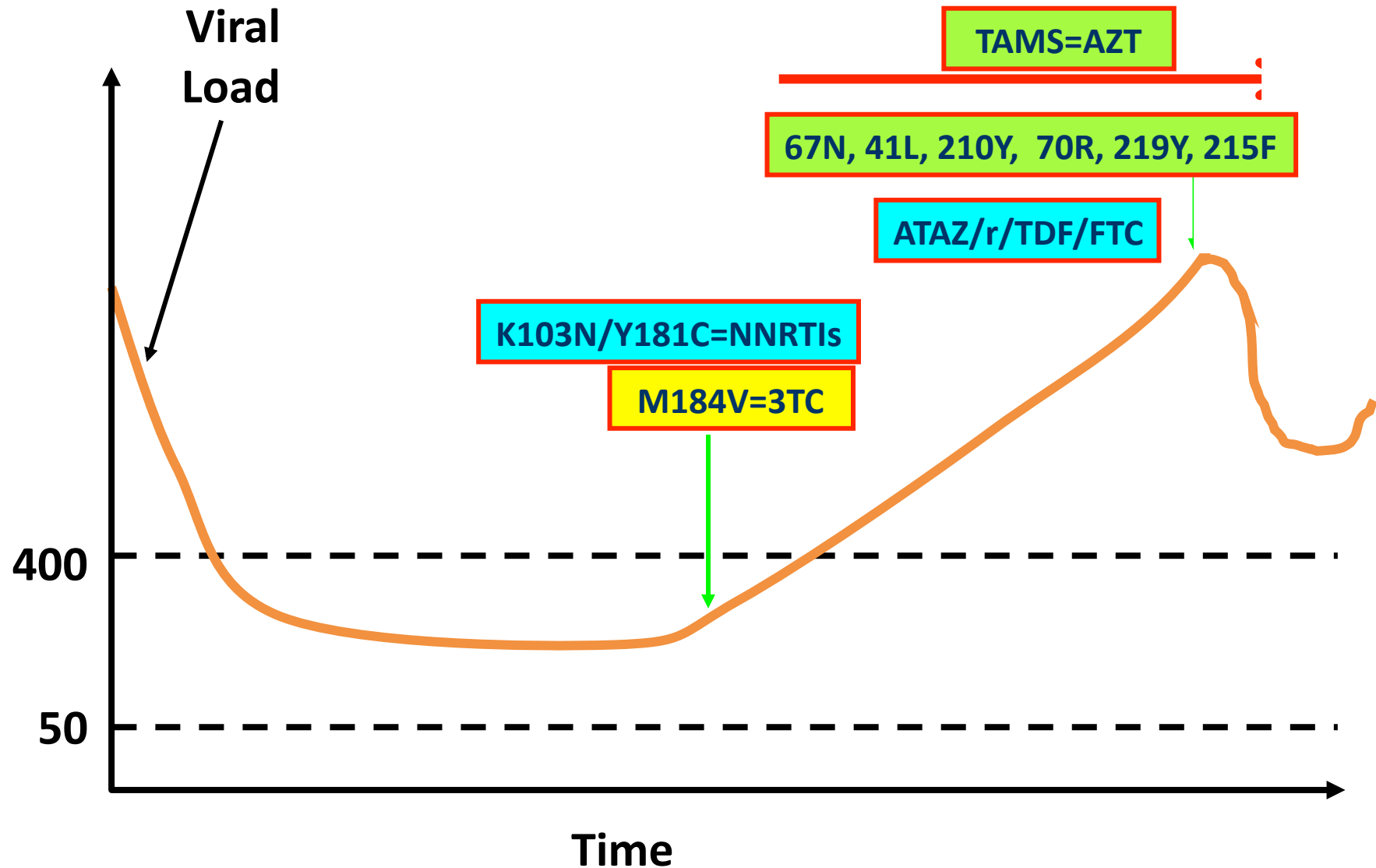
# Emergence and evolution of resistance



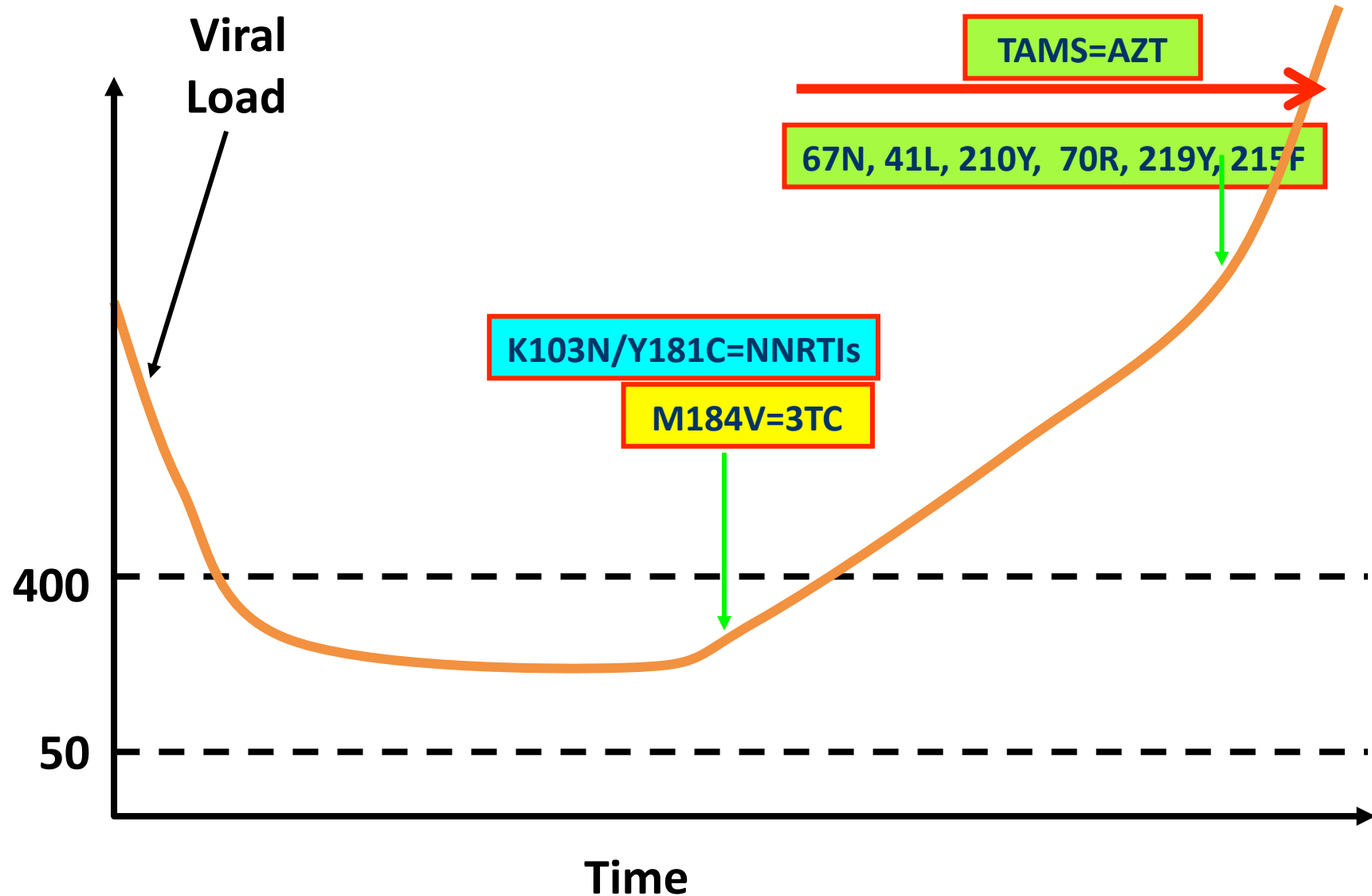
- Increasing number of mutations
- Accumulation of mutations on the same viral genome
- Initially reduced viral fitness
- Compensatory changes restore fitness



# Response to therapy – case 1



# Response to therapy – case 1



# The more mutations the more resistance...

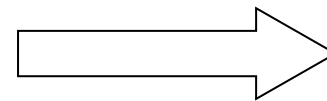
*Accumulation of TAMs:*

M41L, D67N, K70R, L210W, T215Y/F, K219Q/E

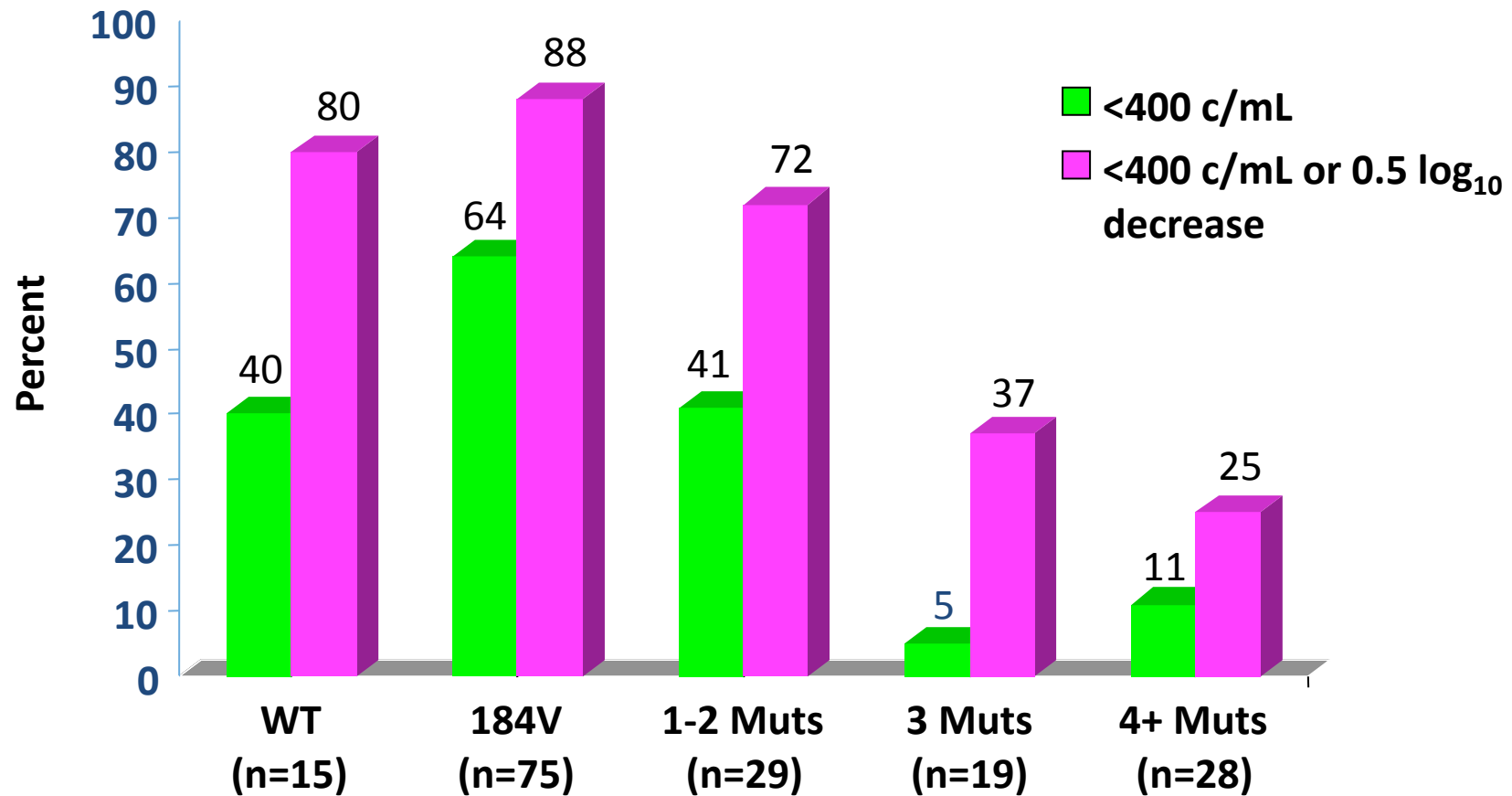
Susceptible      Partial Resistance      Resistance



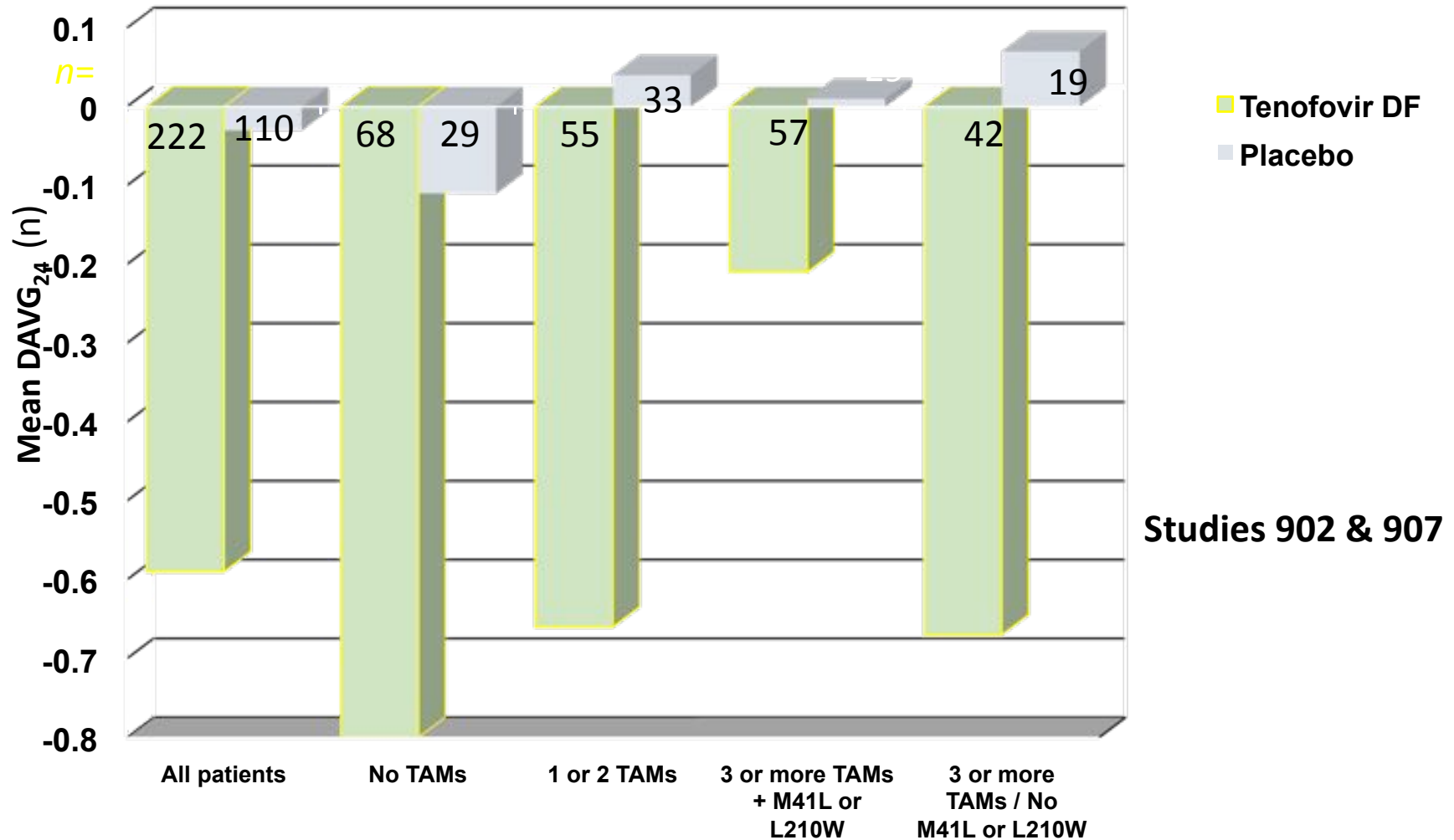
Number of TAMs present



# The more TAMMS the LESS abacavir effect..



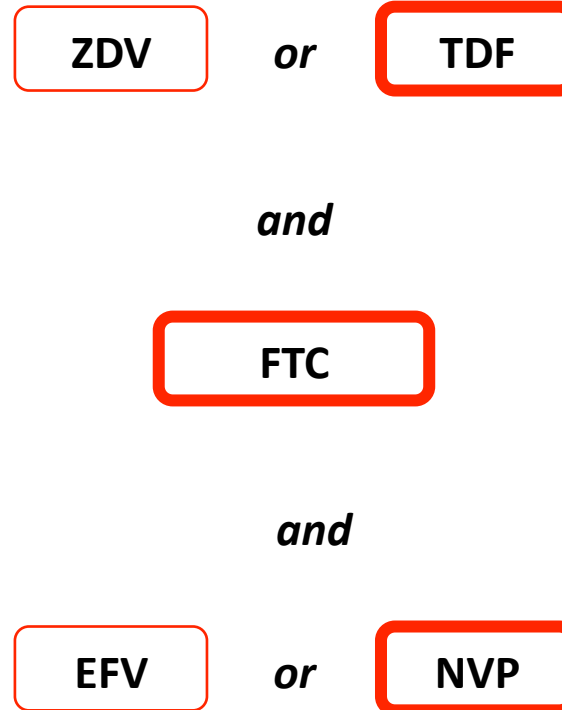
# The more TAMs the LESS tenofovir effect...



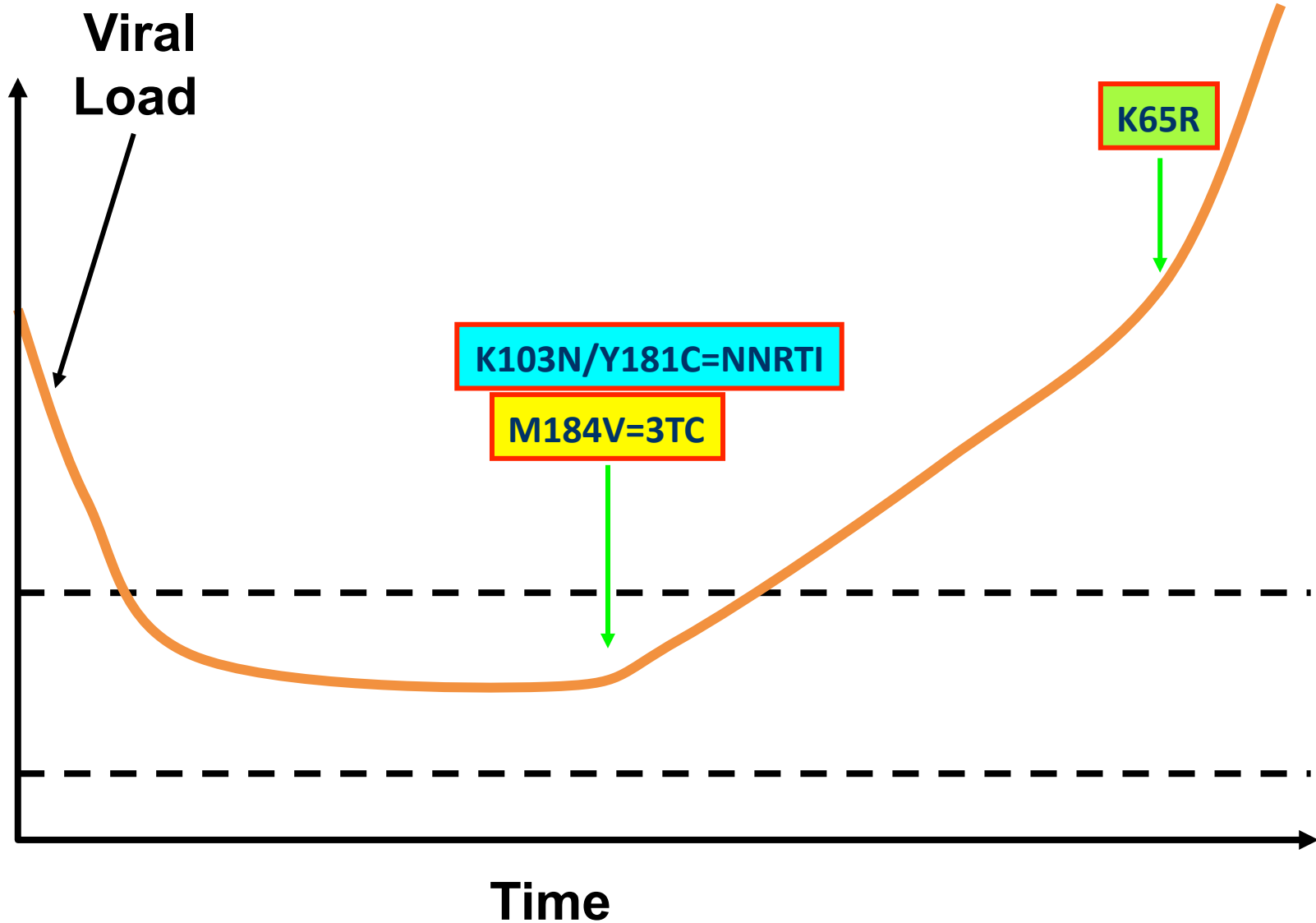


# What if case 1 had been treated differently..

- 33 year old heterosexual male
- Presents with oropharyngeal and oesophageal candidiasis
- CD4 142 cells/mL
- Viral load 37,567 copies/ml
- Started on Septrin

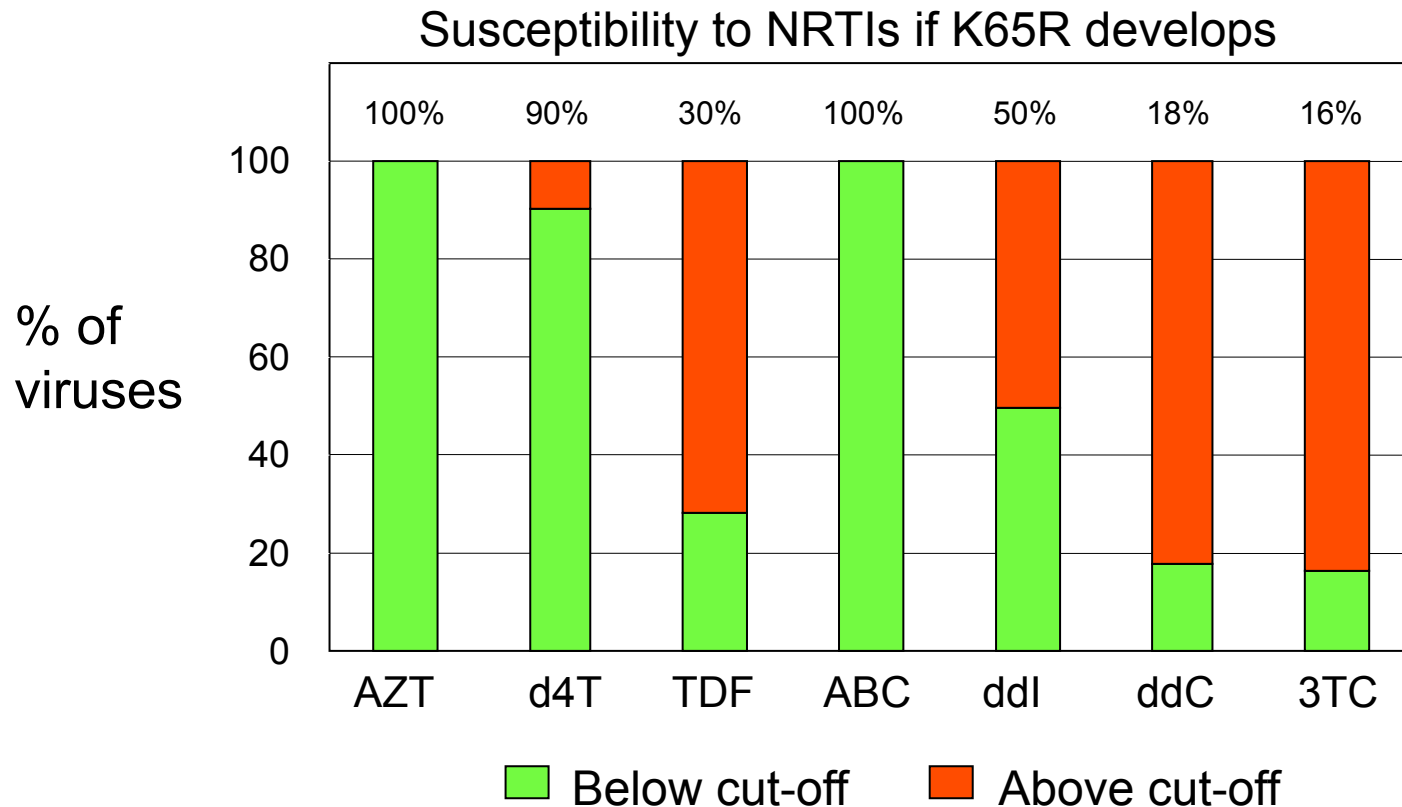


# Case 1 with TDF/FTC...



# Susceptibility to NRTIs if K65R develops

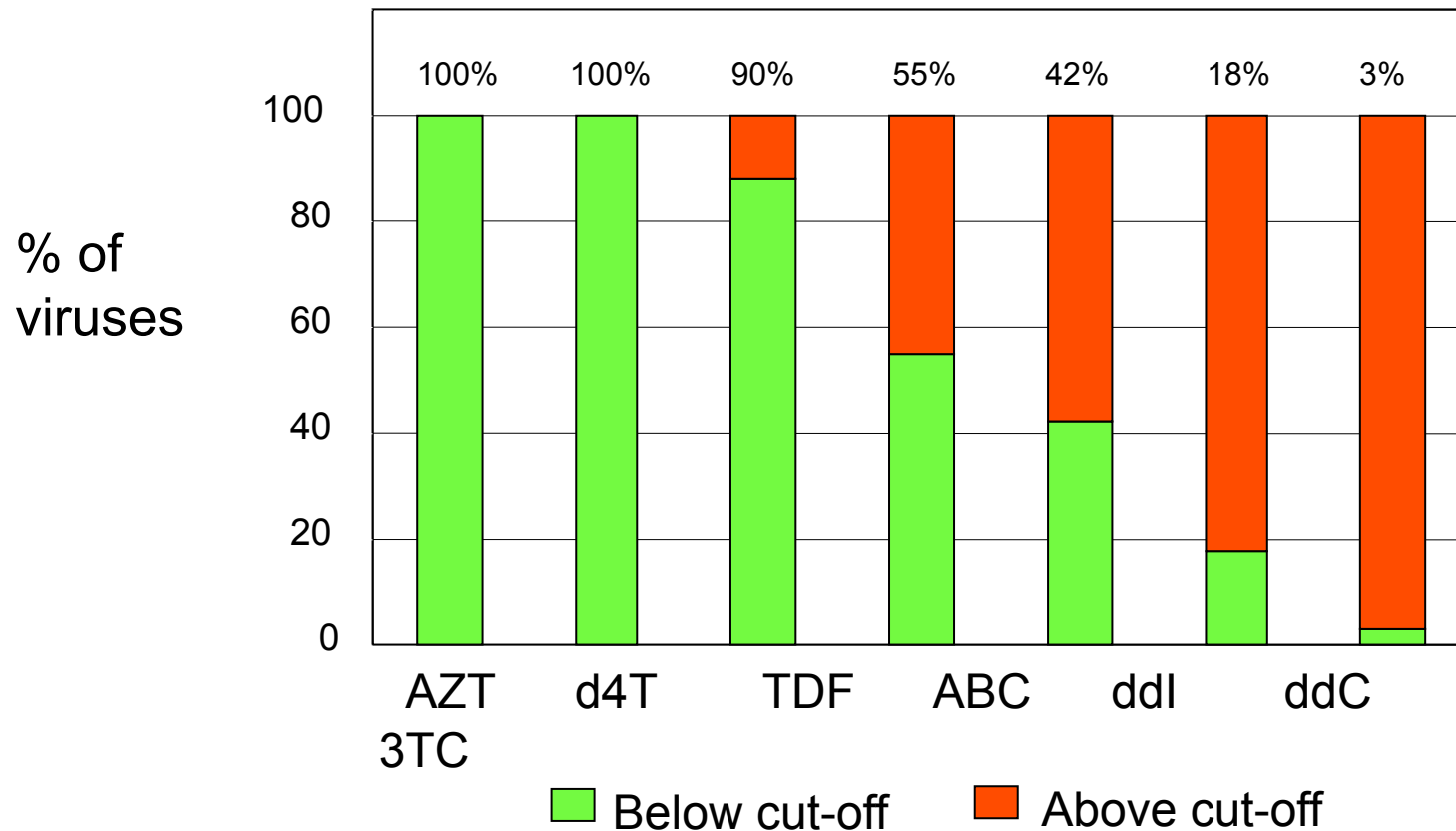
PhenoSense Results for K65R alone (n=50)



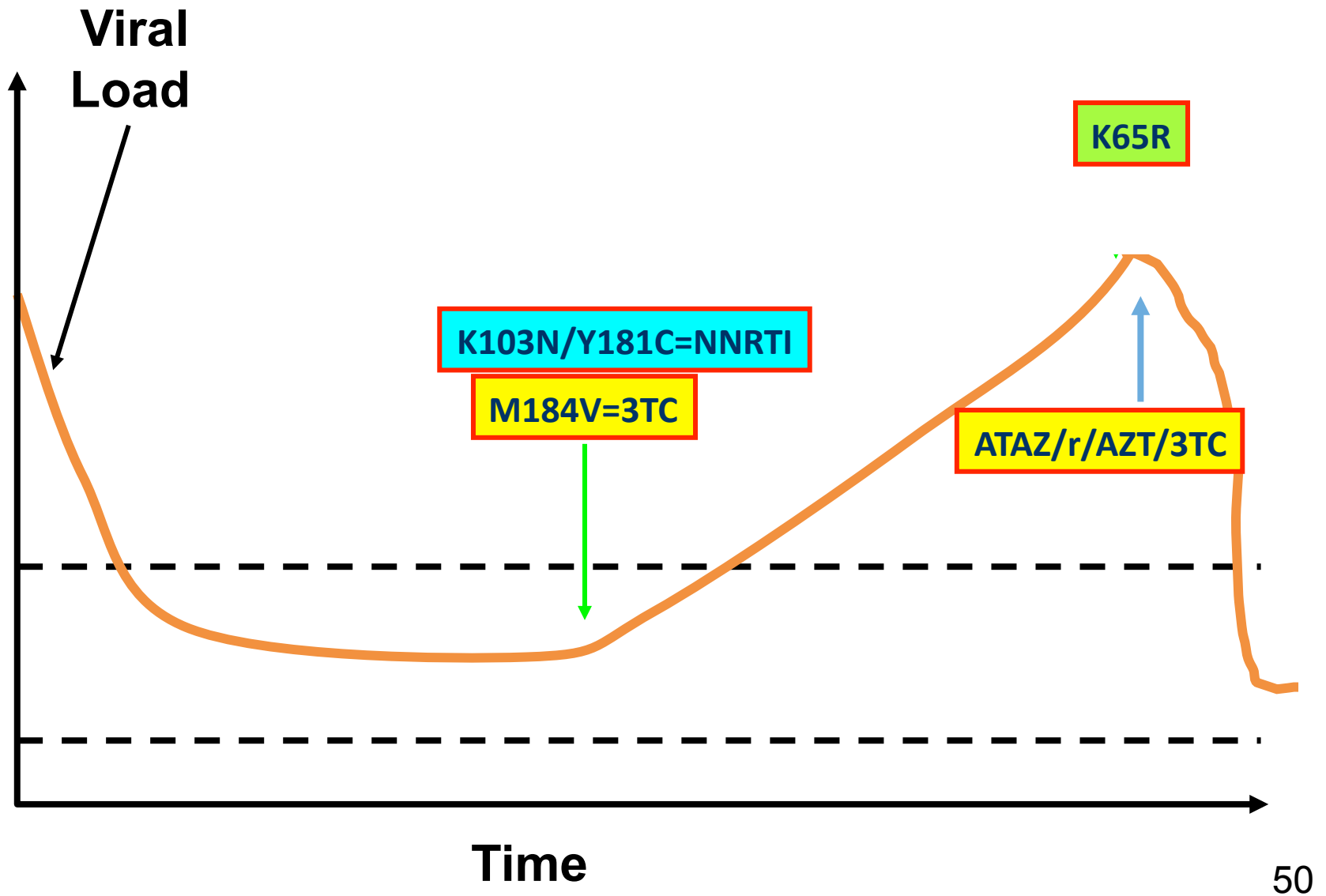
For tenofovir, all viruses were below the 4.0-fold cut off for no response.

# Susceptibility to NRTIs if *K65R and M184V* develop

PhenoSense Results for K65R + M184V (n=58)



# Case 1 with TDF/FTC...



# Hence sequencing Options: PI AND

3TC/  
FTC

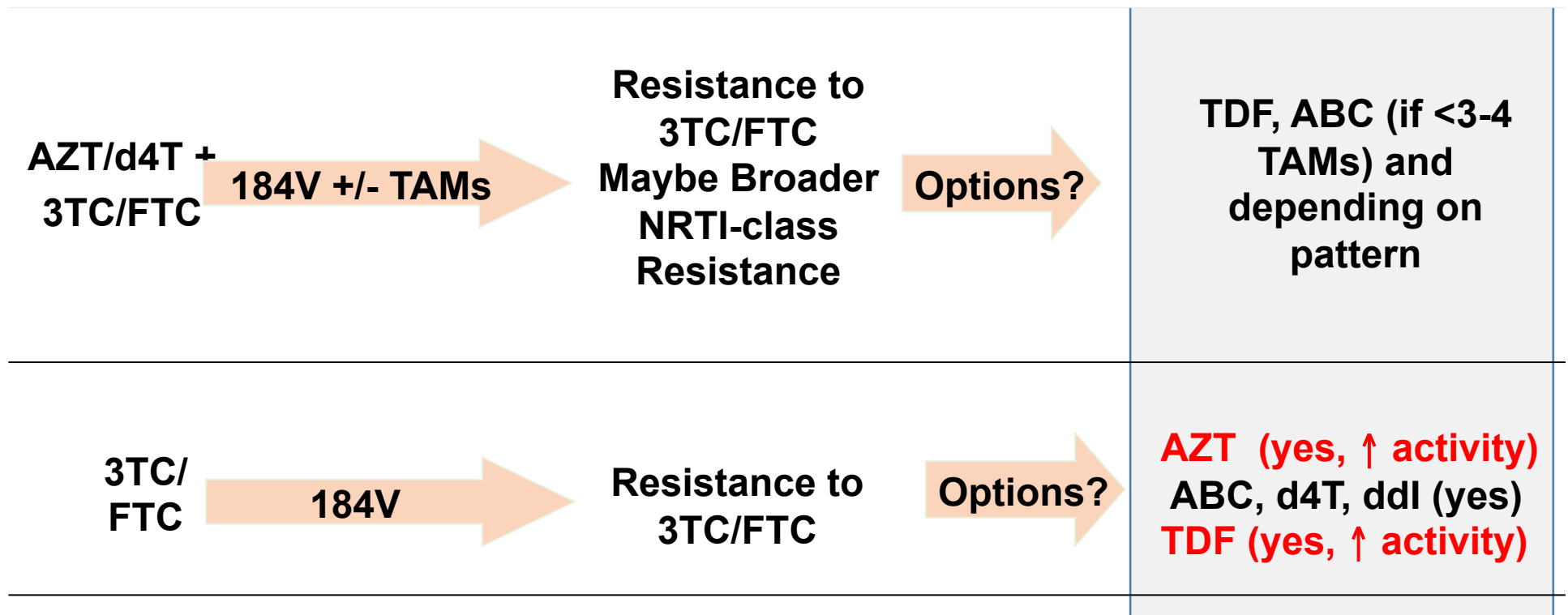
184V

Resistance to  
3TC/FTC

Options?

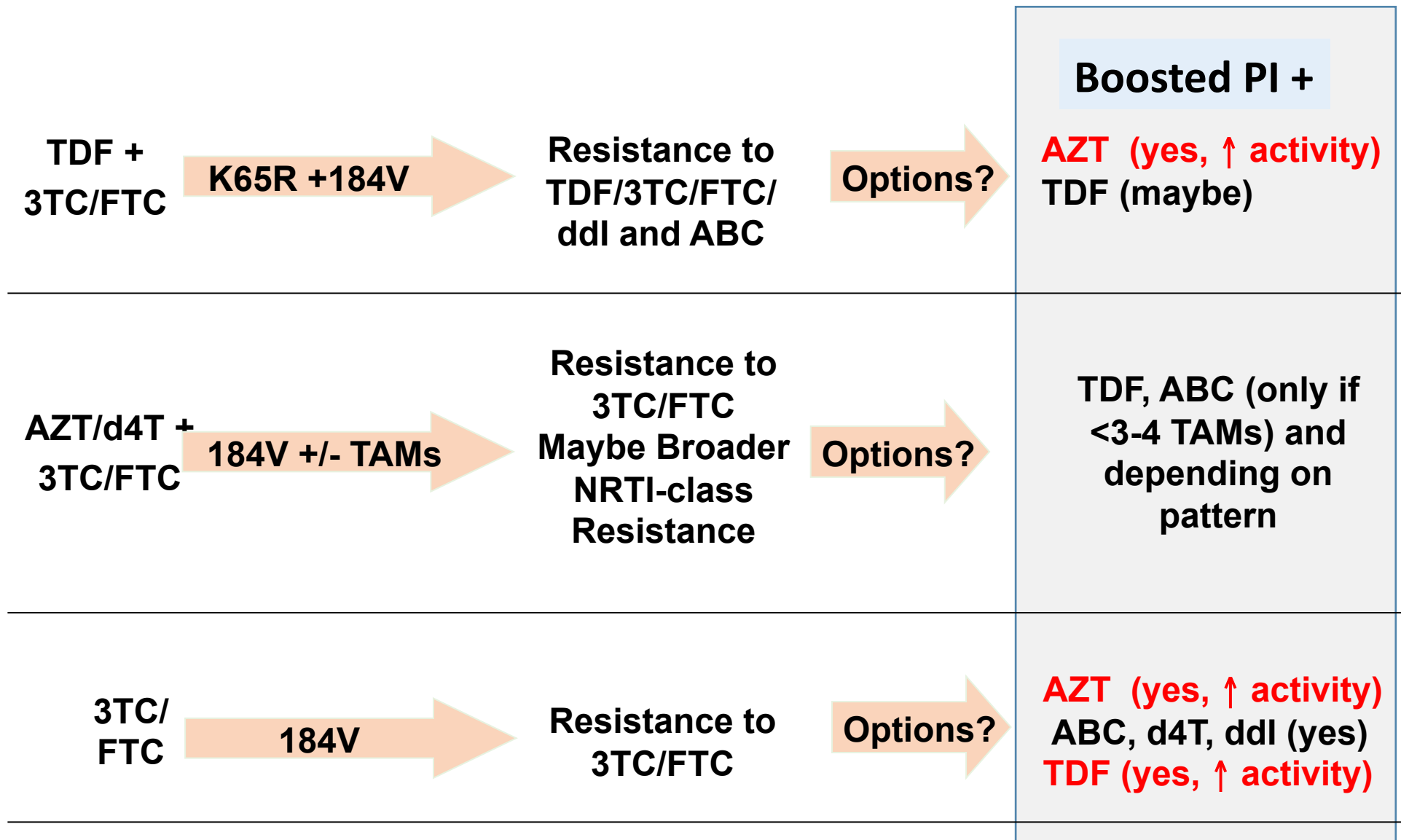
**AZT (yes, ↑ activity)**  
**ABC, d4T, ddl (yes)**  
**TDF (yes, ↑ activity)**

# Hence sequencing Options: PI AND





# Hence sequencing Options: PI AND....



So hands up who will start with...

1. AZT and 3TC

2. TDF and either FTC or 3TC

Audience  
vote

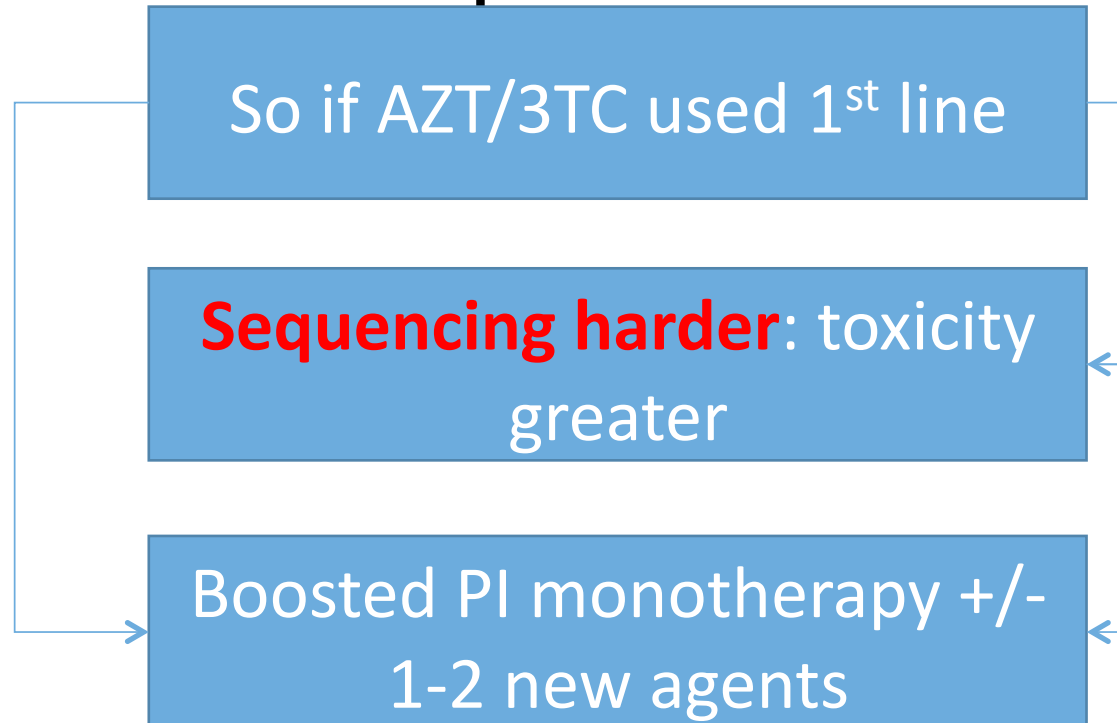
So hands up who will start with...

1. EFV or NVP

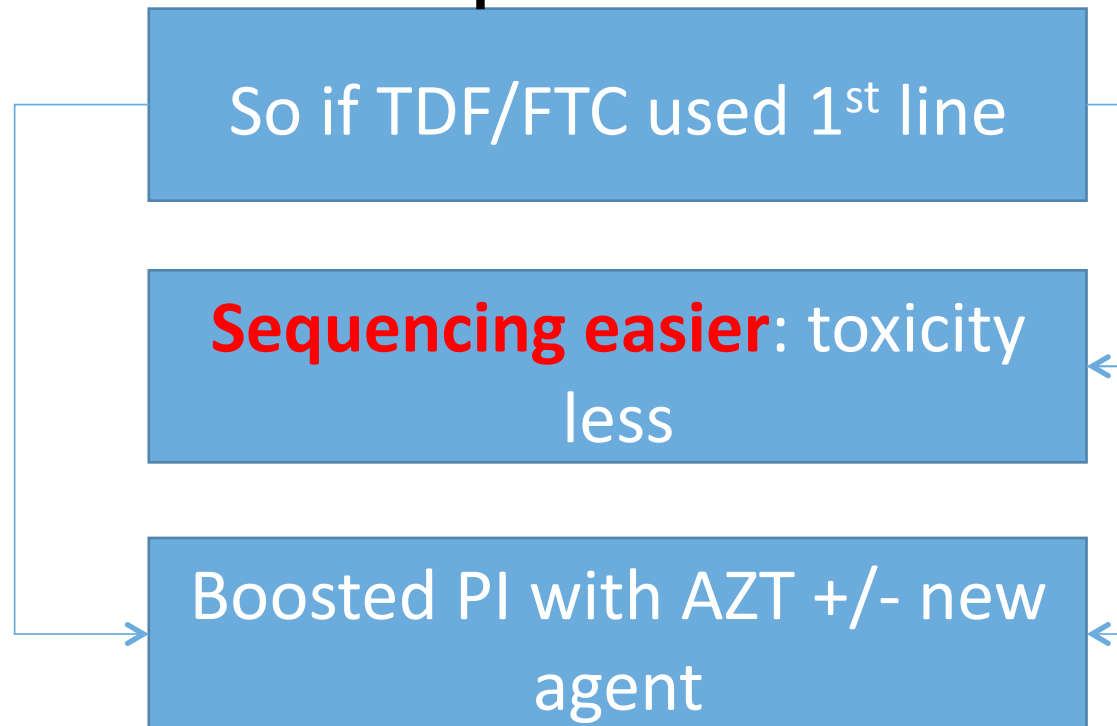
2. ATAZANAVIR boosted by ritonavir  
or KALETRA

Audience  
vote

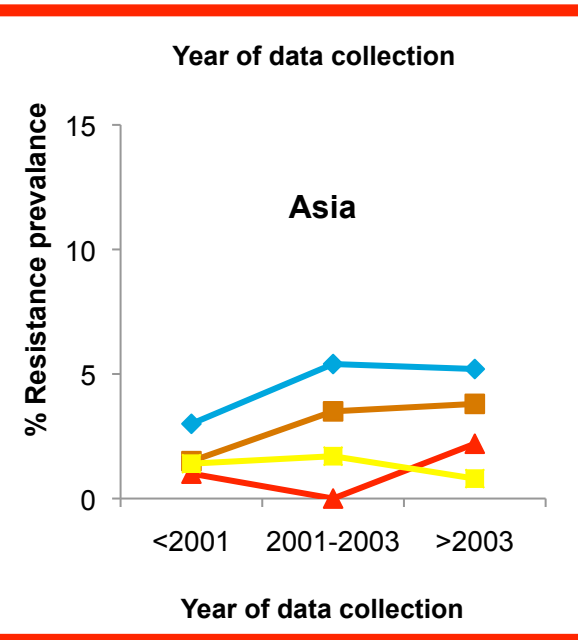
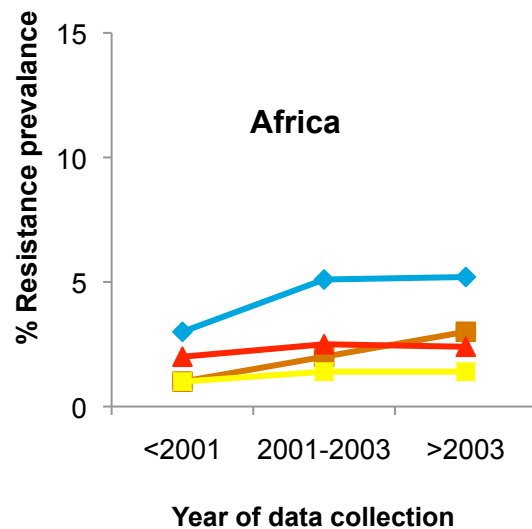
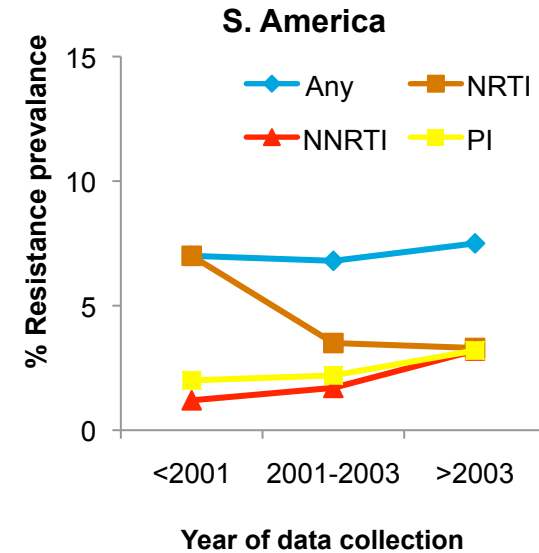
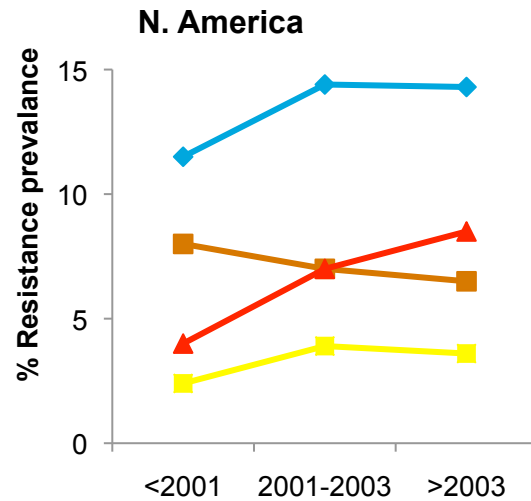
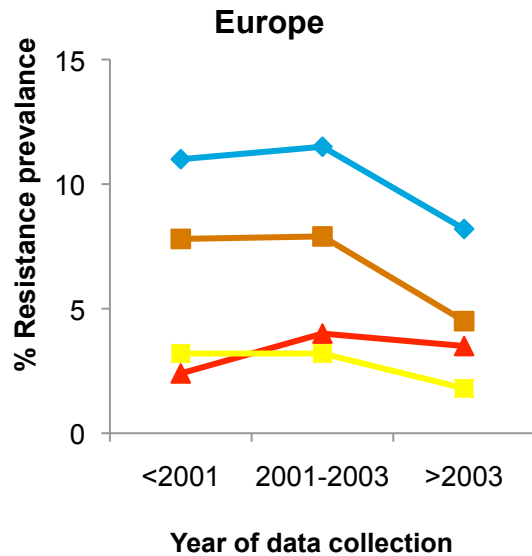
So choice of NRTI backbone is important when sequencing after resistance develops



So choice of NRTI backbone is important when sequencing after resistance develops

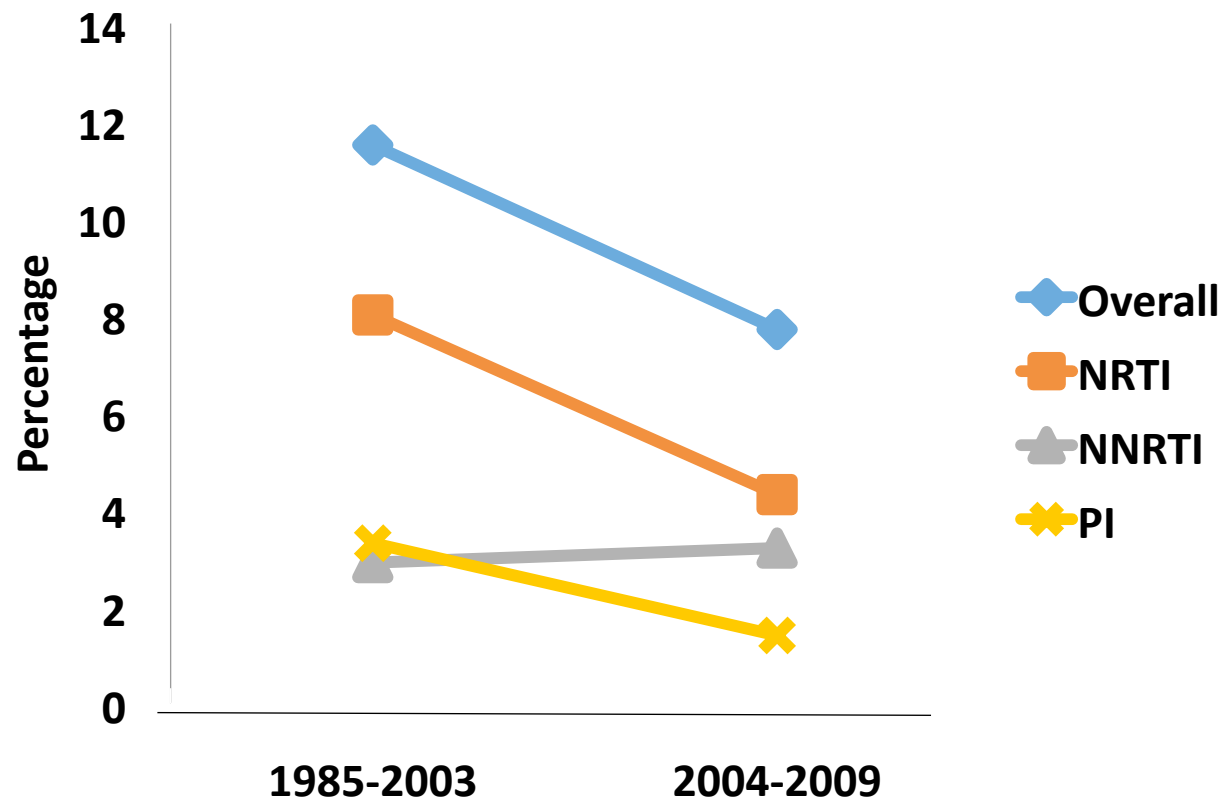


# How common is drug resistance in Myanmar?



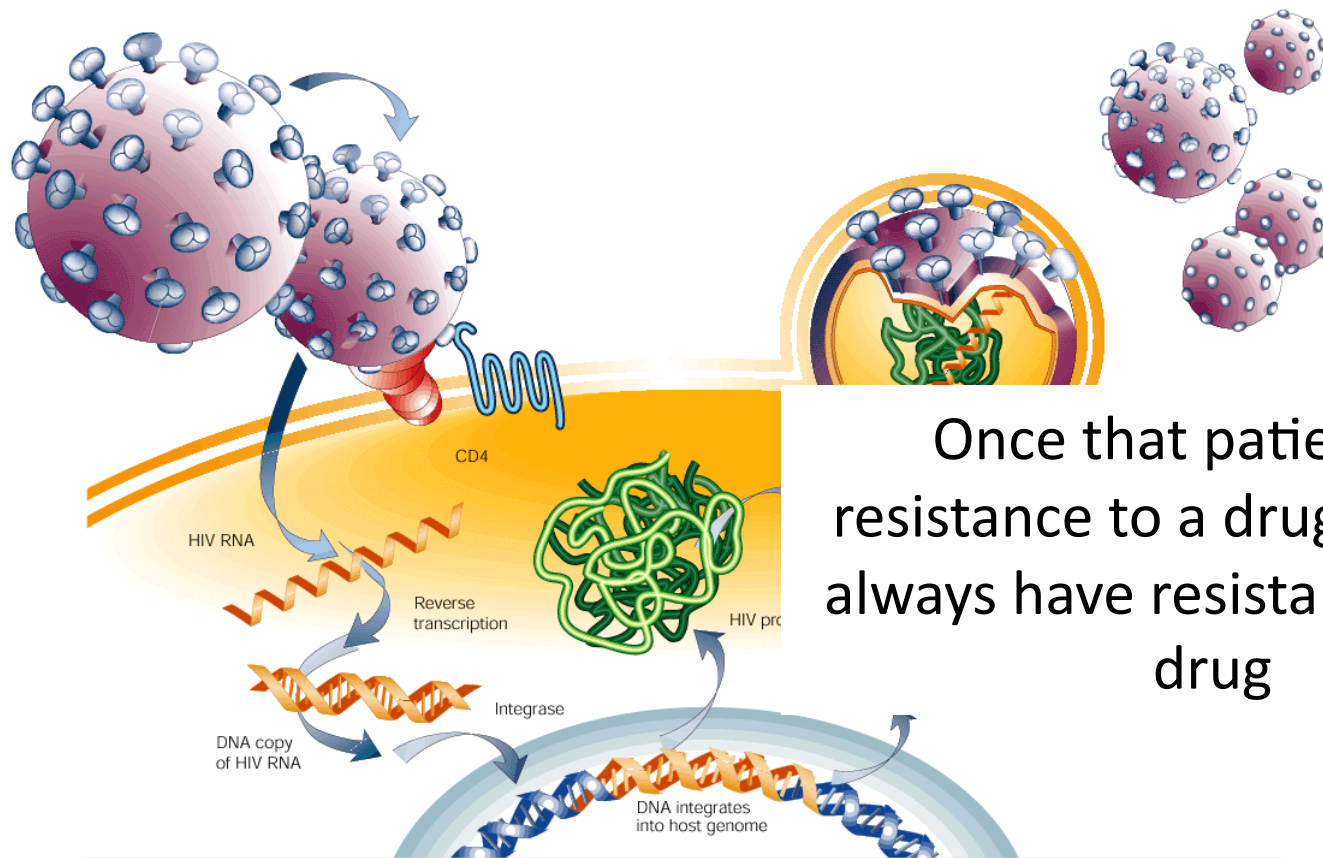
# In Europe around 8%

N=23,000 from 75 studies in 20 countries





# Why are we so bothered?

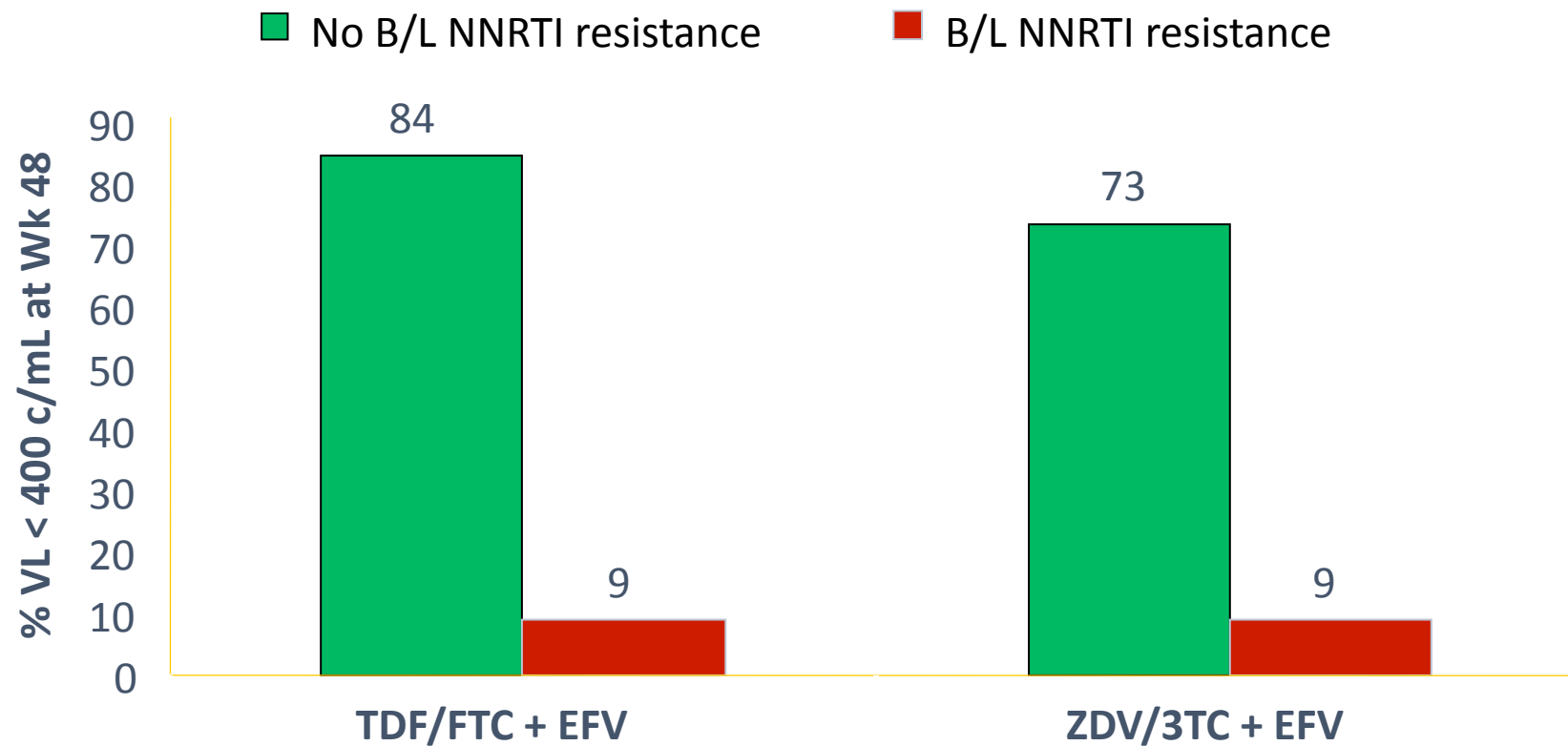


Once that patient has resistance to a drug, they will always have resistance to that drug

# Limitations of resistance testing

- Archived resistance
  - May be so low they cannot be detected.... But they are still there.... and will rapidly re-emerge under drug pressure
  - So you need to look at all previous resistance test results too
  - And maybe make a guess on what might be there.....

# Archived NNRTI Resistance Markedly Reduces Treatment Response



# Why does it occur?

- Viral replication in presence of detectable drug(s)
  - Inadequate combination of drugs
    - Not potent enough
    - Pre-existing resistance
    - **Low levels**
      - **Compliance**
      - **Absorption/metabolism**
      - **Interactions**

Poor compliance often  
results from  
intolerance/toxicity to  
ARVs







..... or too many  
tablets





# The advent of STRs

1996  
30+ Pills a Day



2006  
the first STR



# Drug-drug Interactions

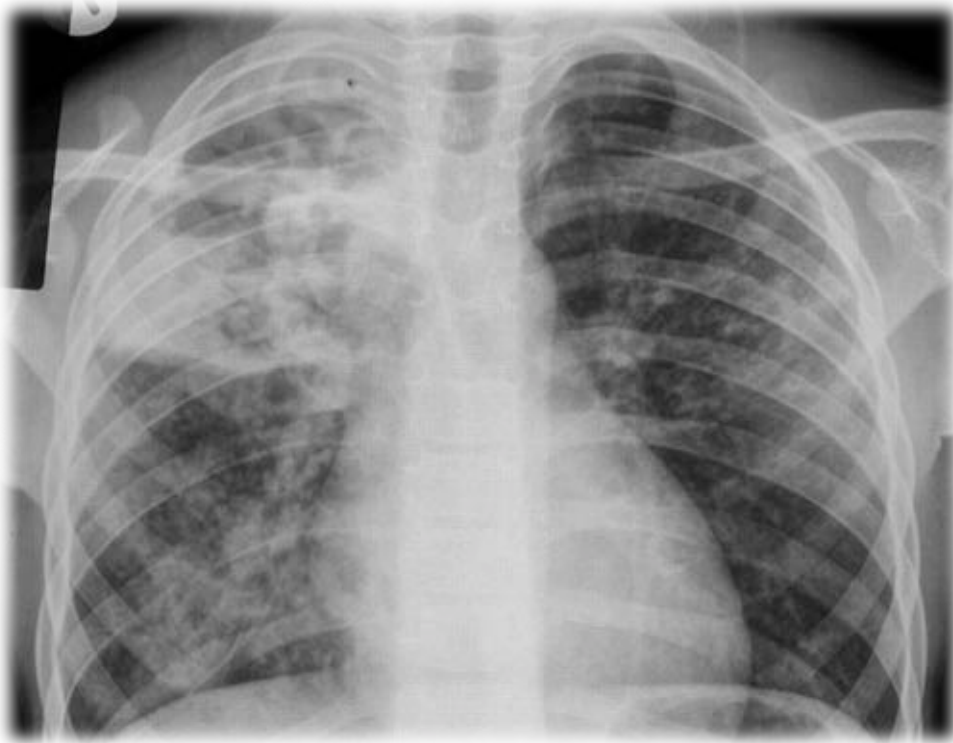


with or  
without food?

take together?

What about my other drugs?

# Drugs for HIV or non-HIV related issues



# Age and illnesses of getting old



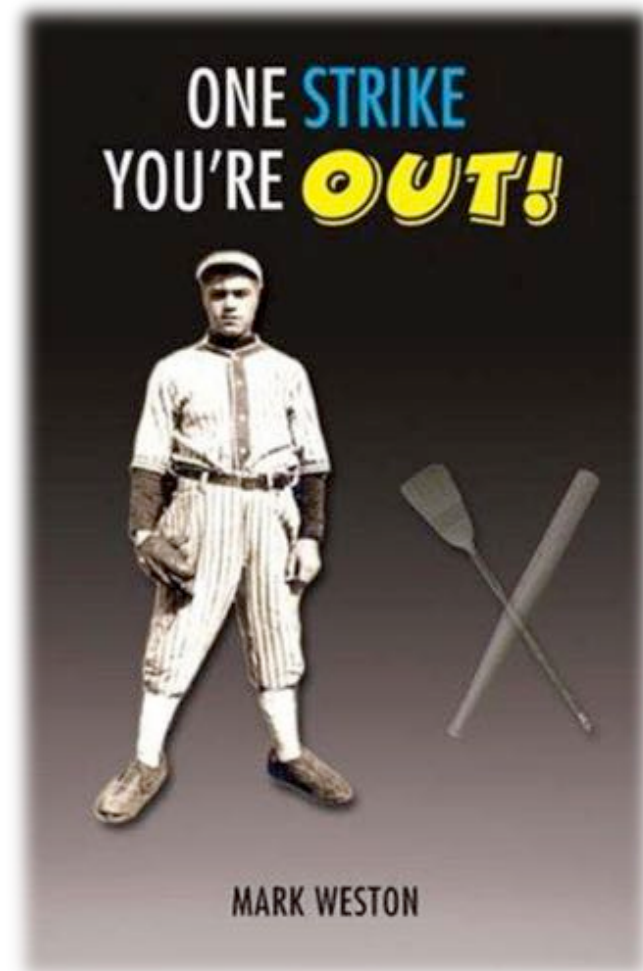
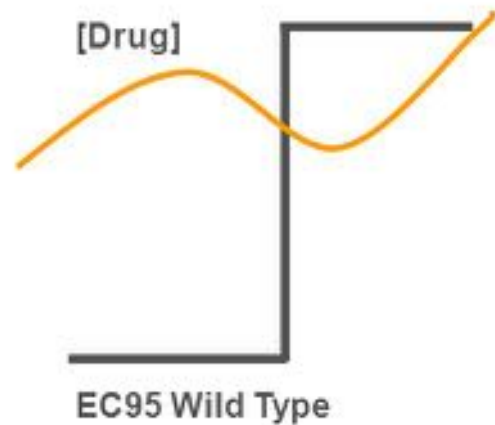
# Why does it occur?

- Viral replication in presence of detectable drug(s)
  - Inadequate combination of drugs
    - Not potent enough
    - Pre-existing resistance
    - Low levels
      - Compliance
      - Absorption/metabolism
      - Interactions
  - Treatment interruption
    - Patient
    - Healthcare system/professional

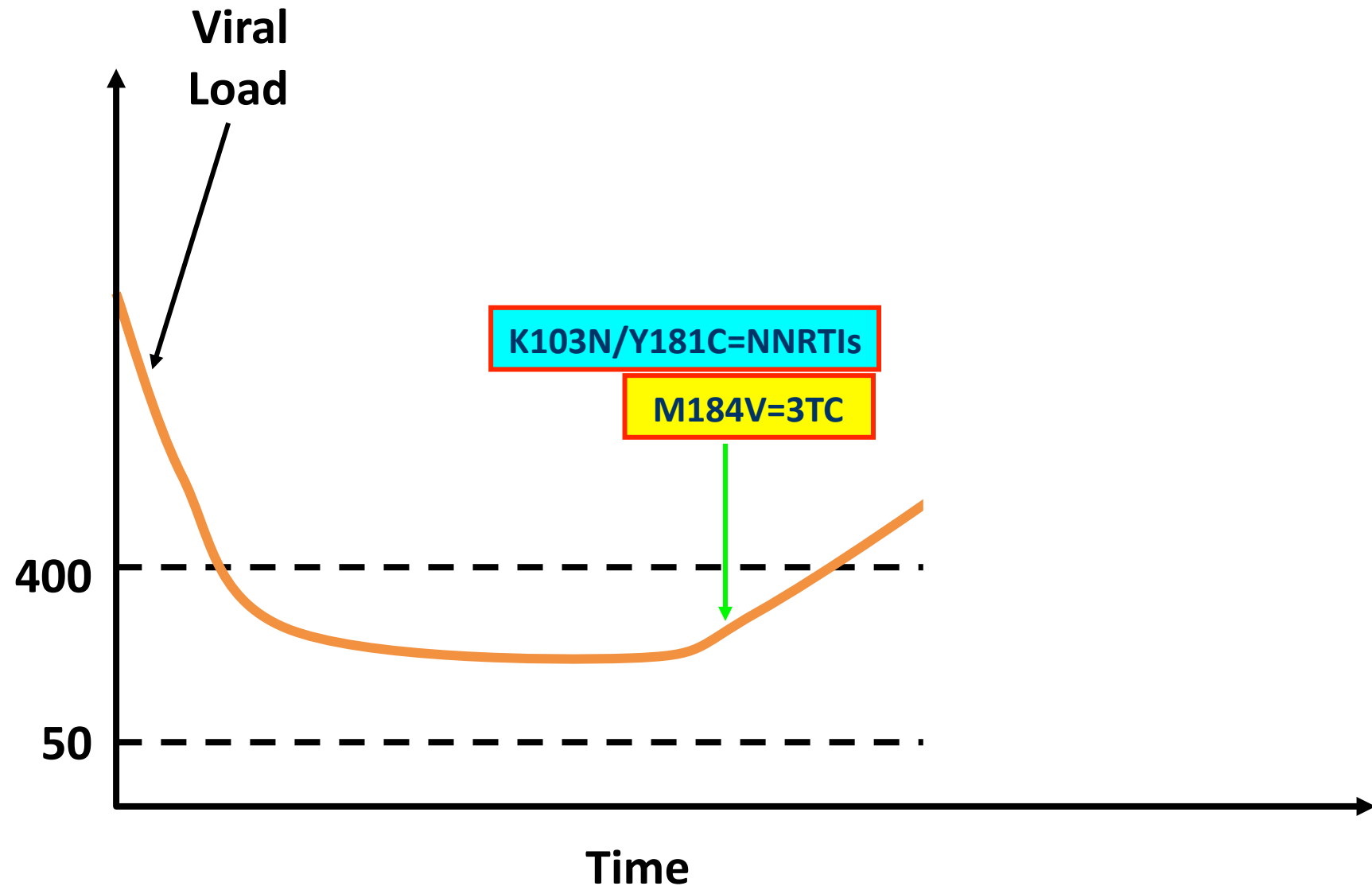


# Resistance - simple

- A single mutation may wipe out activity.....
  - M184V - lamivudine or emtricitabine
  - K103N – efavirenz or nevirapine



# Response to therapy – case 1





# In these situations class cross-resistance is usual

Single point mutations in the NNRTI binding pocket (e.g. K103N) lead to VF

	L	K	K	V	V	Y	Y	G	P
Efavirenz	100	101	103	106	108	181	188	190	225
	I	P	N	M	I	C	L	S	H
			S			I		A	
	L	K	K	V	V	Y	Y	G	
Nevirapine	100	101	103	106	108	181	188	190	
	I	P	N	A	I	C	C	A	
			S	M		L	L		
							H		

- As EFV and NVP share similar binding sites, mutations often lead to cross resistance to the other agent<sup>2</sup>
- NNRTI resistance accumulation can compromise the efficacy of second-generation NNRTIs<sup>3</sup>

1. Johnson VA, et al. Top Antivir Med 2011;19:156–54  
2. Delaugerre C, et al. J Med Virol 2001;65:445–48  
3. Ghosn J, et al. AIDS Rev 2009;11:165–73

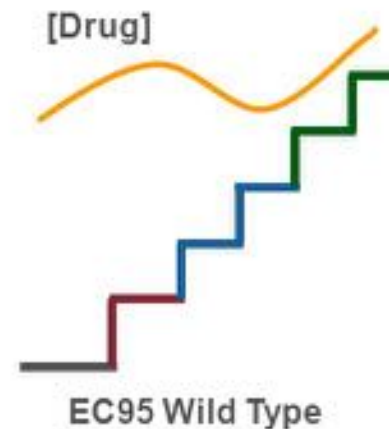
# Resistance not always so simple - NRTIs

*All or nothing;*

- Nevirapine and 3TC

*The more mutations the more resistance:*

- AZT
- Mutations (TAMS): M41L, D67N, K70R, L210W, T215Y/F, K219Q/E/N

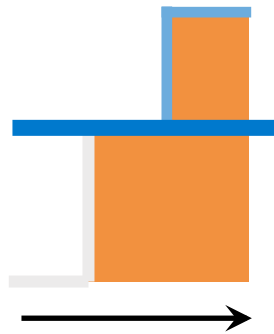


# Resistance: not always so simple – 2<sup>nd</sup> generation NNRTIs (etravirine).

The number of mutations required to substantially decrease the efficacy of an antiviral drug

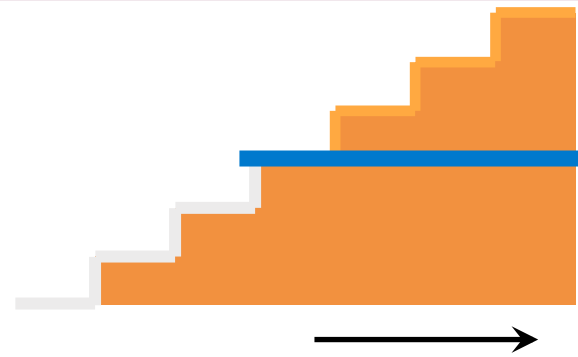
## First-generation NNRTI

One mutation correlates with reduced virological response



## Next-generation NNRTI (ETR)

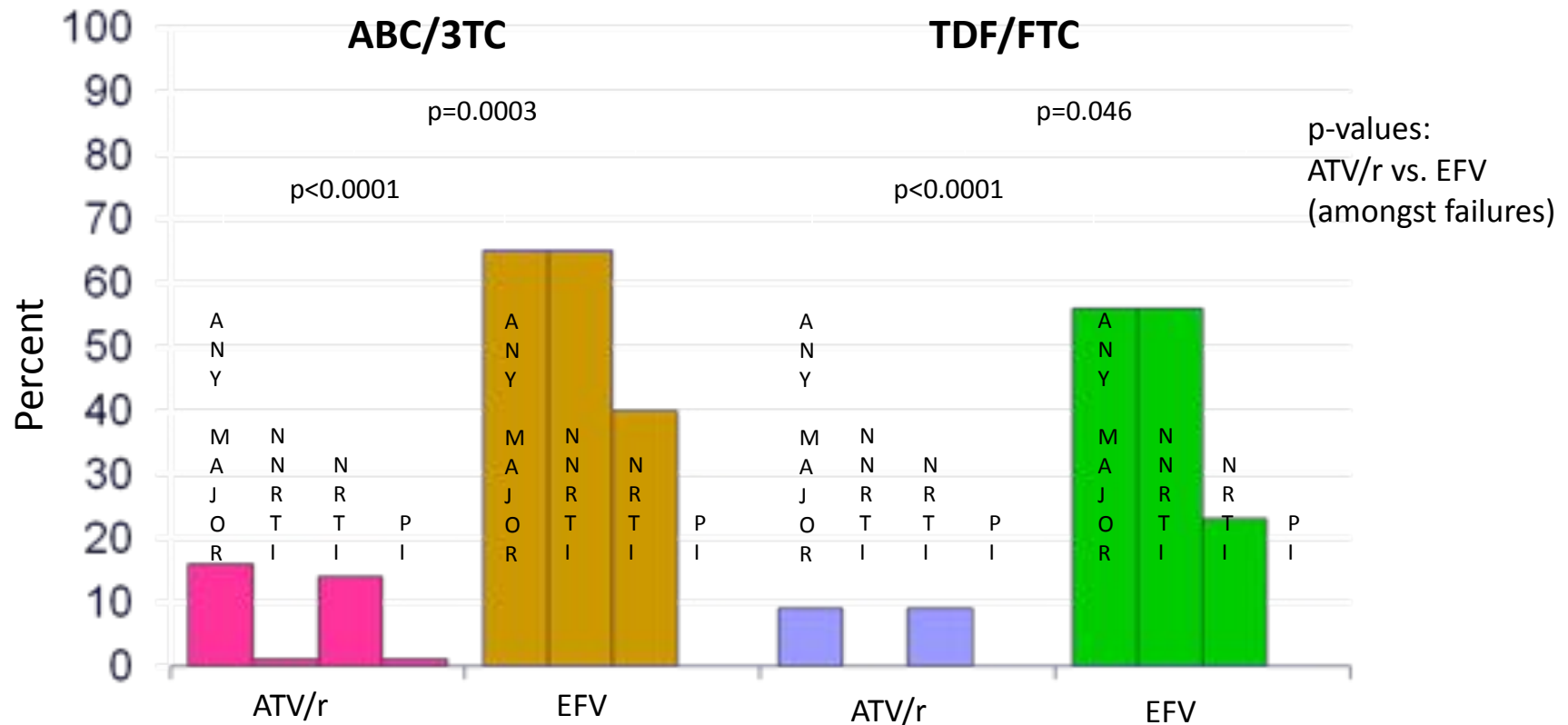
The presence of multiple NNRTI mutations at baseline is usually required to confer a reduced response



Increasing number of mutations at baseline

1. Antinori A, et al. AIDS Res Hum Retroviruses. 2002;18:835–8.
2. Lecossier D, et al. J Acquir Immune Defic Syndr. 2005;38:37–42.
3. Vingerhoets J, et al. 17th IDHRW 2008 [Poster 32].
4. De Béthune MP, et al. 4th EHDRW 2006 [Poster 51].
5. de Mendoza C, et al. HIV Clin Trials. 2006;7:163–71.

# Resistant to resistance: boosted PI's



## Viral failures

No baseline resistance N=      76      63      54      48

\*Major mutations defined by IAS-USA (2008) list plus T69D, L74I, G190C/E/Q/T/V for RT and L24I, F53L, I54V/A/T/S and G73C/S/T/A for PR

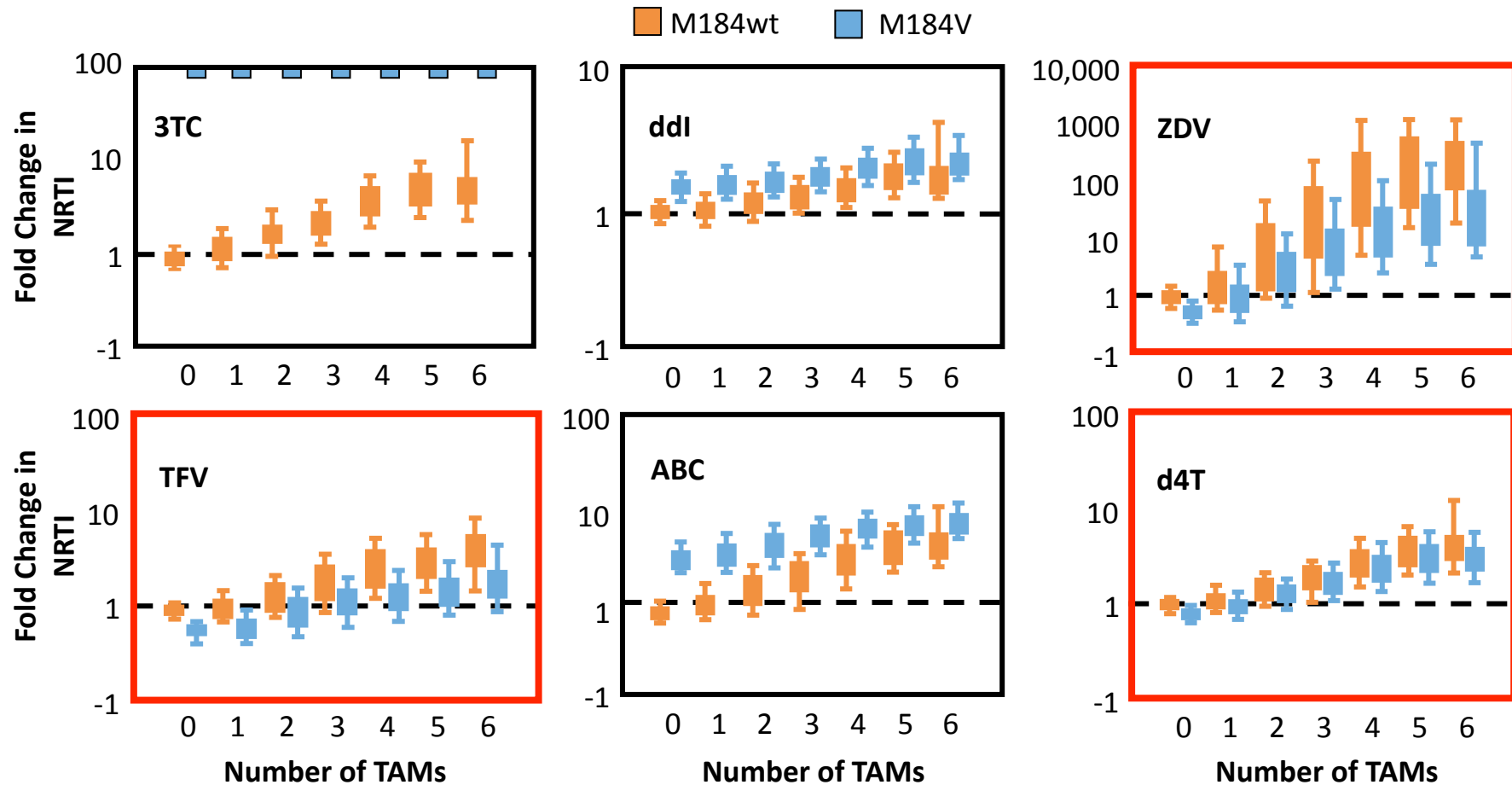
# Using resistance to your advantage

- Hypersusceptibility
  - A resistance to one ARV makes the virus even more susceptible to another.....
- Viral fitness
  - The resistance required to resist a drug interferes with other vital processes in the virus and it is not so 'replication-competent' ....

# M184V Increases Susceptibility to d4T, ZDV, and TDF

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		
NRTI	Abacavir	Ziagen	(4.5 - 6.5)	1.27		Y	Y	Sensitive	
	Didanosine	Videx	(1.3 - 2.2)	0.88		Y	Y	Sensitive	
	Emtricitabine	Emtriva	(3.5)	>MAX		N	N	Resistant	
	Lamivudine	Epivir	(3.5)	>MAX		N	N	Resistant	
	Stavudine	Zerit	(1.7)	0.65		Y	Y	Sensitive	3
	Zidovudine	Retrovir	(1.9)	0.26		Y	Y	Sensitive	2,3
	Tenofovir	Viread	(1.4 - 4)	0.31		Y	Y	Sensitive	2,3
	NRTI Mutations		M184V						

# Change in NRTI Susceptibility and Number of TAMs, $\pm$ M184V



# Doing it without access to a resistance test

The principle aim of the RDI is to provide a treatment decision-making aid free of charge over the Internet such that physicians entering the genotype and other baseline data for a patient will receive a report containing predictions of virological responses to a range of alternative antiretroviral combinations.



# No resistance test...

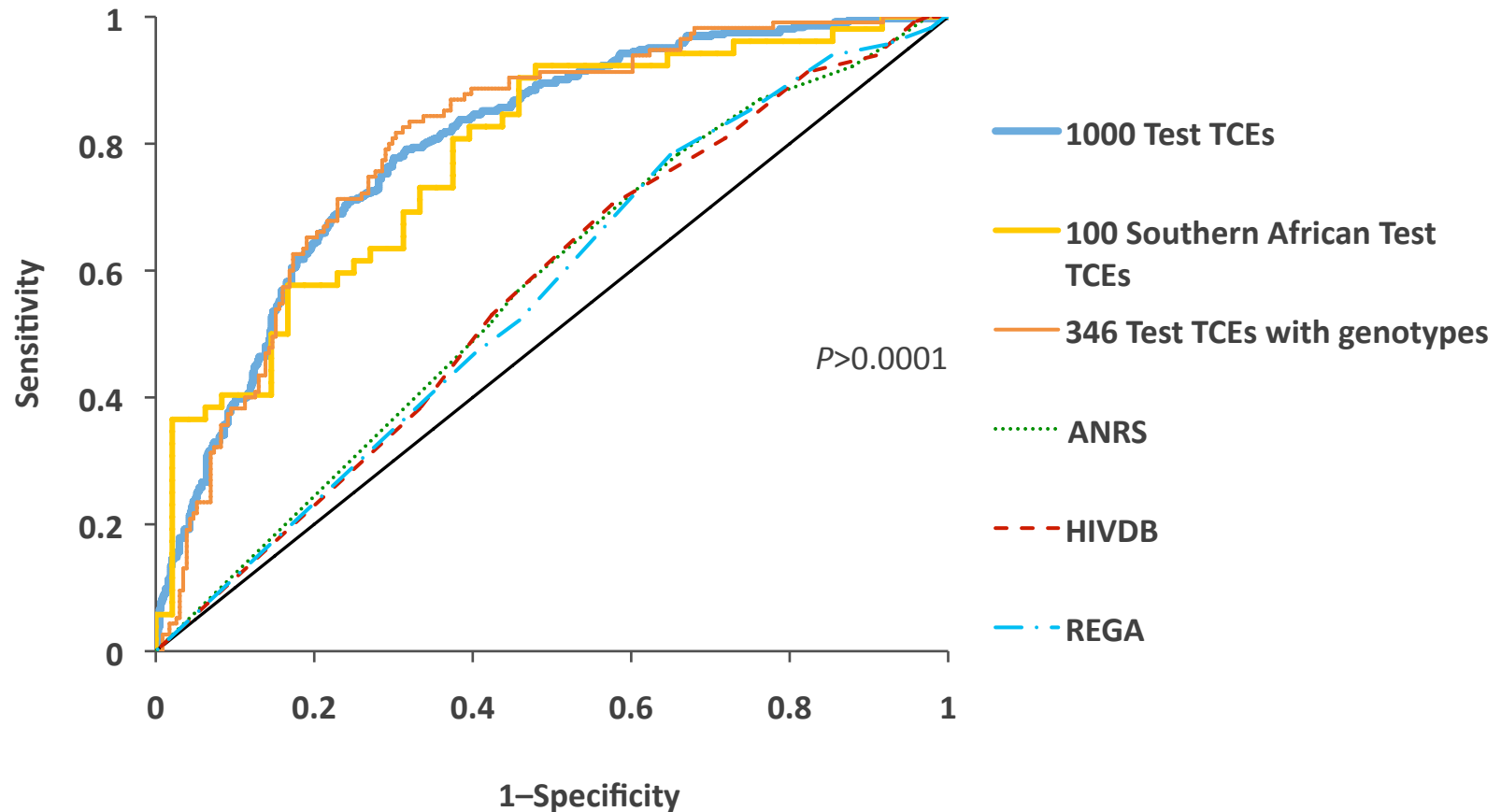
After failing NNRTI/2NRTIs

- Baseline viral load
- Treatment history
- Baseline CD4 count
- Time to follow-up

Resistance	Time	Accuracy (%)
Genotype	12w	66
<b>No genotype</b>	<b>12w</b>	<b>68</b>
Genotype	24w	65
<b>No genotype</b>	<b>24w</b>	<b>64</b>

- With or without genotype

# RDI models: Predicting treatment response without a genotype versus genotyping with interpretation



ANRS, Agence Nationale de Recherches sur le SIDA; RDI, Response Database Initiative; TCE, treatment change episode

Revell AD *et al. J Antimicrob Chemother* 2013; [Epub ahead of print]

# Managing without a resistance test

- Predict virological response to salvage ART accurately (approximately 80%) **without** the use of a genotype<sup>1</sup>
- Significantly more accurate predictors of response than genotyping with rules-based interpretation ( $P<0.001$ )<sup>1</sup>
- As accurate for cases from southern Africa as for other regions<sup>1</sup>
- Identify alternative regimens that are predicted to be effective for the majority of cases where the new regimen in the clinic failed<sup>1,2</sup>
- Identified cost-saving alternatives for most cases of failure in a study of second-line therapy in India<sup>2</sup>

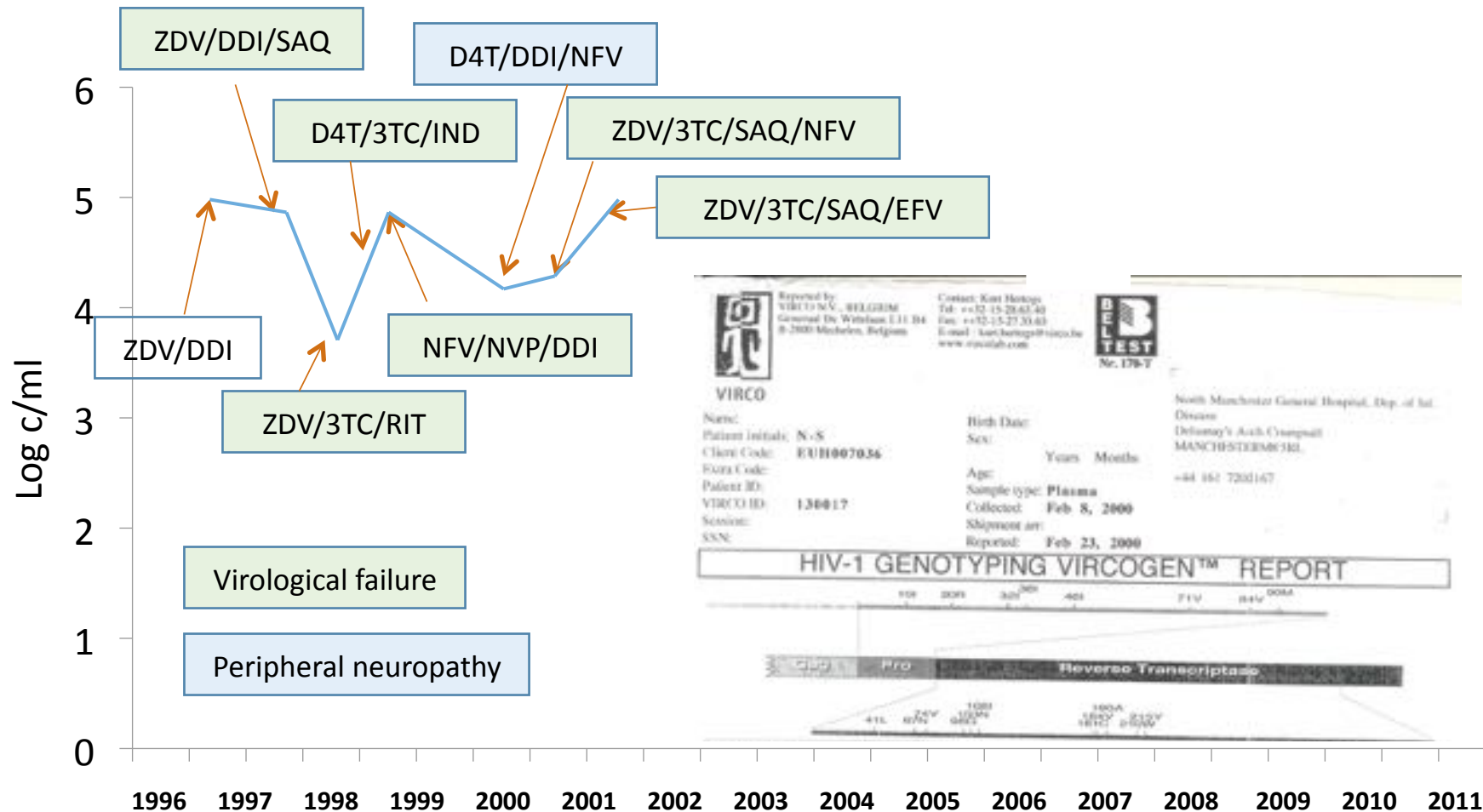
# Preventing resistance

1. Choose a good combination
  - (= 3 active drugs)
2. Ensure good levels
  - Compliance
  - Absorption
  - Interactions
3. Monitor for viral control & react quickly to failure
4. Build a robust new regimen on virological failure
  - Known resistance now
  - All previous resistance
  - *Guesstimate* other resistance
5. Go back to 2....

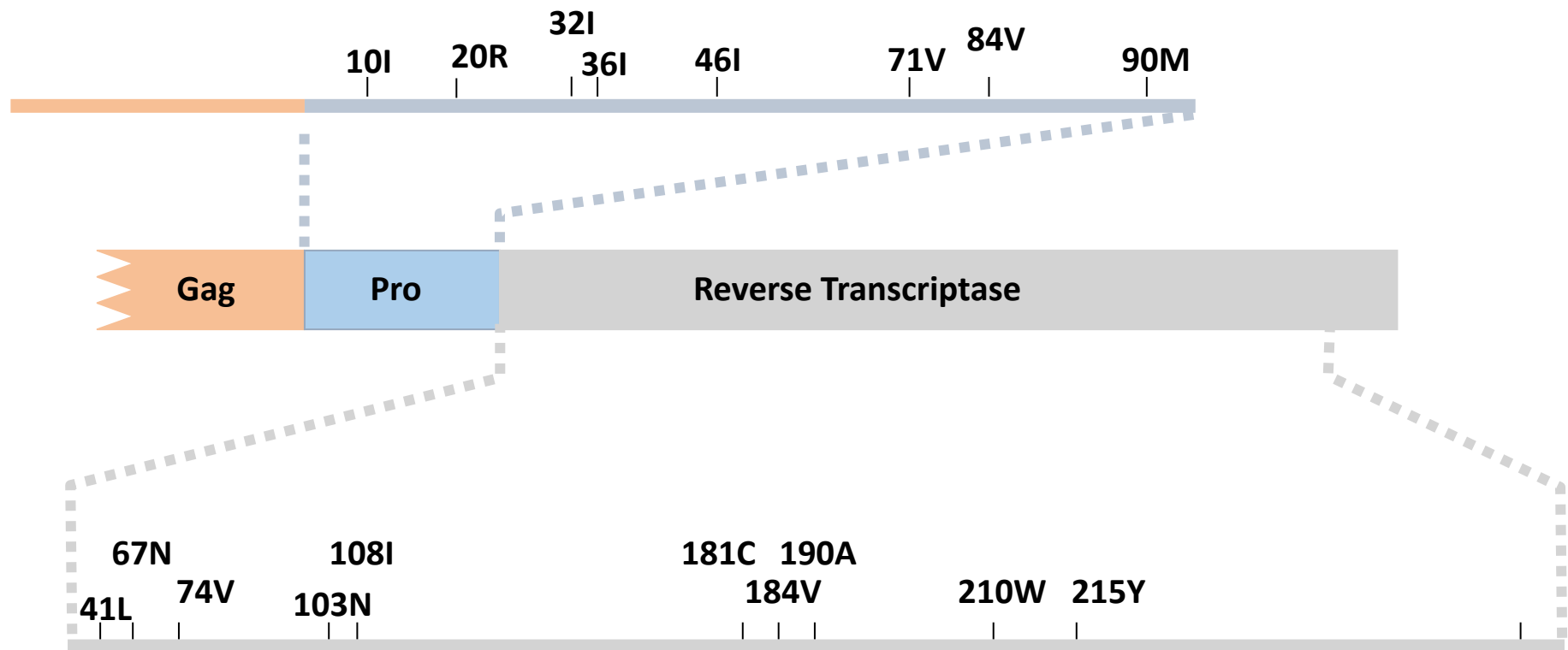
# Case 1

- 49yr old woman diagnosed 1995 (then 32)
  - PCP, CD4 50
  - Weight loss, OPC
- HBV/HCV –ve
- No significant co-morbidity
- CVD risk factors:
  - Non-smoker, No FHx, BMI 19
  - BP 120-5/80
  - TC 4.2, HDL-C 0.9
- Started ZDV/DDI

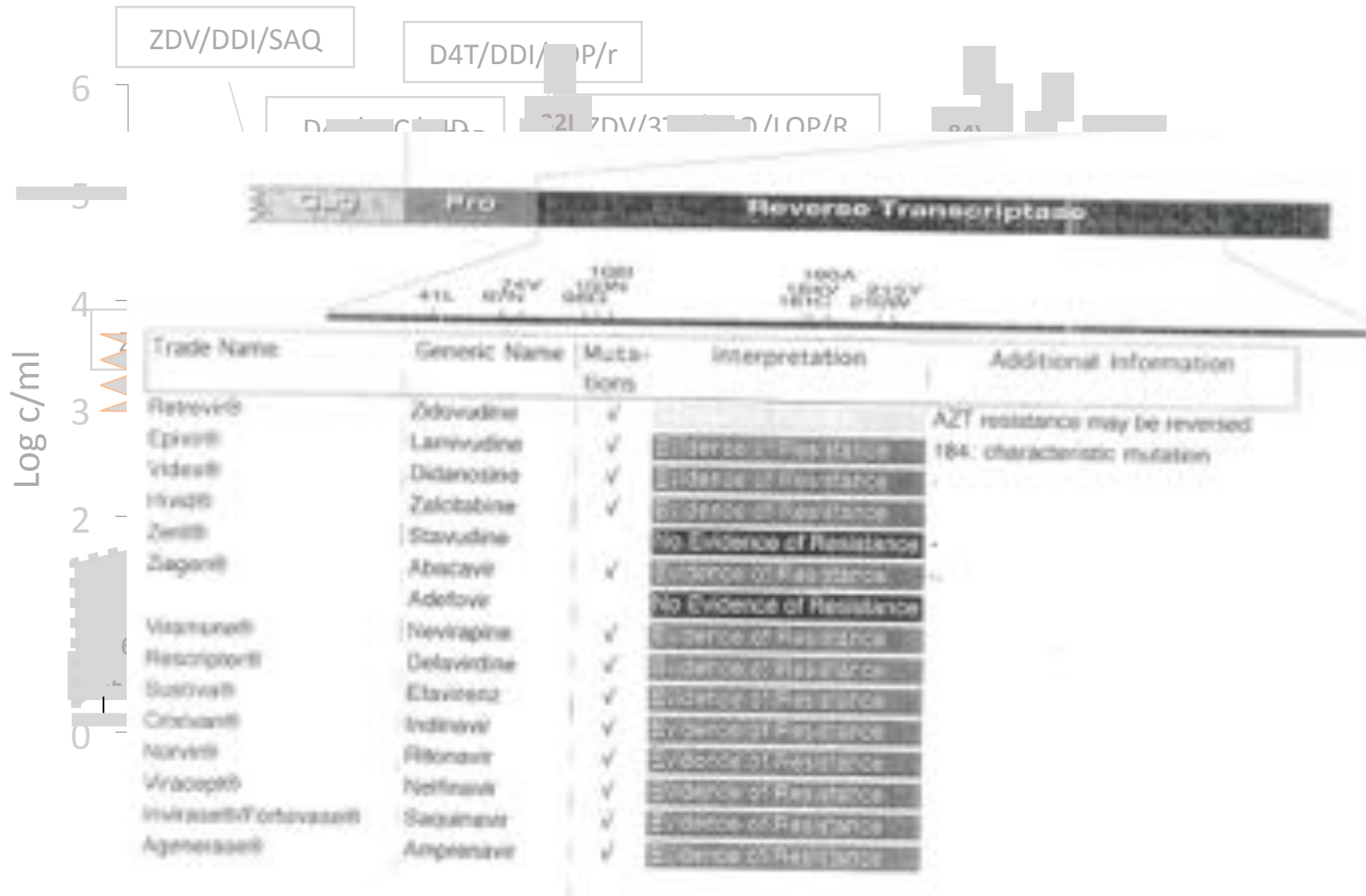
# Case 1



# Genotypic resistance test

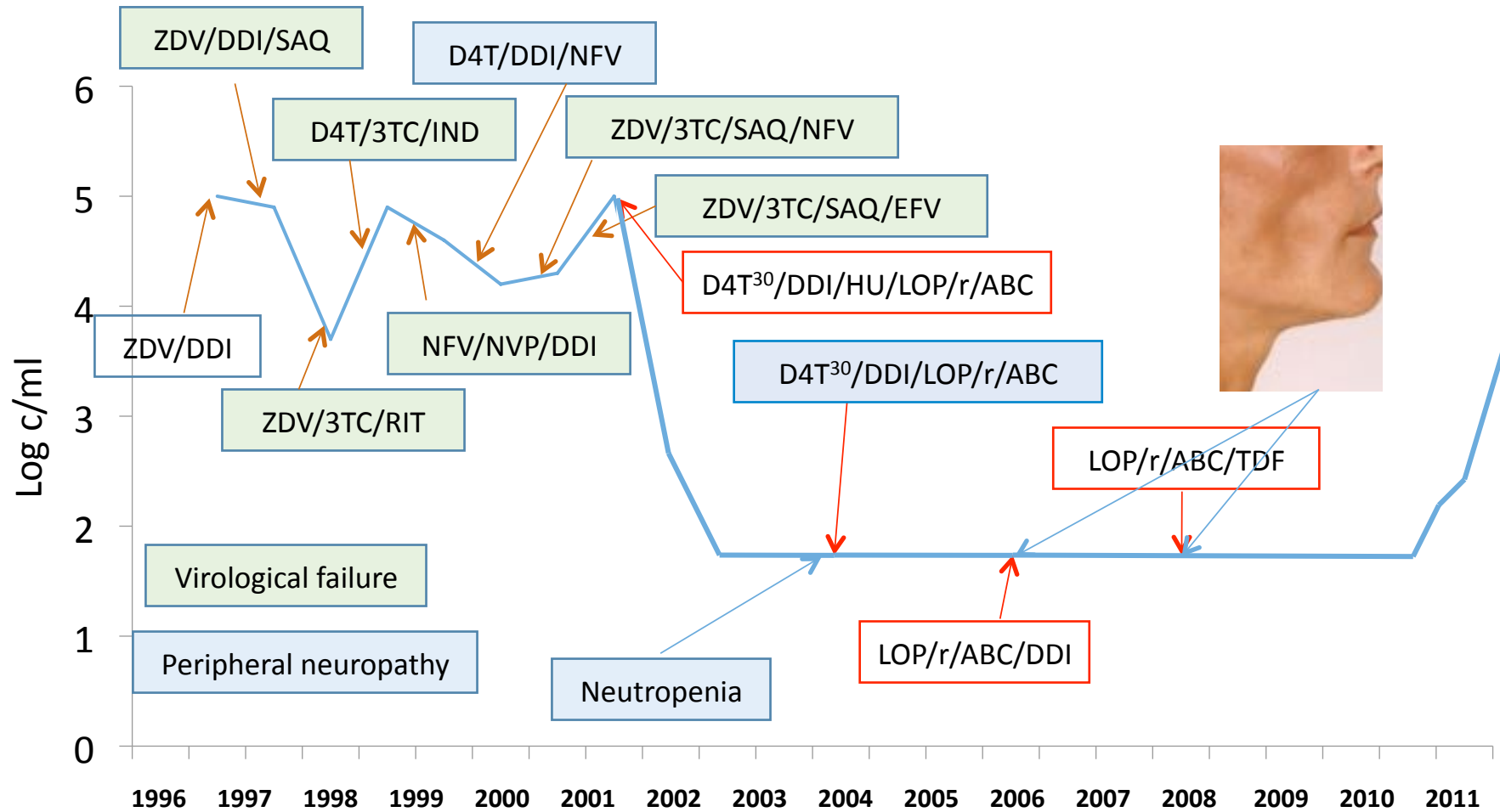


# Case 1





# Viral loads

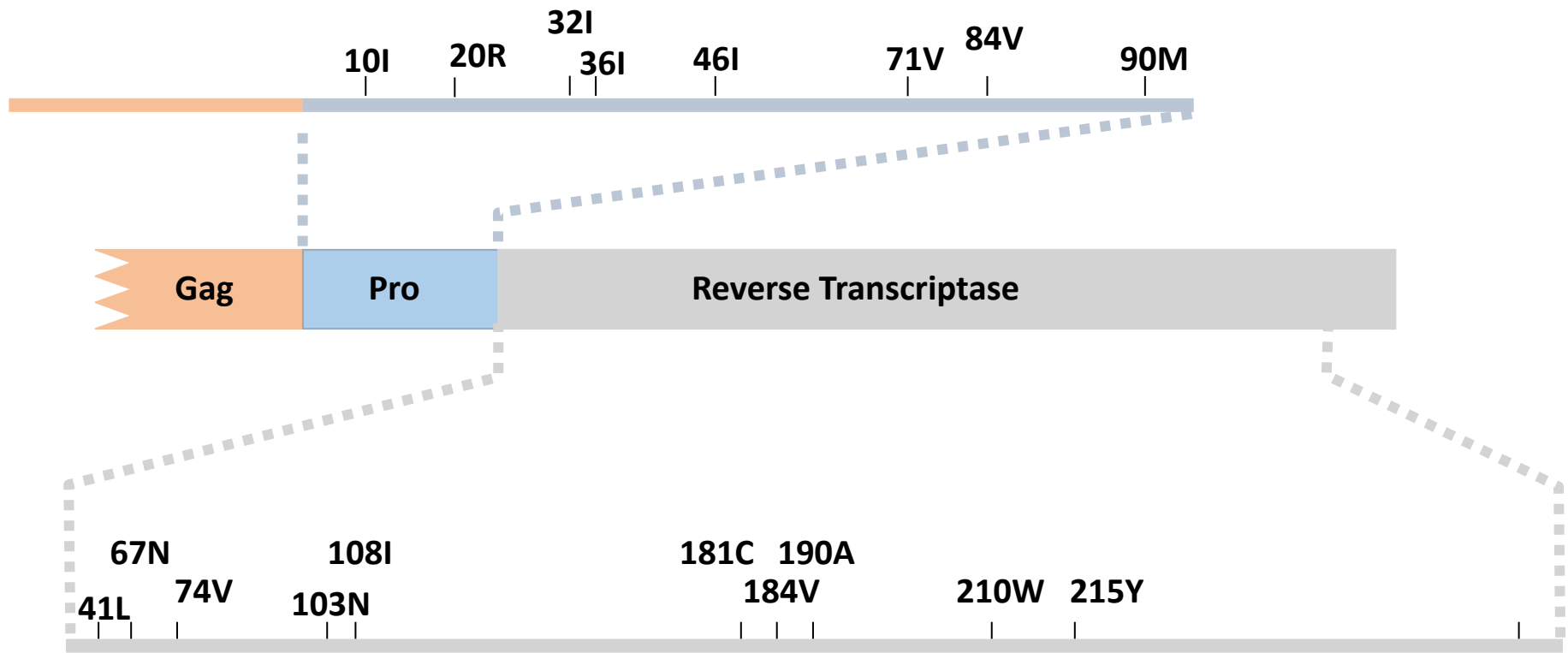


**Most likely cause?**

# Taking only 25% tablets

- Depressed
  - Life in a mess
- Diarrhoea
  - I've had enough!!!
- Aware of new agents
  - Wants to take fewer tablets
- Fed up with current combination
  - CD4 224, viral load 60,000
  - Tropism R5

# Going back to resistance test



# So what's available?

- Definitely active:

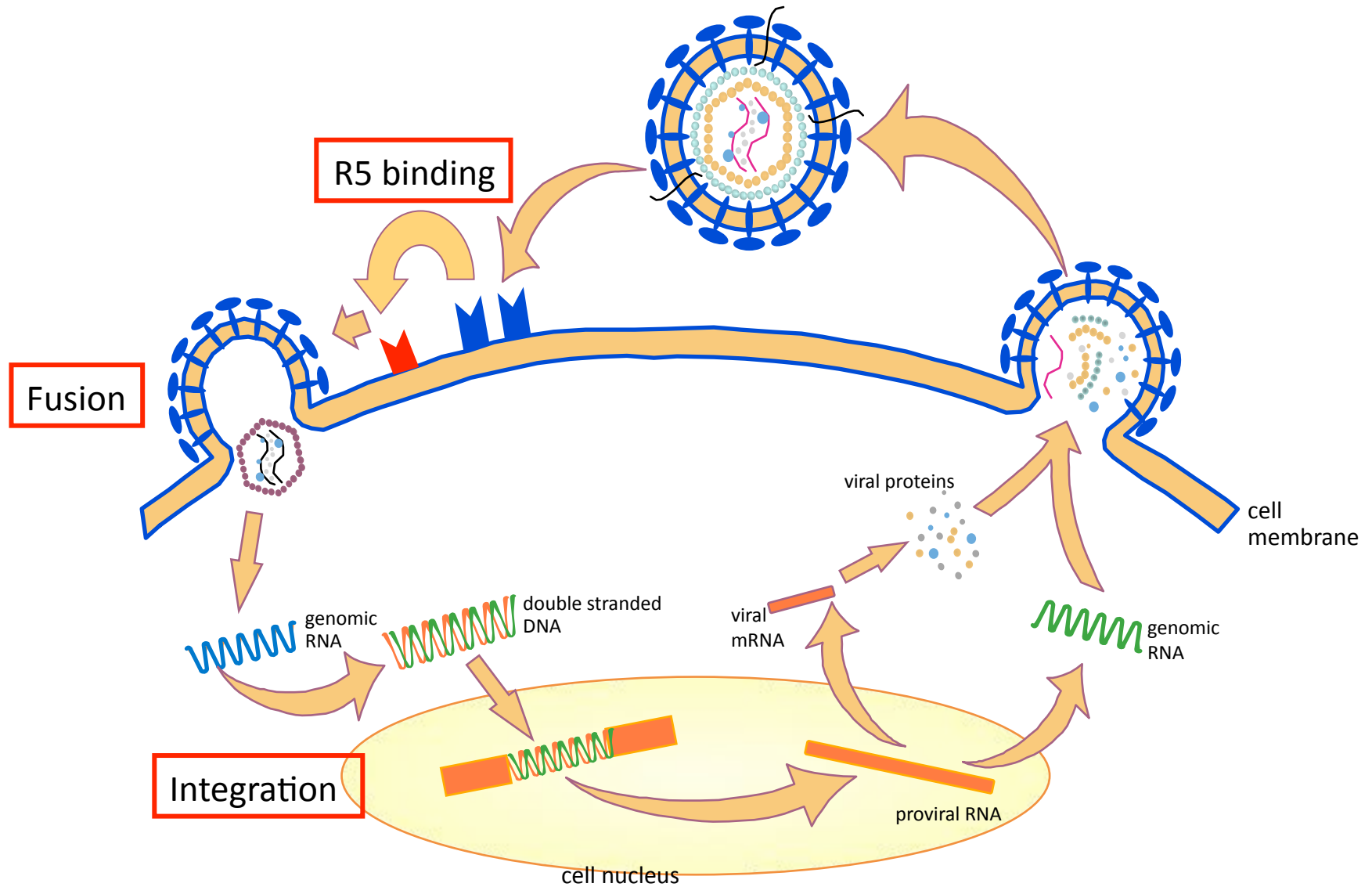
- RAL
- T-20
- MRV

BENCHMMRK

TORO

MOTIVATE

# Where do these act - life cycle of HIV



# So what's available?

- Reduced activity:

### Reverse Transcriptase

Enter Mutation List:

OR

Use The Pulldown Menus:

41	44	62	65
67	69	70	74
75	77	98	100
101	103	106	108
115	116	118	138
151	179	181	184
188	190	210	215
219	225	227	230
236	238	318	333

### Protease

Enter Mutation List:

OR

Use The Pulldown Menus:

10	11	13	15
20	23	24	30
32	33	35	36
43	45	47	48
50	53	54	58
60	62	63	69
71	73	74	78
77	82	83	84
85	88	89	90
93			

Identifier (Optional):

Date (Optional):

Output Analysis:

Mutation Scores       Mutation Comments



# Increasing predictive accuracy to DRV score by weighting mutations

Estimated increase in FC	<2	2 to 3	3 to 4	>4
Mutations	V11I <u>I54L</u> G73S L89V	<u>V32I</u> L33F I47V	I54M <u>L76V</u> <u>I84V</u>	<u>I50V</u>

Add mutations up for fold-change

Example: = ~ 5–7 fold-change = **Intermediate activity**

# Increasing predictive accuracy to TPV score by weighting mutations

- 1	0.5	1	2
L24I	L10V/I	V11L	I47V
D30N	I13V	V32I	I54A
I50L/V	K20R	A71L	V82T
154L	M46L	G73T	I84V
L76V	L90M	L89V	
V82I			

Add mutations up for fold-change

Example: = ~ 4 fold-change = **Intermediate activity**



STANFORD UNIVERSITY

# HIV DRUG RESISTANCE DATABASE

A curated public database designed to represent, store, and analyze the divergent forms of data underlying HIV drug resistance.

HOME

GENOTYPE-RX

GENOTYPE-PHENO

GENOTYPE-CLINICAL

HIVdb PROGRAM

## HIVdb: Genotypic Resistance Interpretation Algorithm

SeqID: Date Feb 2011

### Drug Resistance Interpretation

PI Major Resistance Mutations: V32I, M40I, I84V, L50M

PI Minor Resistance Mutations: L10L, A11V

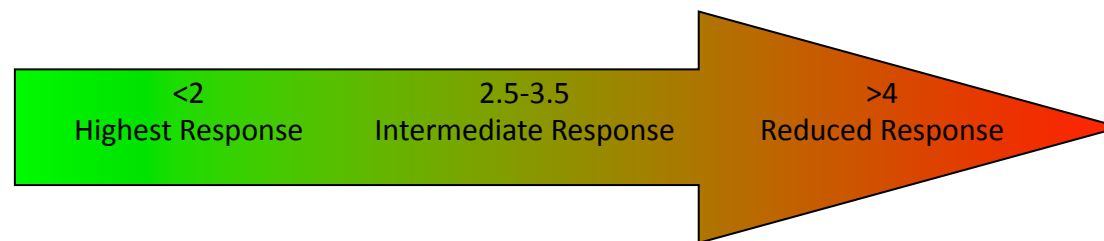
Other Mutations: K20R, M36I

	Protease Inhibitors
atazanavir (ATV)	High-level resistance
darunavir (DRV)	Intermediate resistance
fosamprenavir (FPV)	High-level resistance
indinavir (IDV)	High-level resistance
lopinavir (LPV)	Intermediate resistance
nelonavir (NFV)	High-level resistance
saquinavir (SQV)	High-level resistance
tipranavir (TPV)	Intermediate resistance



# Increasing predictive accuracy to ETR score by weighting mutations

0	1	1.5	2.5	3
<b>K103N</b>	<b>V90I</b>	<b>V106I</b>	<b>L100I</b>	<b>Y181I</b>
	<b>A98G</b>	<b>E138A</b>	<b>K101P</b>	<b>Y181V</b>
	<b>K101E</b>	<b>V179F</b>	<b>Y181C</b>	
	<b>K101H</b>	<b>G190S</b>	<b>M230L</b>	
	<b>V179D</b>			
	<b>V179T</b>			
	<b>G190A</b>			





## HIVdb: Genotypic Resistance Interpretation Algorithm

SeqID: Date Feb 2011

## Drug Resistance Interpretation

NRTI Resistance Mutations: M18I, D67N, L74V, L210W, T215Y  
 NNRTI Resistance Mutations: A30G, K103N, V108I, Y181C, G190A  
 Other Mutations: None

## Nucleoside RTI

## Non-Nucleoside RTI

lamivudine (3TC)	Potential low-level resistance
abacavir (ABC)	High-level resistance
zidovudine (AZT)	High-level resistance
stavudine (D4T)	High-level resistance
didanosine (DDI)	High-level resistance
emtricitabine (FTC)	Potential low-level resistance
tenofovir (TDF)	Intermediate resistance

delavirdine (DLV)	High-level resistance
efavirenz (EFV)	High-level resistance
etravirine (ETR)	High-level resistance
nevirapine (NVP)	High-level resistance



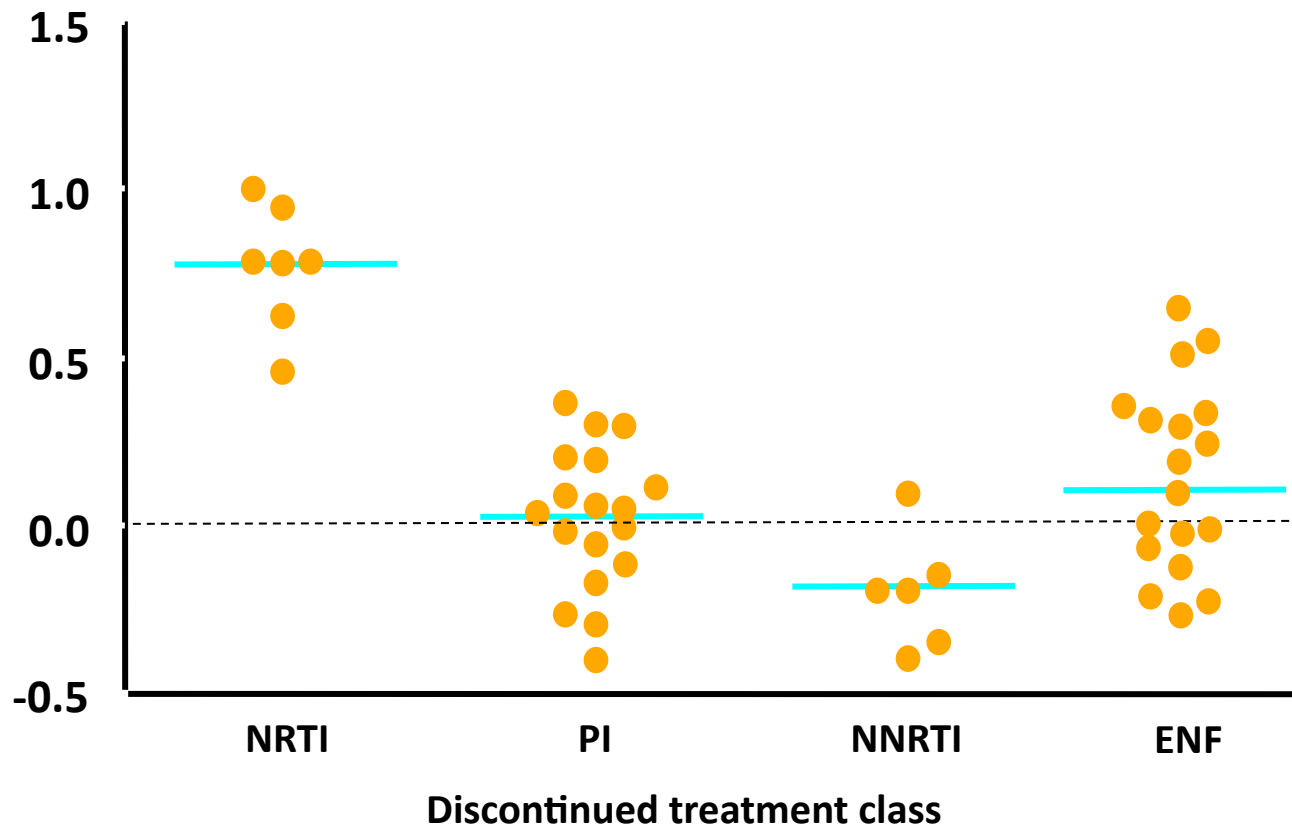
# So what's available?

- Definitely active:
  - RAL, MRV, T-20
- Reduced activity:
  - DAR/r & TIP/r
- Some benefit or not?
  - NRTI's, 3TC/FTC

# Partial treatment interruption studies

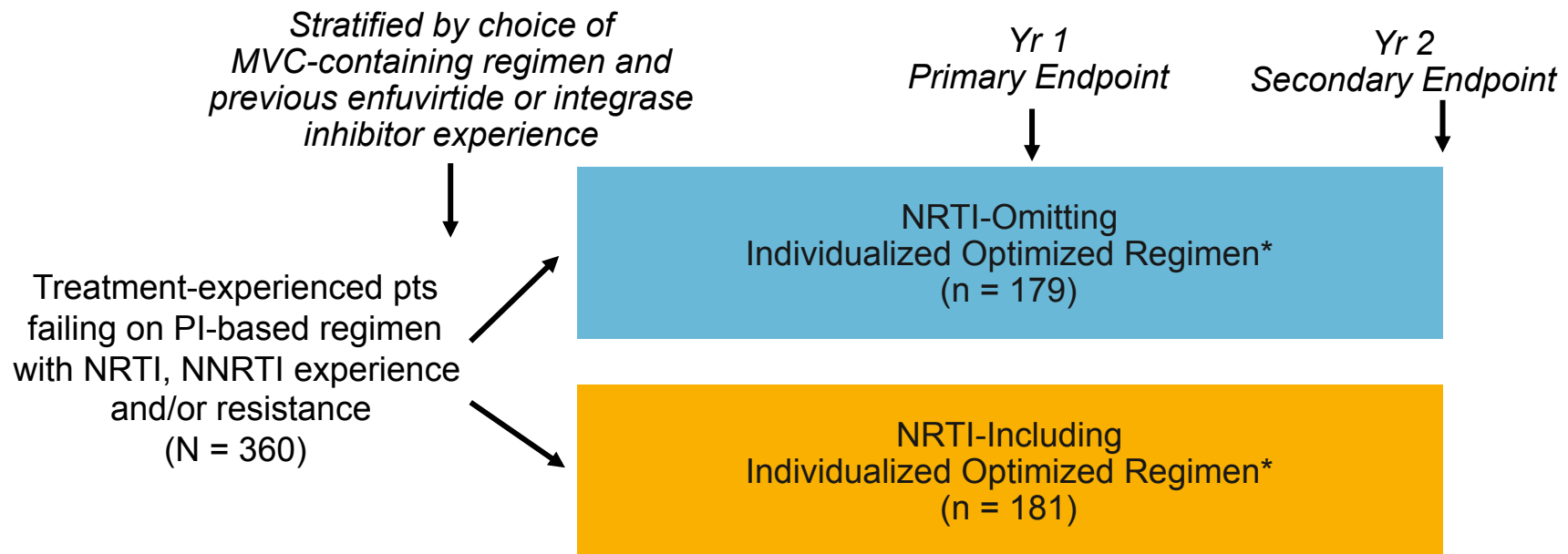
## Maintain NRTIs

Week 2 viral load increase after drugs in one class interrupted;  
other ARVs maintained



# OPTIONS: NRTIs vs No NRTIs in Regimens for Highly ART-Experienced

- Randomized, noninferiority, multicenter trial (ACTG A5241)
  - Primary endpoint: regimen failure (VF or divergence from NRTI assignment, whichever occurred first)

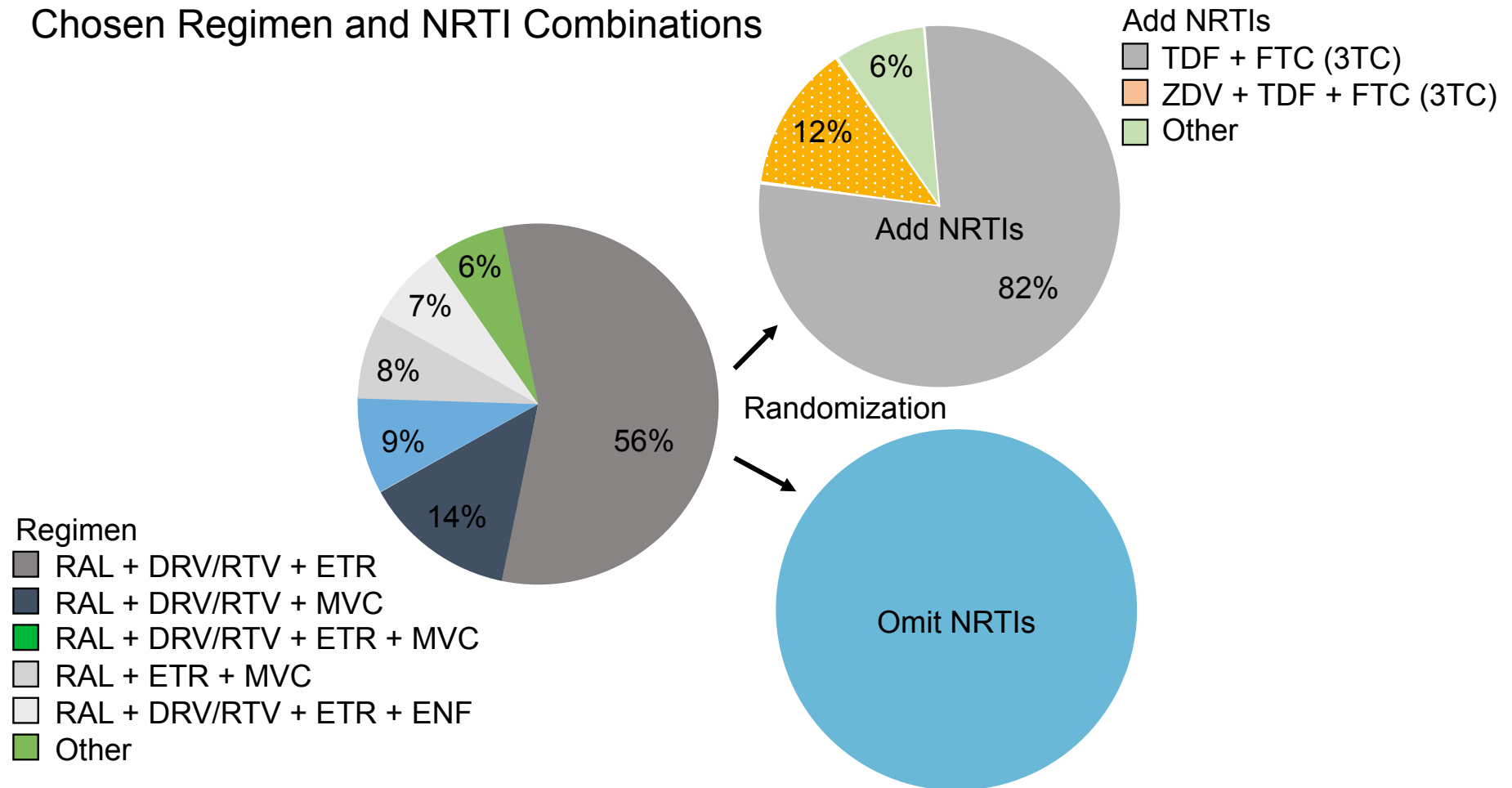


\*20 potential 3- to 4-drug combinations including DRV/RTV, ENF, ETR, MVC, RAL, TPV/RTV. Individualized selection of regimens with PSS > 2.



# OPTIONS: Pt-Specific Regimen Selected Then Randomized to $\pm$ NRTI

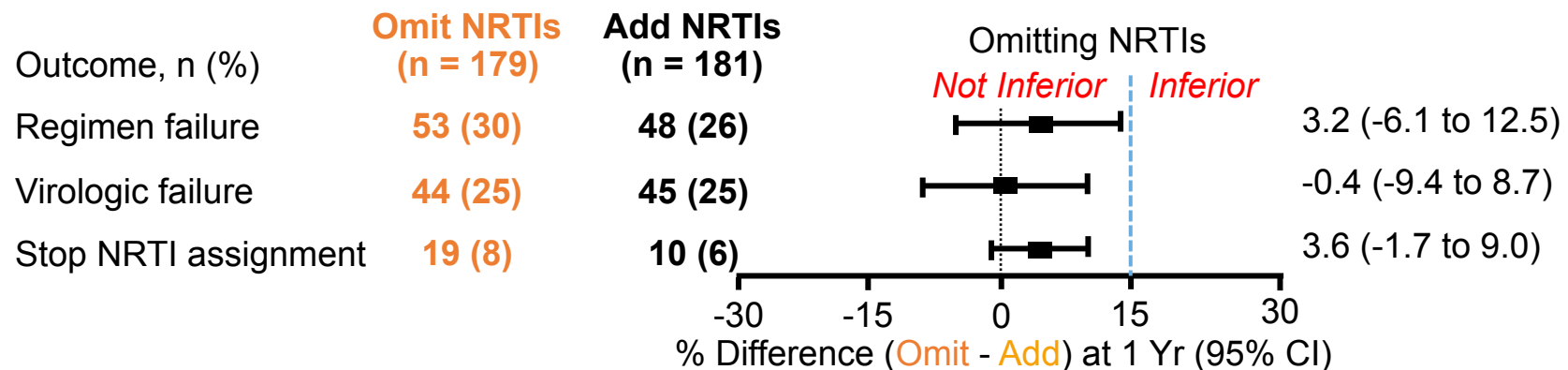
Chosen Regimen and NRTI Combinations



Tashima K, et al. CROI 2013. Abstract 153LB. Graphic used with permission.

# OPTIONS: Omitting NRTIs Non-inferior to Adding NRTIs to Optimized Regimen

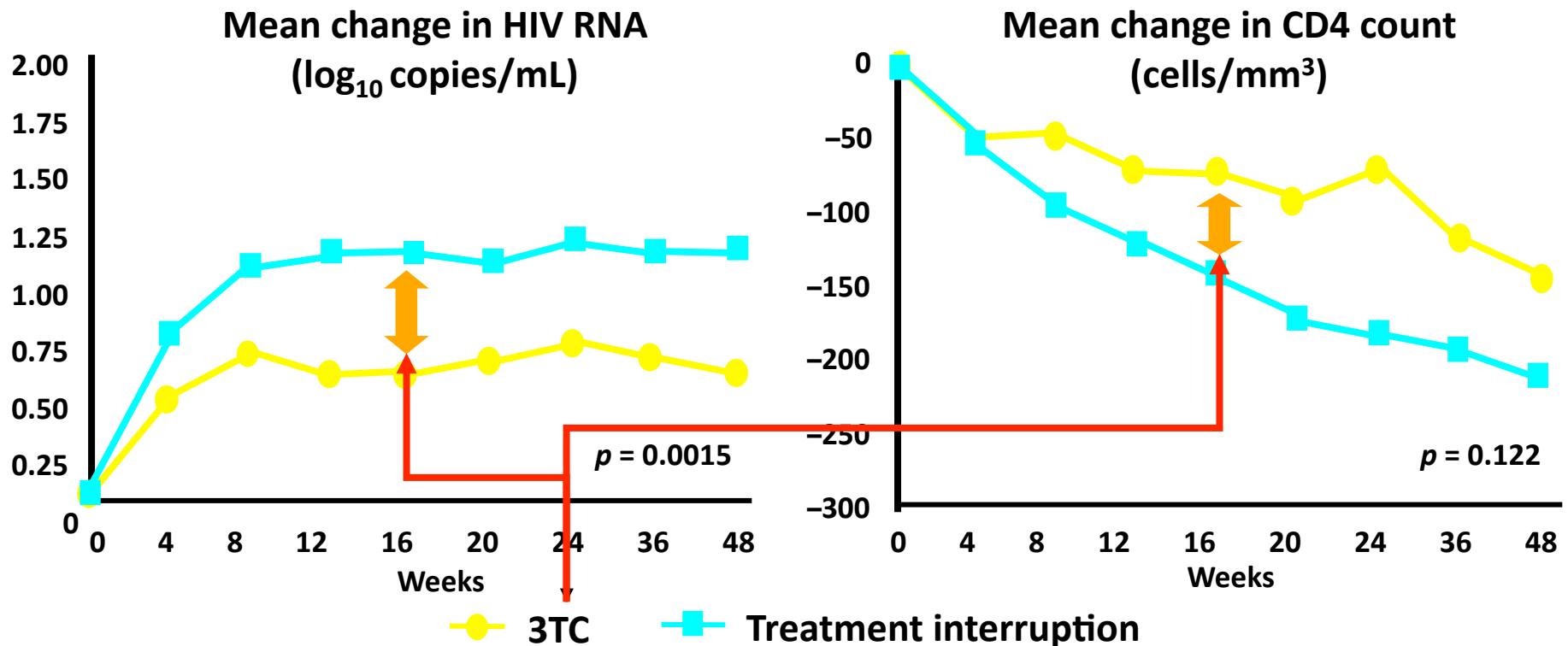
*Primary Efficacy Outcome Comparisons*



- Similar virologic suppression (HIV-1 RNA < 50 c/mL) in each arm (~ 65%)
- Similar CD4+ cell count increases in each arm (90-106 cells/mm<sup>3</sup>)
- No significant difference in any safety outcome when globally evaluating symptoms and laboratory abnormalities
  - However, mortality significantly higher in NRTI-added arm ( $P < .001$ )
  - 6 deaths in NRTI arm, 2 possibly due to ART drug

# Partial treatment interruption: 3TC monotherapy benefits over STI

- HIV RNA > 1000 copies/mL
- CD4 > 500 cells/mm<sup>3</sup>
- Have M184V mutation



# What are you going to do?

She is currently on LOP/r/TDF/ABC

She wants to change her ART

# Choices?

Fully active

Partially active

Some benefit

RAL

DAR/r

MRV

TIP/r

3/FTC

T-20

# Data for >1 active drug + OBR in triple-class failure

Overall Efficacy Data

Efficacy by ARTs in OBT

Enfuvirtide      Darunavir

+

+

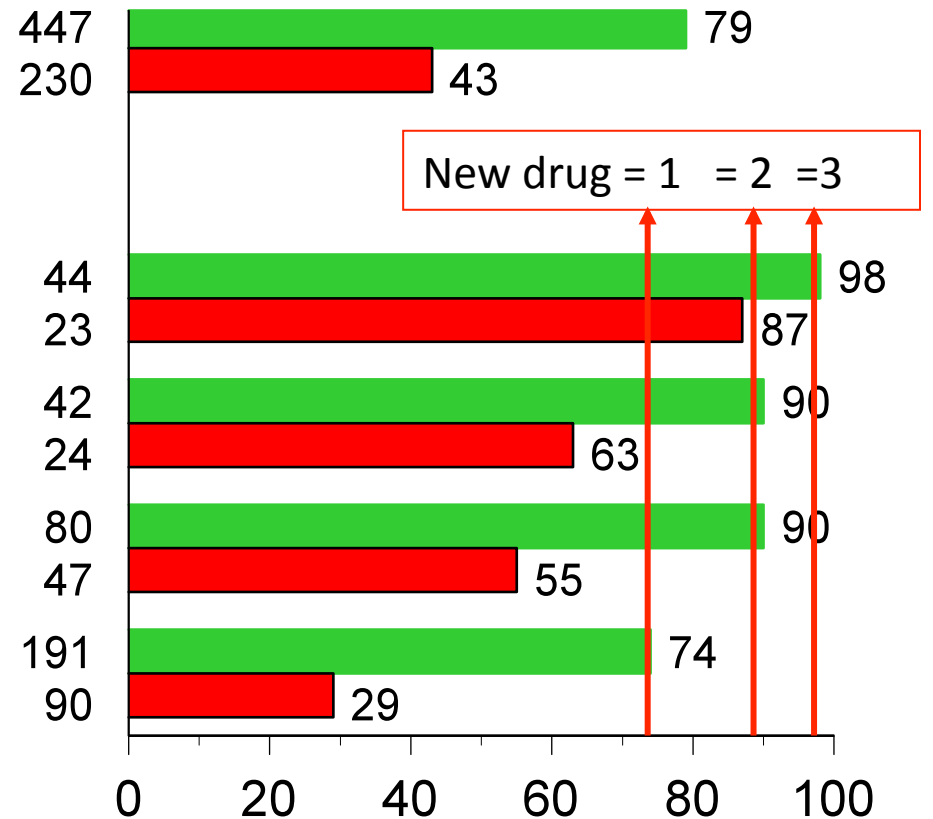
-

-

+ : First Use in OBT

- : No Use in OBT

\* Virological failures carried forward



■ Raltegravir + OBT      ■ Placebo + OBT

# Patient

- Received adherence support
- DAR/r
- RAL/MAR
- TDF/FTC
- Has been undetectable since

# Summary

- Antiretroviral resistance can seem confusing and complex
- A lot of it is actually quite simple
- People might bluff you with lots of numbers
  - You don't really need to know them as there are programmes to help
- It is important to understand:
  - How resistance develops
  - That it is archived forever
  - That you may not detect resistance even if it is there (look at previous regimens and if failed)



**If time for discussion**

# Case 1

- 49 yr old MSM
- HIV diagnosed 2006
  - CD4 = 95
- Commenced on Nevirapine and Truvada (Tenofovir/FTC)
  - Poor compliance → virologic failure
- Resistance testing:
  - K103N – NNRTI resistance
  - M184V – 3TC/FTC resistance

# Case 1 cont.

- Intensive counselling
  - Compliance issues addressed
- Commenced on Atazanavir/ritonavir & Truvada
  - Remains undetectable since, good CD4
- Wants a single tablet regimen....
  - We have:
    - Atripla = Efavirenz, Tenofovir, FTC
    - Eviplera = Rilpivirine, Tenofovir, FTC
    - QUAD in the future (Elvitegravir, Cobicistat, Tenofovir, FTC)
- Can he have any of these?
- What are the issues?

# Case 2

- 37 year old woman
- HIV positive, CD4 157
  - Referred from elsewhere
  - Diagnosed 6 weeks before, all other tests and baselines 'normal'
- Commenced on Atripla (Efavirenz, Tenofovir, FTC)
- Viral load decreased from 780,000 to 12,000 over 6 weeks but then rebounded...
- Why?

# Case 2 cont.

- Possibilities:
  - Poor compliance
  - Poor absorption / PK interactions
  - Virus was resistant already....
  - Something else
- In fact baseline resistance test was abnormal
  - Primary K103N resistance...
    - Which knocked out Efavirenz

# Case 3

- 47 yr MSM
- HIV positive since 1991
  - Regimens:
    - AZT 1992 for two years
    - D4T, 3TC and Indinavir Oct '97 to Nov '97
    - D4T, 3TC and Saquinavir Nov '97 to March '98
    - DDI, Nevirapine, AZT and Indinavir Oct '98 to Feb '99
    - DDI, Nevirapine, AZT, Nelfinavir and Ritonavir Feb '99 to Apr '99
    - DDI, Nevirapine, AZT and Nelfinavir Apr '99 to Dec 2000
    - AZT, 3TC and Abacavir Feb '01 to Aug '02
    - Lopinavir/ritonavir, 3TC and Abacavir till 1 month ago, when stopped treatment
- Resistance test now wild-type
- What more do I need to know and what can I use?

# Case 3 cont.

- Any previous resistance tests?
  - No
- Why did they switch previously - ?intolerance ? AEs
  - Generally from failure
- Therefore have to guesstimate resistance...
  - Multiple NRTI resistance
  - First-line NNRTI resistance
  - (Possible some PI resistance)

# Case 4

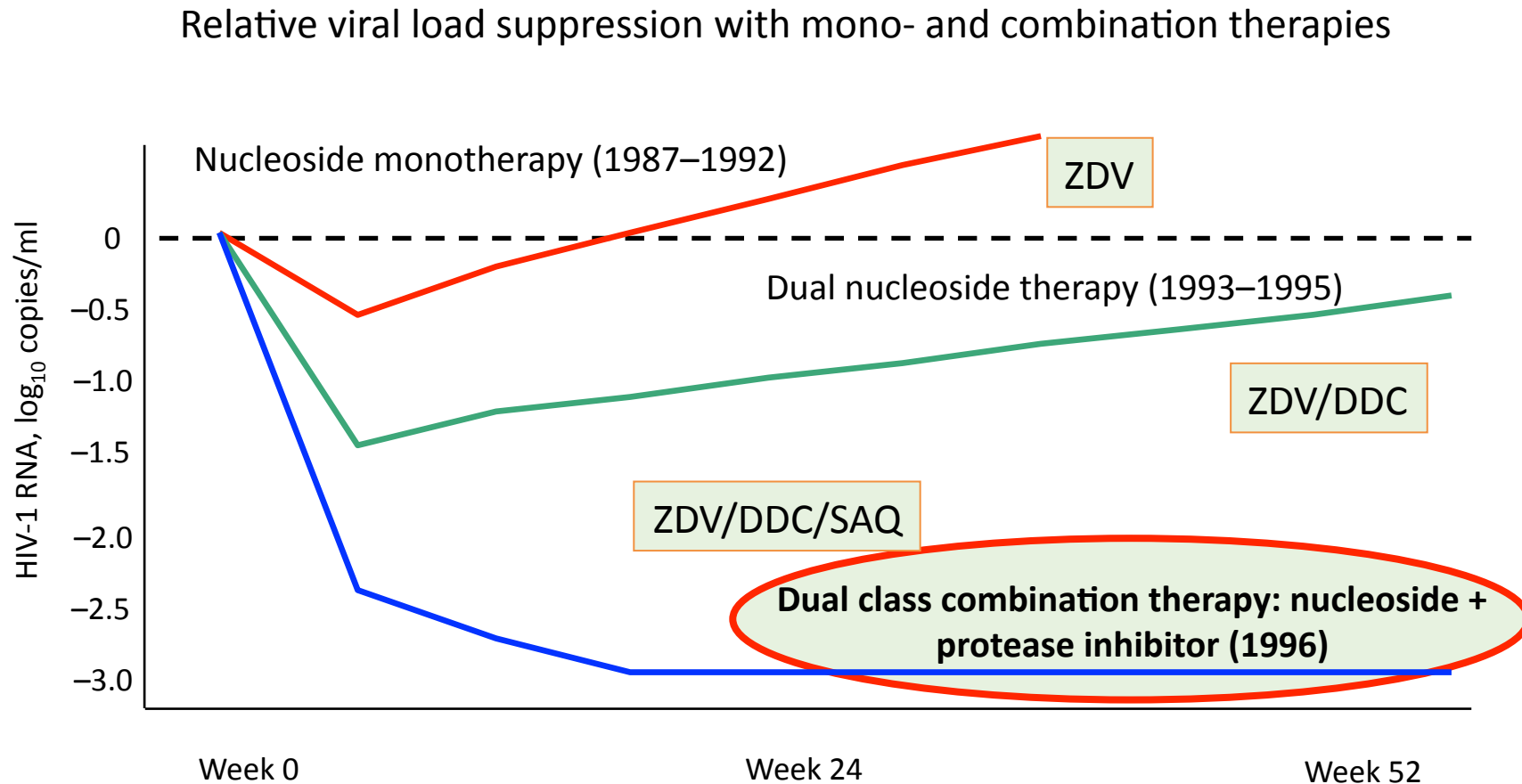
- 27 yr woman
  - HIV positive 2009
- Commenced Efavirenz, AZT, 3TC
  - Complied and was undetectable for 2 years
- Then lost job, got fed up etc. and stopped therapy
  - Now off therapy for 1 year
  - Resistance test wild-type
- Do I have to be wary of any resistance?
- May have had a 'functional monotherapy' for a while...



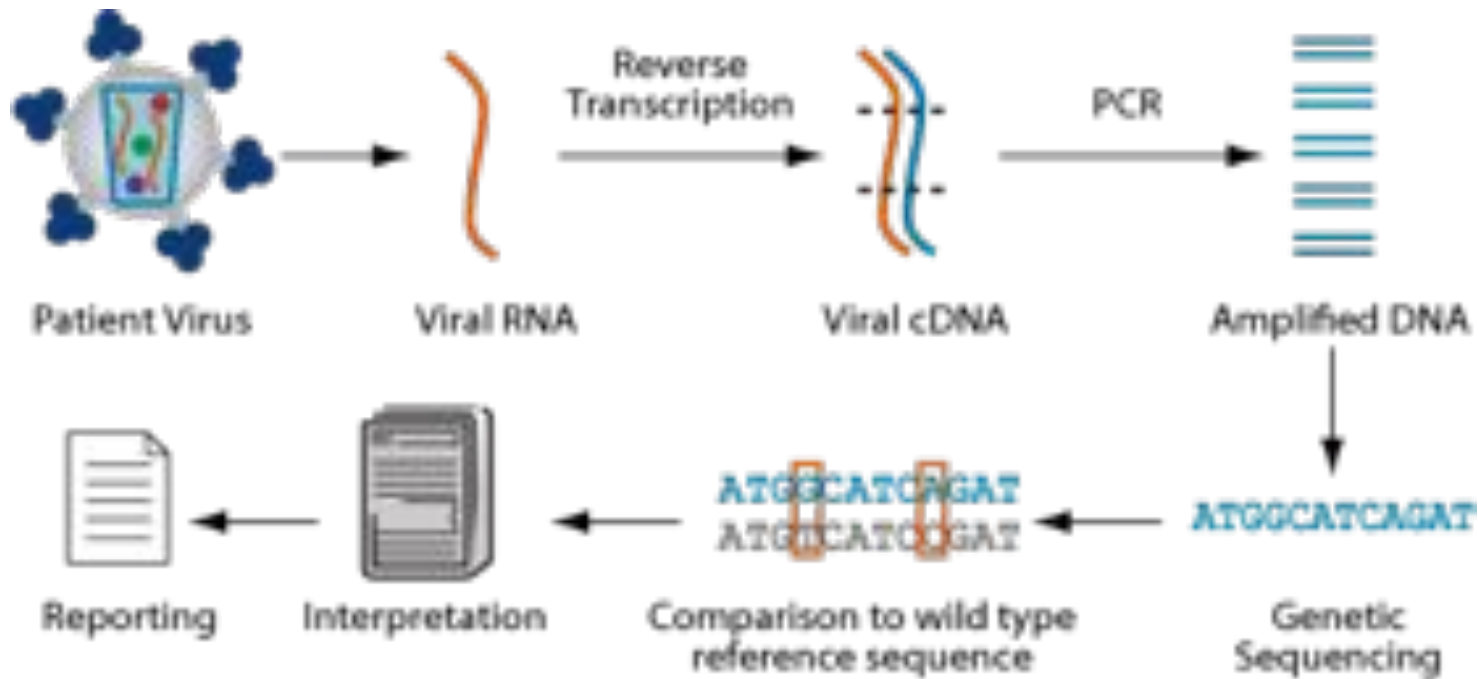
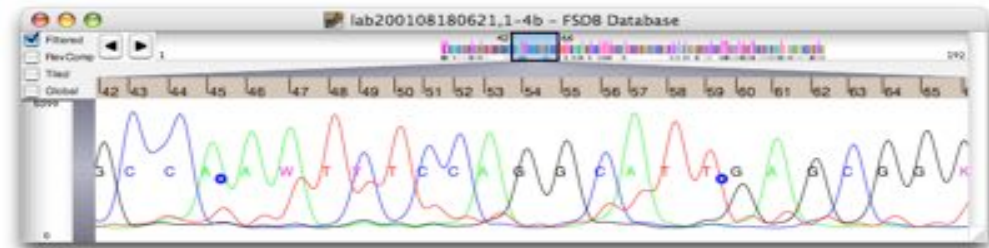
Data was limited



# And ART was still a pipedream



# You request a resistance test - how does it work?



# And how do you interpret?

- **Genotypic Testing:** Prediction of phenotype based on sequence

**STANFORD UNIVERSITY HIV DRUG RESISTANCE DATABASE**  
*Expanded public database designed to represent, store, and analyze the divergent forms of data underlying HIV drug resistance.*

HOME GENOTYPE-SEX GENOTYPE-PRR80 GENOTYPE-CLINICAL HIVdb PROGRAM

### HIVdb Program: Mutation List Analysis

Protease, RT, and integrase mutations can be entered using either the text box or pull-down menus (Detailed usage is found below).

The output can then be customized to display mutation comments, mutation scores, and an optional identifier and date. For further explanations and sample datasets please see the [Detailed Usage](#).

Reverse Transcriptase	Protease	Integrase																																																												
Enter Mutation List	Enter Mutation List	Enter Mutation List																																																												
OR	OR	OR																																																												
Use The Pull-down Menus:	Use The Pull-down Menus:	Use The Pull-down Menus:																																																												
<table border="1"> <tr><td>41</td><td>44</td><td>53</td><td>65</td></tr> <tr><td>67</td><td>69</td><td>78</td><td>80</td></tr> <tr><td>75</td><td>77</td><td>88</td><td>92</td></tr> <tr><td>100</td><td>101</td><td>106</td><td>114</td></tr> <tr><td>108</td><td>115</td><td>118</td><td>119</td></tr> </table>	41	44	53	65	67	69	78	80	75	77	88	92	100	101	106	114	108	115	118	119	<table border="1"> <tr><td>10</td><td>11</td><td>13</td><td>16</td></tr> <tr><td>20</td><td>23</td><td>24</td><td>30</td></tr> <tr><td>32</td><td>33</td><td>35</td><td>36</td></tr> <tr><td>43</td><td>45</td><td>47</td><td>48</td></tr> <tr><td>50</td><td>53</td><td>54</td><td>58</td></tr> </table>	10	11	13	16	20	23	24	30	32	33	35	36	43	45	47	48	50	53	54	58	<table border="1"> <tr><td>51</td><td>54</td><td>55</td><td>68</td></tr> <tr><td>74</td><td>92</td><td>96</td><td>97</td></tr> <tr><td>114</td><td>121</td><td>125</td><td>128</td></tr> <tr><td>138</td><td>140</td><td>143</td><td>145</td></tr> <tr><td>146</td><td>147</td><td>148</td><td>151</td></tr> </table>	51	54	55	68	74	92	96	97	114	121	125	128	138	140	143	145	146	147	148	151
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**TRUGENE<sup>®</sup> HIV-1**  
*Genotyping Test*  
**Guidelines<sup>™</sup> Rules 13.0**  
**RESISTANCE REPORT**

Sample ID: 185C060124  
 Patient ID: 1013drew  
 Patient Name: John Doe  
 Date Drawn: 20030606  
 Physician: Dr. Jane Doe  
 Institution: City Hospital  
 Report Date: 2007/11/12

State Central Labs  
 22 State St. Stateville, IL 61878

Bayer Reference Testing Laboratory  
 Example Report  
 725 Potter Street (APC3)  
 Berkeley, CA 94710  
 Tel: 900-434-2447  
 Fax: 510-705-5902

Resistance associated RT Mutations: L100I, K103N, T215S\*Y

Nucleoside and Nucleotide RT Inhibitors	Resistance Interpretation
abacavir (ABC)	No Evidence of Resistance
didanosine (ddI)	No Evidence of Resistance
lamivudine (3TC)/emtricitabine (FTC)	No Evidence of Resistance
stavudine (d4T)	Resistance
tenofovir (TDF)	No Evidence of Resistance
zidovudine (AZT)	Resistance

NonNucleoside RT Inhibitors	Resistance Interpretation
efavirenz (EFV)	Resistance
nevirapine (NVP)	Resistance

Resistance associated PR Mutations: L19I, M46L\*, L63P, A71T

Protease Inhibitors	Resistance Interpretation
amprenavir (APV)/fosamprenavir (FPV)	Resistance
APV or FPV**	Resistance
atazanavir (ATV)	No Evidence of Resistance
ATV**	No Evidence of Resistance
darunavir + ritonavir (DRV/r)	No Evidence of Resistance
indinavir (IDV)	Resistance
IDV**	Possible Resistance
lopinavir + ritonavir (LPV/r)	No Evidence of Resistance
nefinavir (NFV)	Possible Resistance
saquinavir + ritonavir (SQV/r)	No Evidence of Resistance
tipranavir + ritonavir (TPV/r)	No Evidence of Resistance

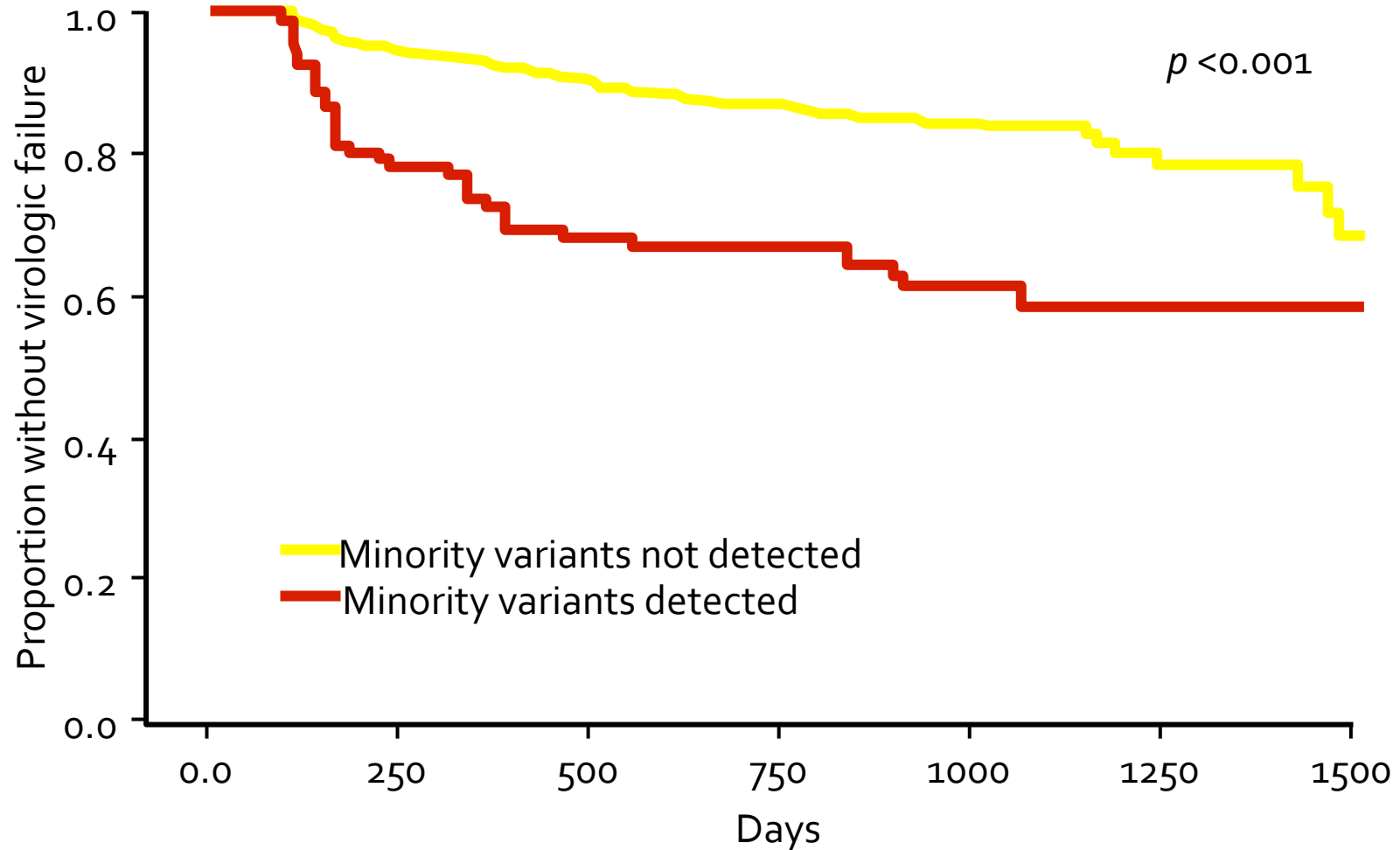
\*\* Protease Inhibitors administered with low-dose ritonavir for pharmacological boosting.

Resistance interpretation is based upon interpretation by an international expert panel (The Consensus Panel) of all *in vitro* and *in vivo* data including phenotypic and virologic response data available as of June 2007 for combination of Protease and RT sequences to antiretroviral drug resistance. These include primary and secondary mutations.

# Limitations of resistance testing

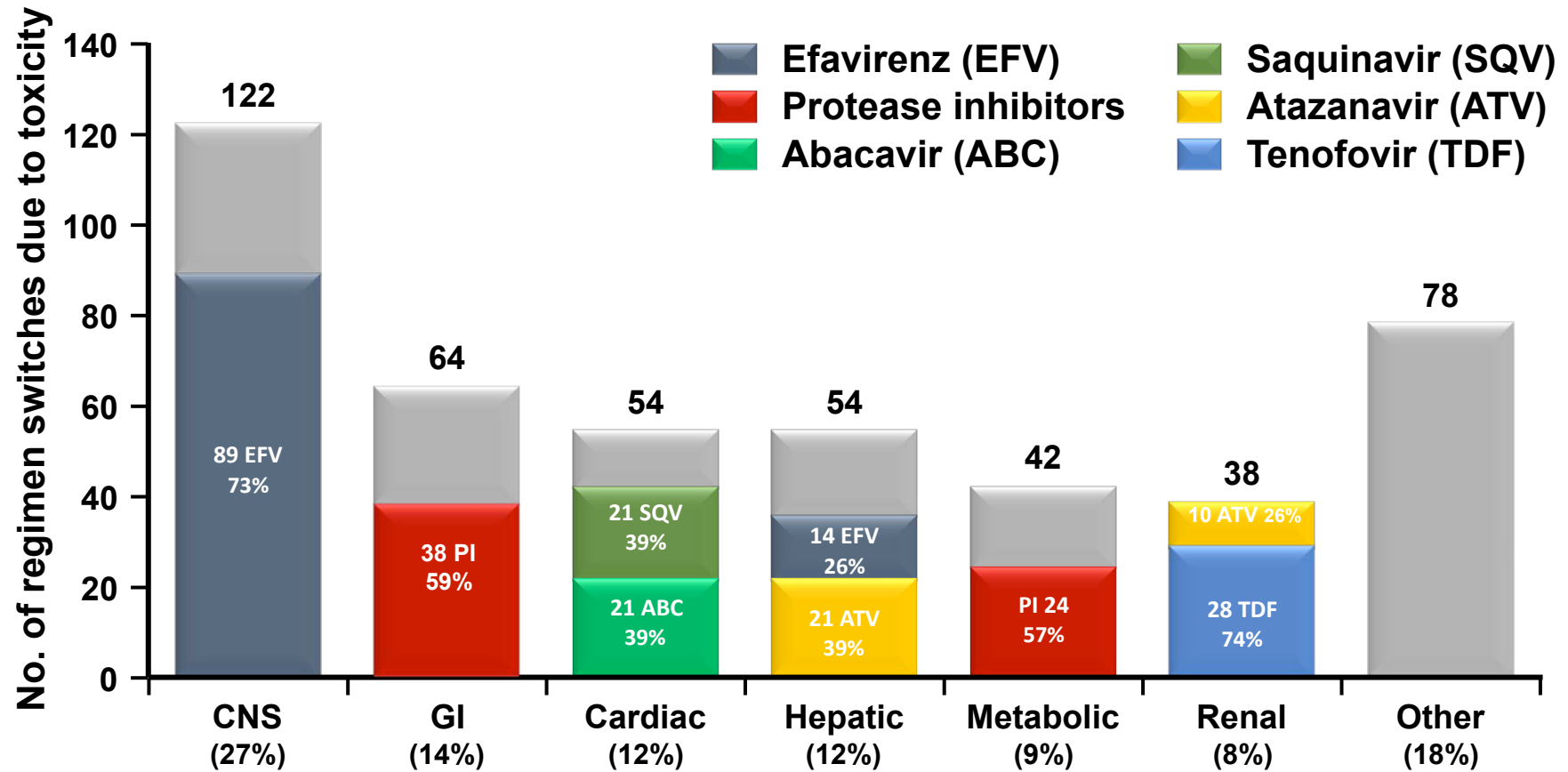
- Population sequencing
  - Standard resistance testing will only detect mutations that are in >20% of the circulating virus

# Kaplan-Meier curves for the proportion of patients without virologic failure



# Drugs still main reason to switch...

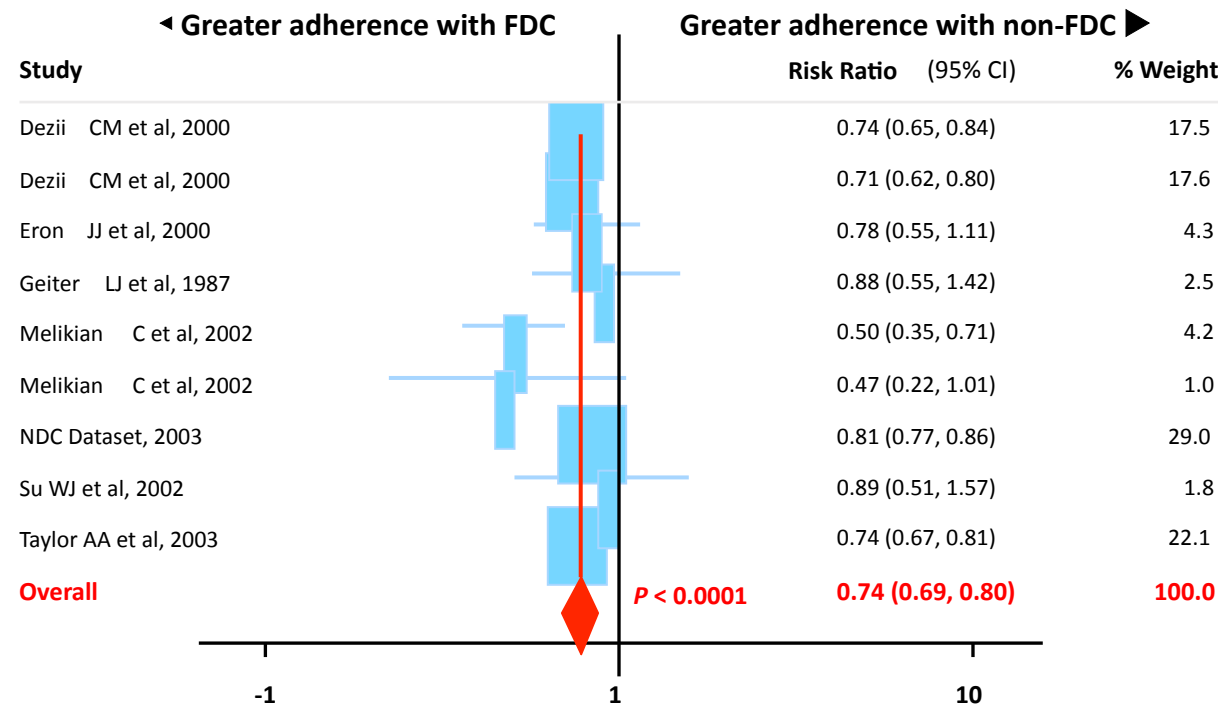
- 452 switches due to toxicity/perceived toxicity



# Adherence benefits of STR/ FDC

Meta-analysis of 9 clinical trials in 4 therapeutic areas (TB, HTN, HIV, DM)  
11,925 FDC patients and 8317 non-FDC patients

## Effect of FDCs versus non-FDC on risk of non-adherence

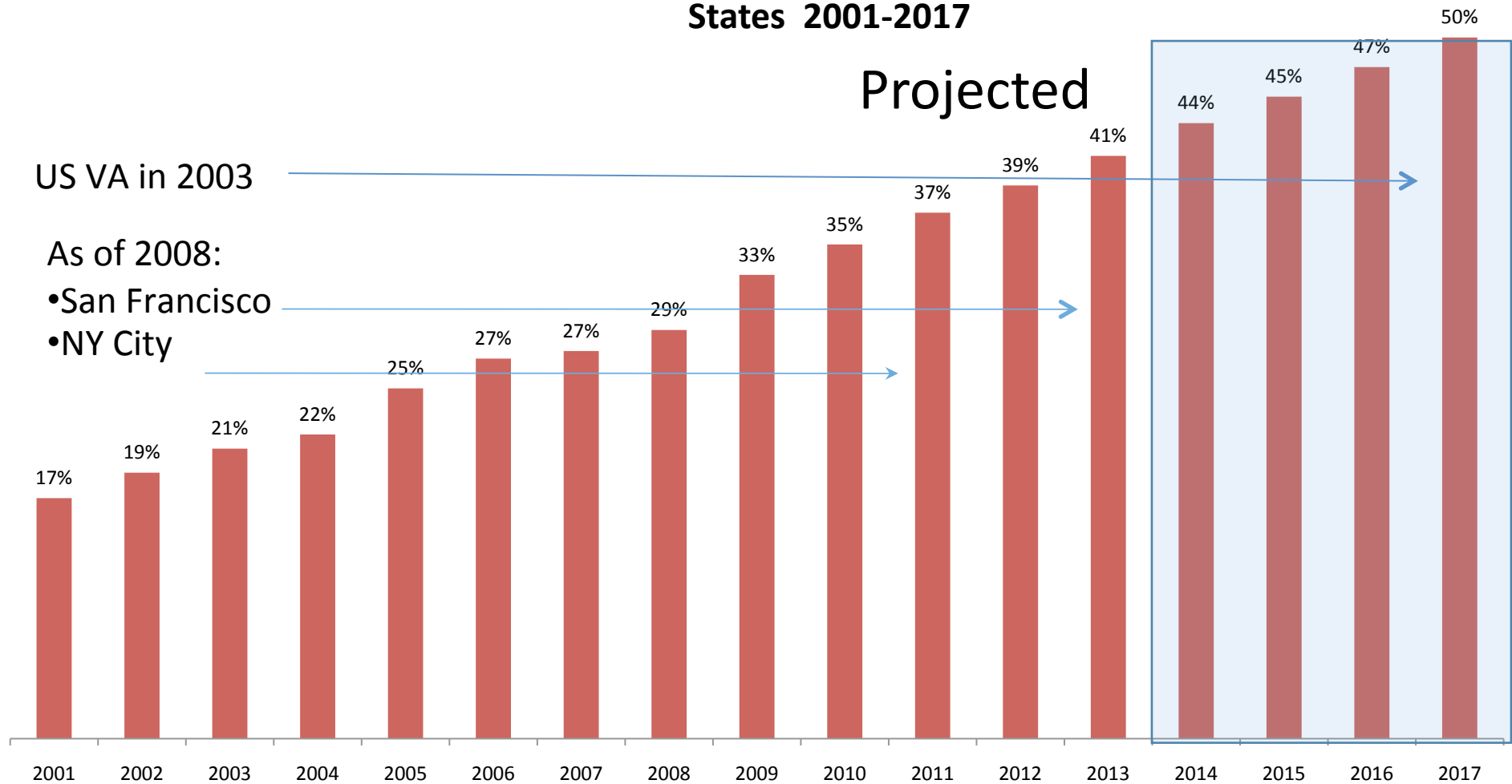


FDC regimens reduce risk of non-adherence by ~25% (compared to dosing with individual pills)



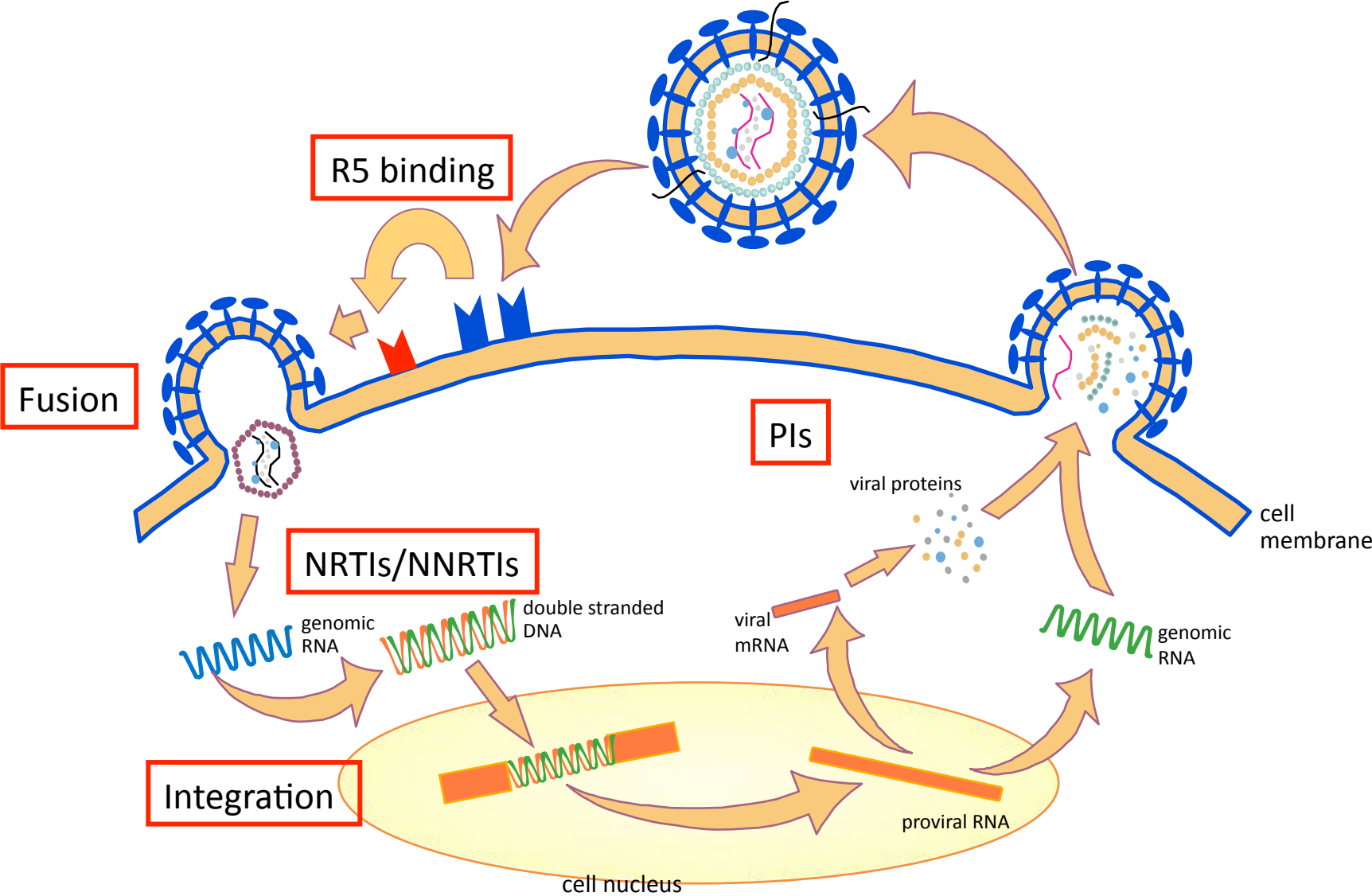
# Drugs associated with ageing

Projected Proportion of those 50+ Years of Age\* Living With HIV in United States 2001-2017



\*Data from 2008, onward projected based on 2001-2007 trends (calculated by author), 2001-2007 data from CDC Surveillance Reports 2007. New York and San Francisco data from their Departments of Public Health.

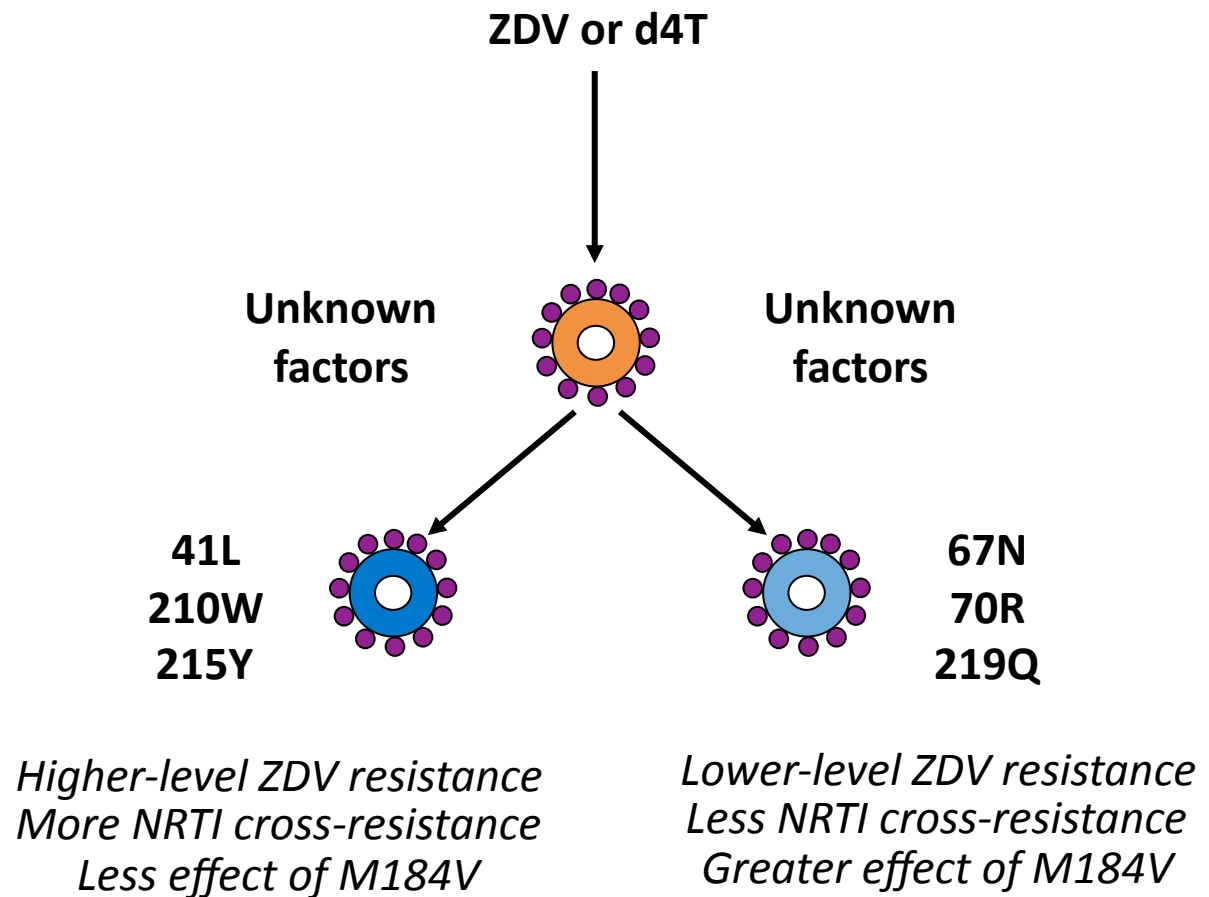
# HIV lifecycle and where ARVs act...




# Resistance: not always so simple – NRTIs. Dichotomous Pathways to Resistance

TAMs emerge  
sequentially with ZDV-  
and  
d4T-containing  
regimens after M184V

6 identified:  
**M41L, D67N, K70R,  
L210W, T215Y/F,  
K219Q/E/N/R**



# Resistance not always so simple – NNRTIs Weighting mutation system helps predict response

Relative weight* for individual ETR RAMs					Total weighted genotypic score	
1	1.5	2.5	3			
V90I	V106I	L100I	Y181I		0–2	Highest response
A98G	E138A	K101P	Y181V			
K101E	V179F	Y181C				
K101H	G190S	M230L			2.5–3.5	Intermediate response
V179D						
V179T						
G190A						
					≥4	Reduced response

Example: K101H + G190A = Weighted score of 2 = Highest response

\*When the genotype report shows a mixture of two or more different substitutions at the same position, only the highest of the individual weight factors for these substitutions is counted when calculating the weighted genotypic score.

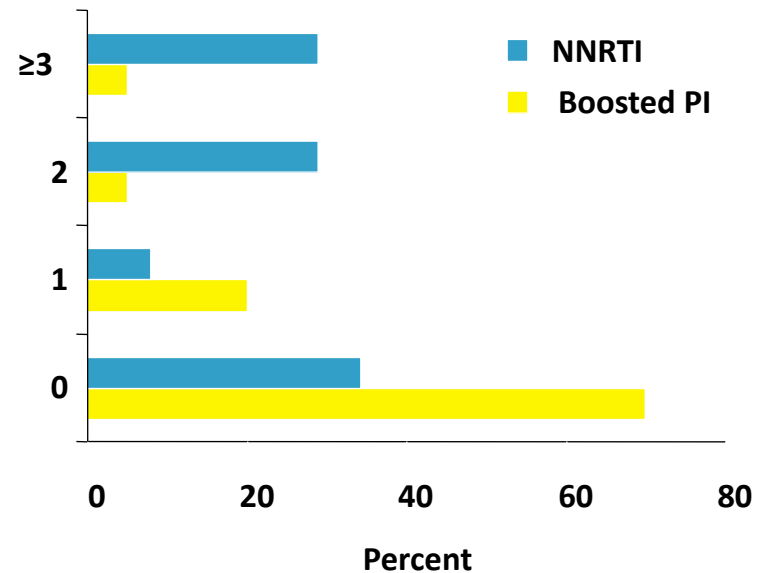
# Resistant to resistance: boosted PI's

	Quad (n=353)	ATV/r + FTC/TDF (n=355)
<b>Subjects Analyzed for Resistance<sup>a</sup>, n (%)</b>	<b>12 (3)</b>	<b>8 (2)</b>
<b>Subjects with Resistance to ARV Regimen, n (%)</b>	<b>5 (1)</b>	<b>0</b>
<b>Any Primary Integrase-R, n</b>	<b>4</b>	<b>-</b>
E92Q	1	-
T66I	1	-
Q148R	2	-
N155H	2	-
<b>Any Primary PI-R, n</b>	<b>-</b>	<b>0</b>
<b>Any Primary NRTI-R, n</b>	<b>4</b>	<b>0</b>
M184V/I	4	
K65R	1	

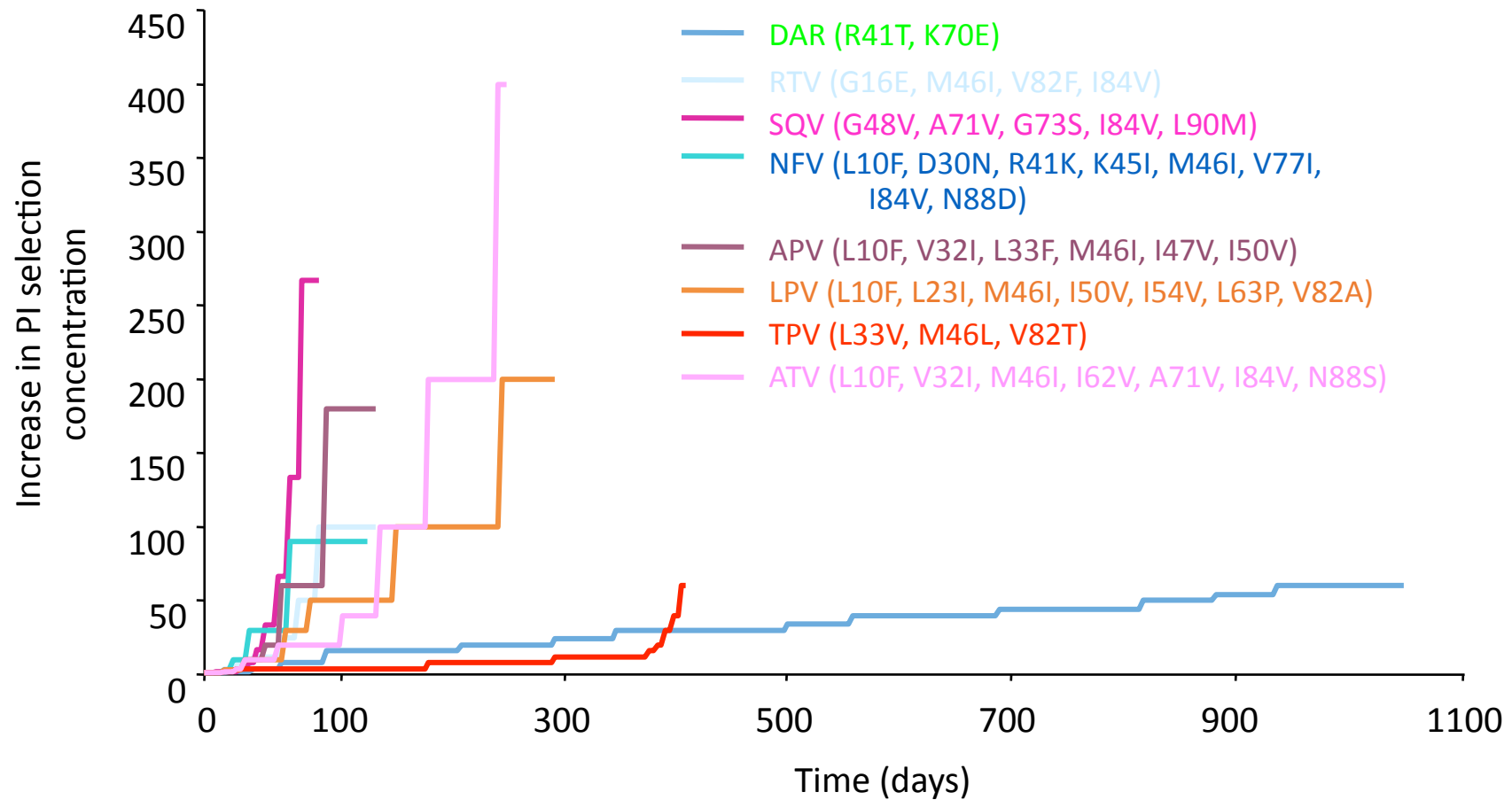
# Boosted PIs protect against emergence of drug resistance after 1<sup>st</sup> line ART

- Combination ARV therapy started Jan 1999 – Dec 2005 (n=1323)
  - Boosted PI (n=518)
  - NNRTI (n=805)
- Viral failure (defined as HIV RNA >500 c/mL after more than 180 days of treatment) by third agent
  - Boosted PI (n=4.6%); NNRTI (n=5.6%)
  - No difference by regimen but more resistance emerged with NNRTI-based regimens

Mutations at time of virologic failure\*



# But also differences...



De Meyer S, et al. *Antimicrob Agents Chemother.* 2005;49: 2314-21.

De Meyer S, et al. *XV IHDRW*, 2006, Poster 19.

# Resistance not always so simple – PIs

DRV Weighting mutation system helps predict response

Estimated increase in FC	<2	2 to 3	3 to 4	>4
Mutations	V11I <u>I54L</u> G73S L89V	<u>V32I</u> L33F I47V	I54M <u>L76V</u> <u>I84V</u>	<u>I50V</u>

Add mutations up for fold-change

Example: = ~ 5–7 fold-change = **Intermediate activity**



# And for tipranavir....

- 1	0.5	1	2
L24I	L10V/I	V11L	I47V
D30N	I13V	V32I	I54A
I50L/V	K20R	A71L	V82T
154L	M46L	G73T	I84V
L76V	L90M	L89V	
V82I			

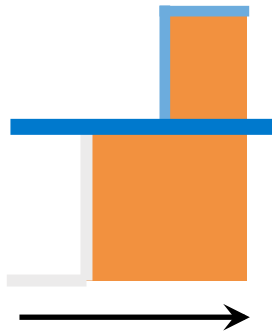
Add mutations up for fold-change

Example: = ~ 4 fold-change = **Intermediate activity**

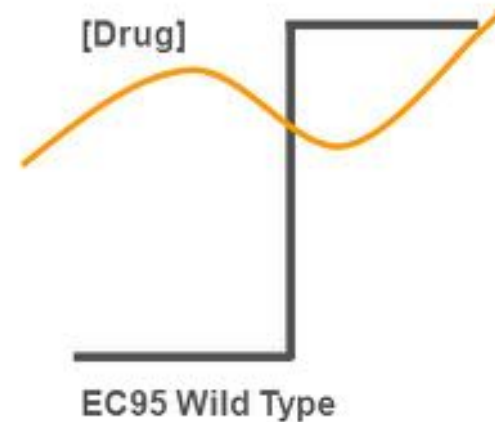
# Single mutation leading to resistance – all or nothing

## First-generation NNRTI

One mutation correlates with reduced virological response



Increasing number of mutations at baseline



1. Antinori A, et al. AIDS Res Hum Retroviruses. 2002;18:835–8.
2. Lecossier D, et al. J Acquir Immune Defic Syndr. 2005;38:37–42.
3. Vingerhoets J, et al. 17th IDHRW 2008 [Poster 32].
4. De Béthune MP, et al. 4th EHDRW 2006 [Poster 51].
5. de Mendoza C, et al. HIV Clin Trials. 2006;7:163–71.

# Thank you

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