

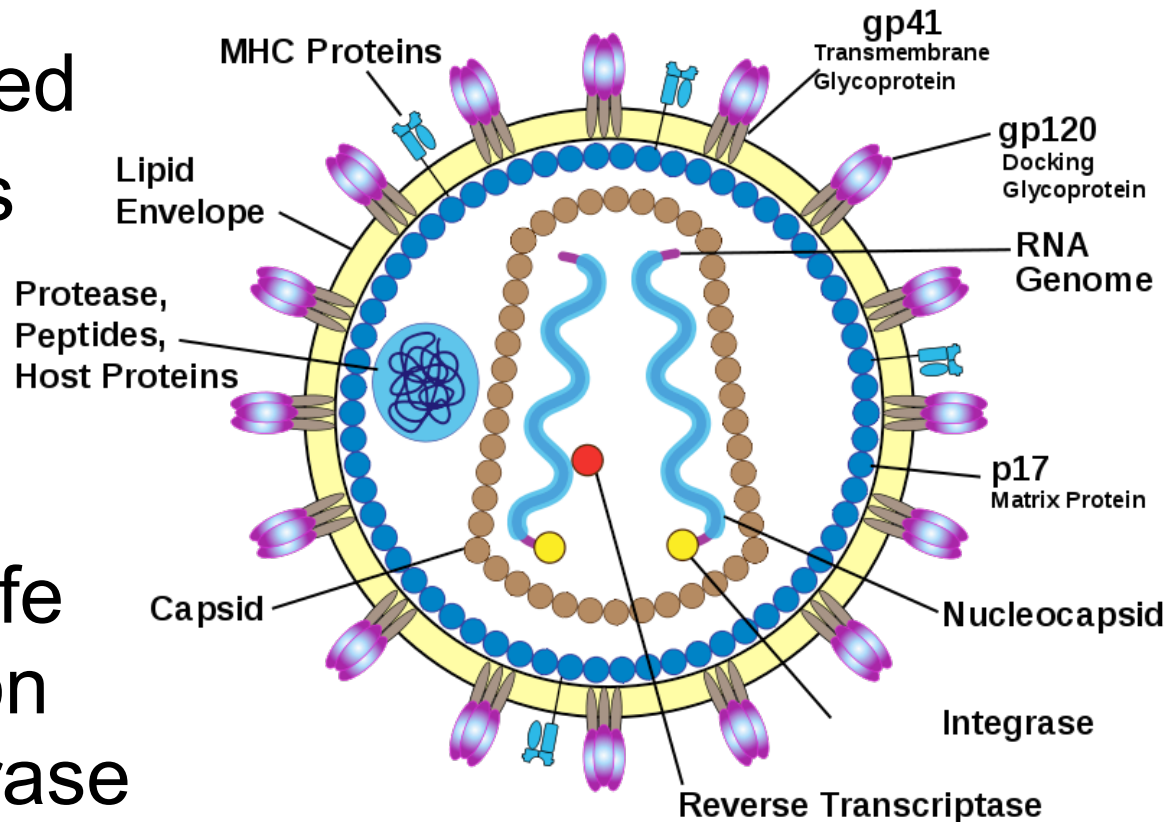
# Outline

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- Biology of HIV and drug resistance
- Mathematical model of induction therapy
- Impulsive differential equations
- Determining resistance thresholds
- Calculating the length and number of drug holidays
- Comparison with clinical results
- Implications.

# HIV/AIDS

- 33 million infected
- Infection causes a depletion of CD4<sup>+</sup> T cells
- No cure
- Drugs prolong life (RTIs, PIs, fusion inhibitors, integrase inhibitors).

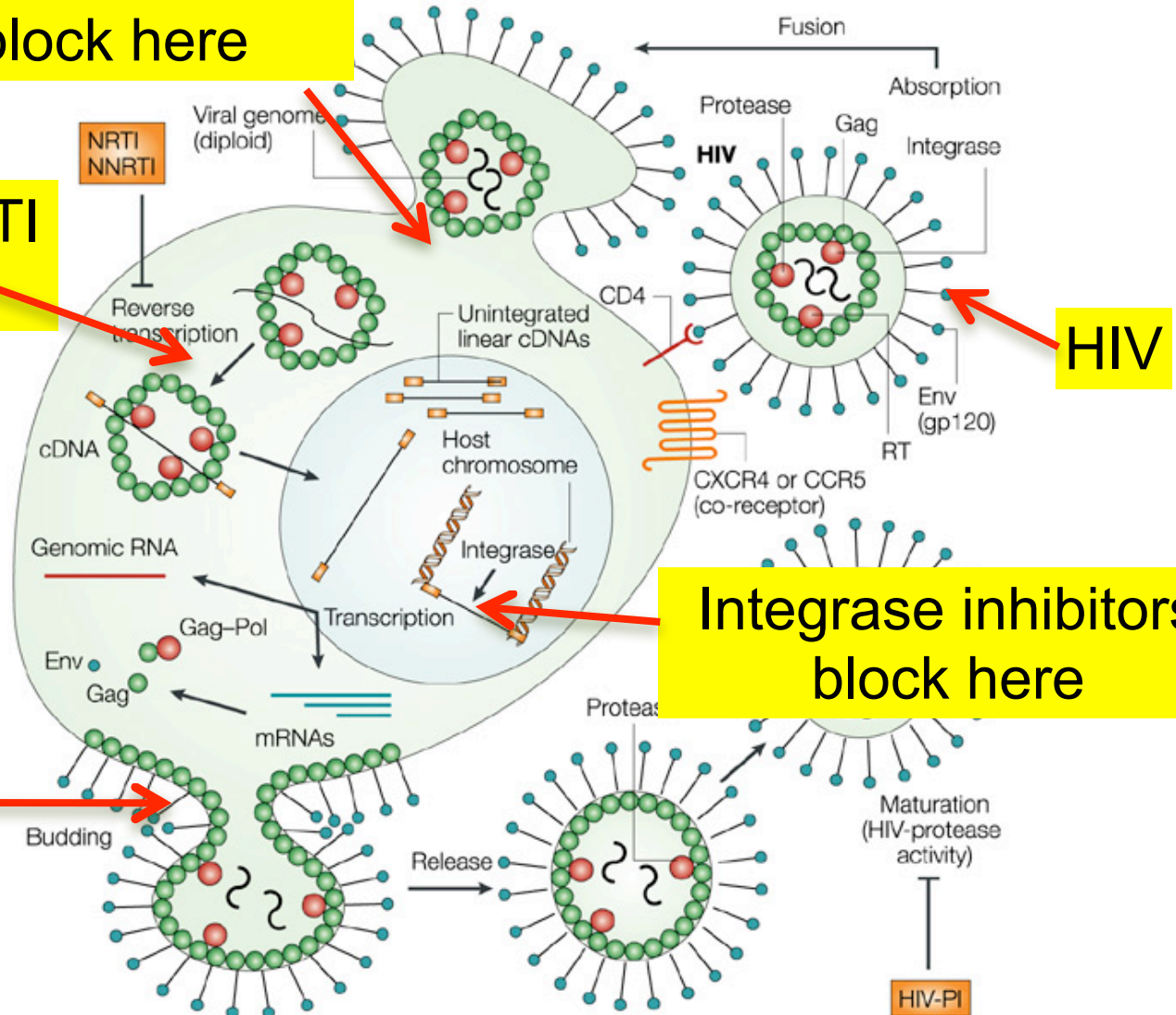


*RTI=reverse transcriptase inhibitors*  
*PIs=Protease inhibitors*

Fusion inhibitors  
block here

NRTI and NNRTI  
block here

PIs block here.



HIV

Integrase inhibitors  
block here

*NRTIs=nucleoside reverse transcriptase inhibitors*  
*NNRTIs=nonnucleoside reversetranscriptase inhibitors*  
*PIs=protease inhibitors*

# Adherence

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- Studies have shown that 40-60% of patients are less than 90% adherent to their drugs
- Adherence also decreases over time
- Lack of adherence promotes the development of drug-resistant mutations.



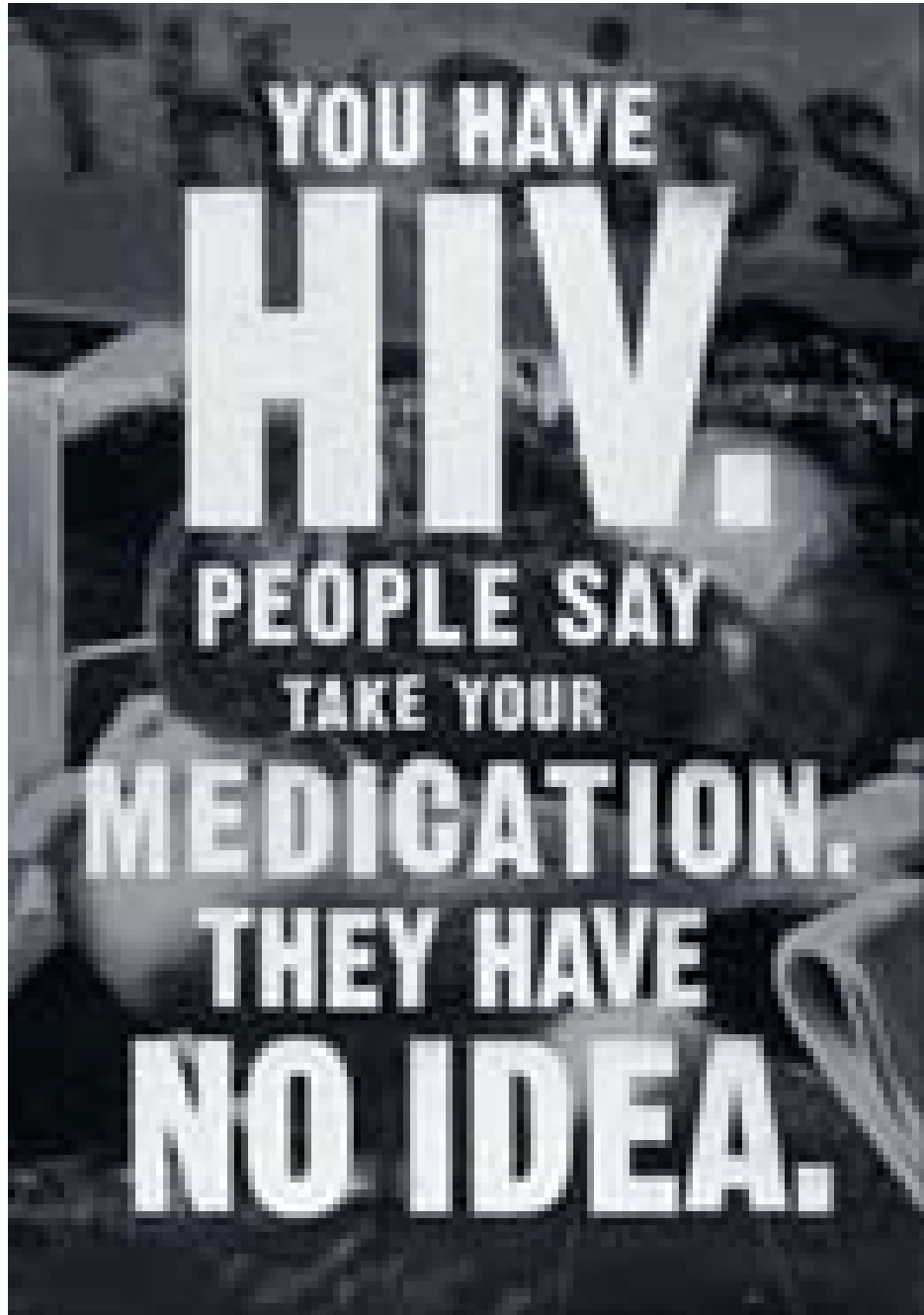
# Drug holidays

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"Drug holidays" are extended breaks from the drugs. They may occur due to:

- lifestyle factors
- relief from side effects
- economic implications (especially in the developing world).





YOU HAVE

**HIV.**

PEOPLE SAY

TAKE YOUR

**MEDICATION.**

THEY HAVE

**NO IDEA.**

# Induction Therapy

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- New HIV/AIDS treatment regime that hopes to benefit patients by decreasing drug resistance and reducing the overall number of drugs that must be taken
- Two phases:
  - Induction phase: Period of intensified antiretroviral therapy
  - Maintenance phase: Long-term regimen
- The induction phase lasts about six months.



*RTI=reverse transcriptase inhibitors*  
*PIs=Protease inhibitors*

# Adherence to Induction Therapy

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- Since induction therapy has a higher chance of pill fatigue than maintenance therapy, it is likely that patients will take some holidays during this period
- If a holiday occurs at the end, induction therapy finished too early, which isn't acceptable
- However, a few breaks in the middle may be acceptable...  
...assuming subsequent therapy was undertaken to control the virus.



# Research Questions

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**For all PI-sparing drugs, how long can a drug holiday be?**

**How many doses must be taken after a holiday to return drug levels to before?**

**How many drug holidays can be taken during the entire induction phase?**

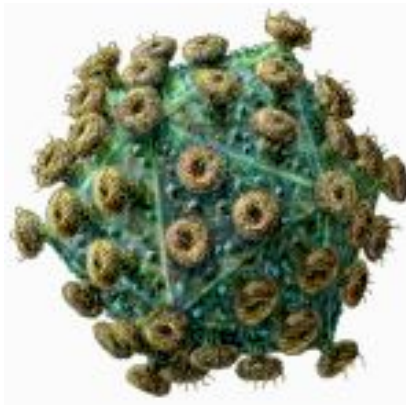
*PI sparing drugs=reverse transcriptase inhibitors, fusion inhibitors, integrase inhibitors*

# Modelling Drug Therapy

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We assume two strains of the virus:

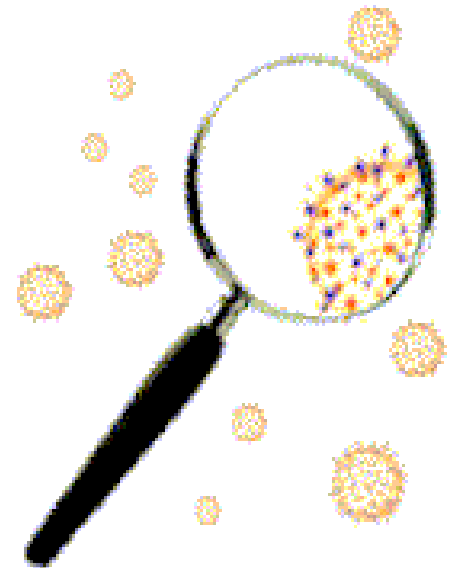
1. The wild-type strain will dominate in the absence of drugs
2. There is also a mutant strain that is a less efficient competitor, but more resistant to the drugs.



# The basic idea

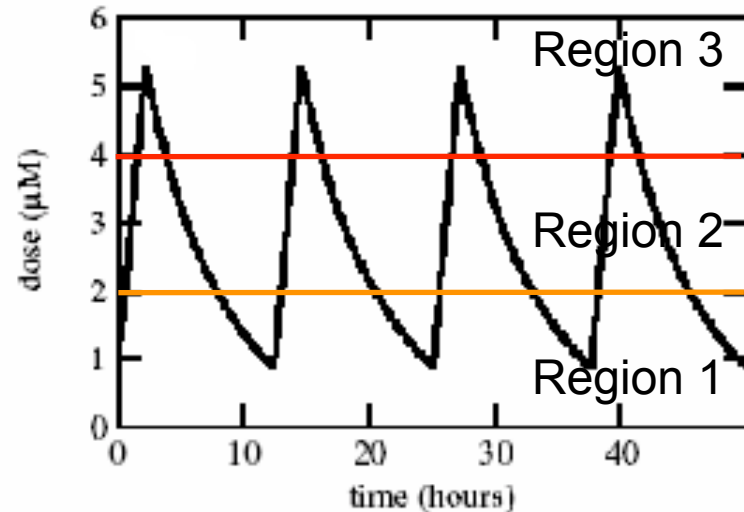
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- Resistant mutants are not impervious to the drugs
- Rather, “resistance” confers a degree of resistance to the drugs
- Thus, if drug concentrations in the cell were sufficiently high, then even the mutant would be controlled.



# Drug dependence

- As drug levels fall, the wild-type strain is controlled, but the mutant may take hold
- When the drug falls to trough levels, the wild-type strain can regain its advantage
- The amount of drug will determine how and when one, the other or neither strain gains dominance.



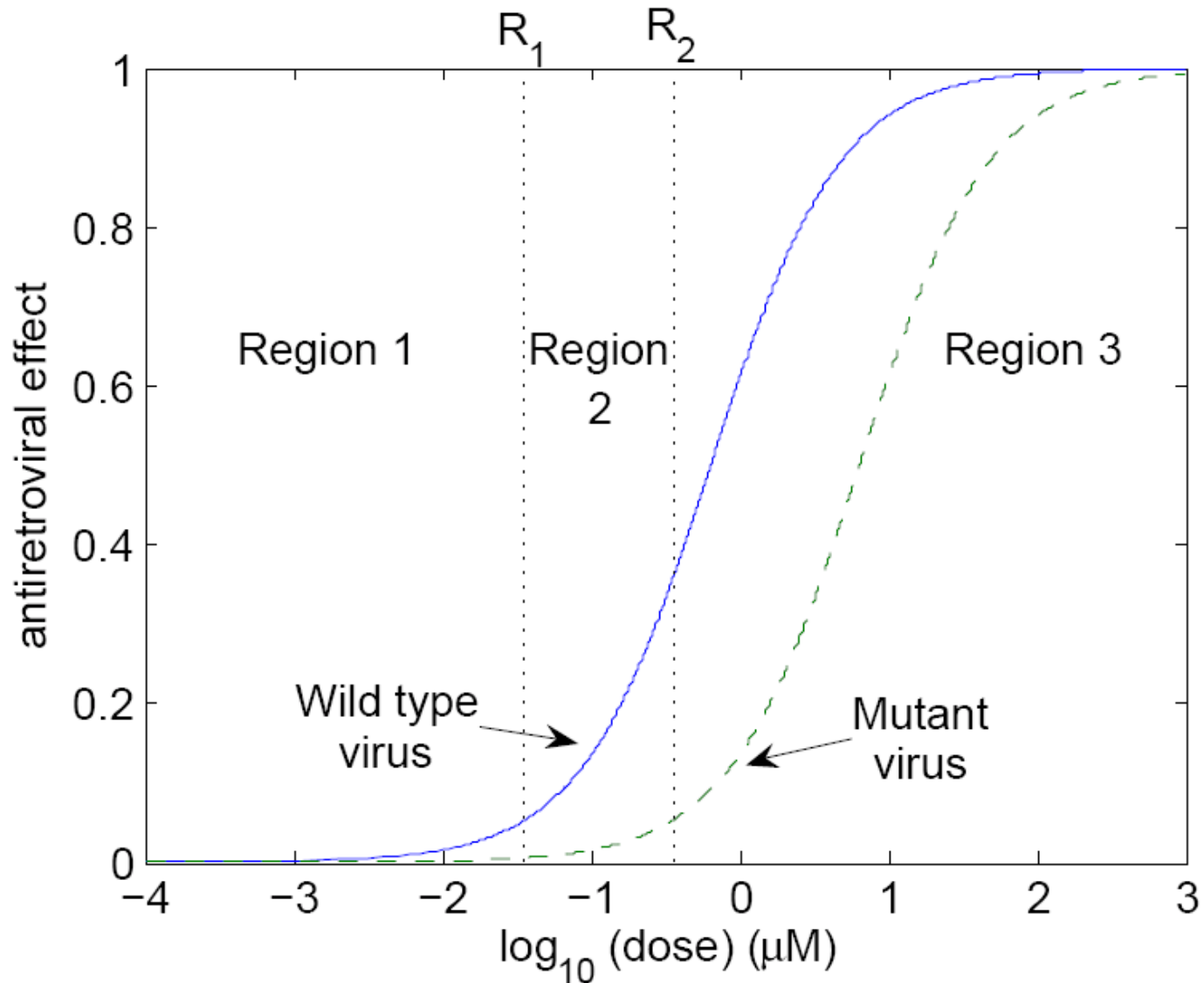
# The three regions

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We consider three drug regimes:

- Region 1 (low): drugs are not sufficient to inhibit either strain
- Region 2 (medium): drugs will inhibit the wild type, but not the mutant
- Region 3 (high): drugs will inhibit both strains.

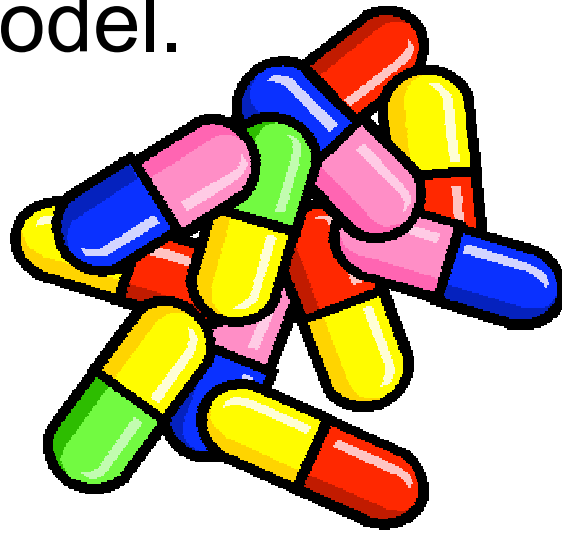
# The region thresholds



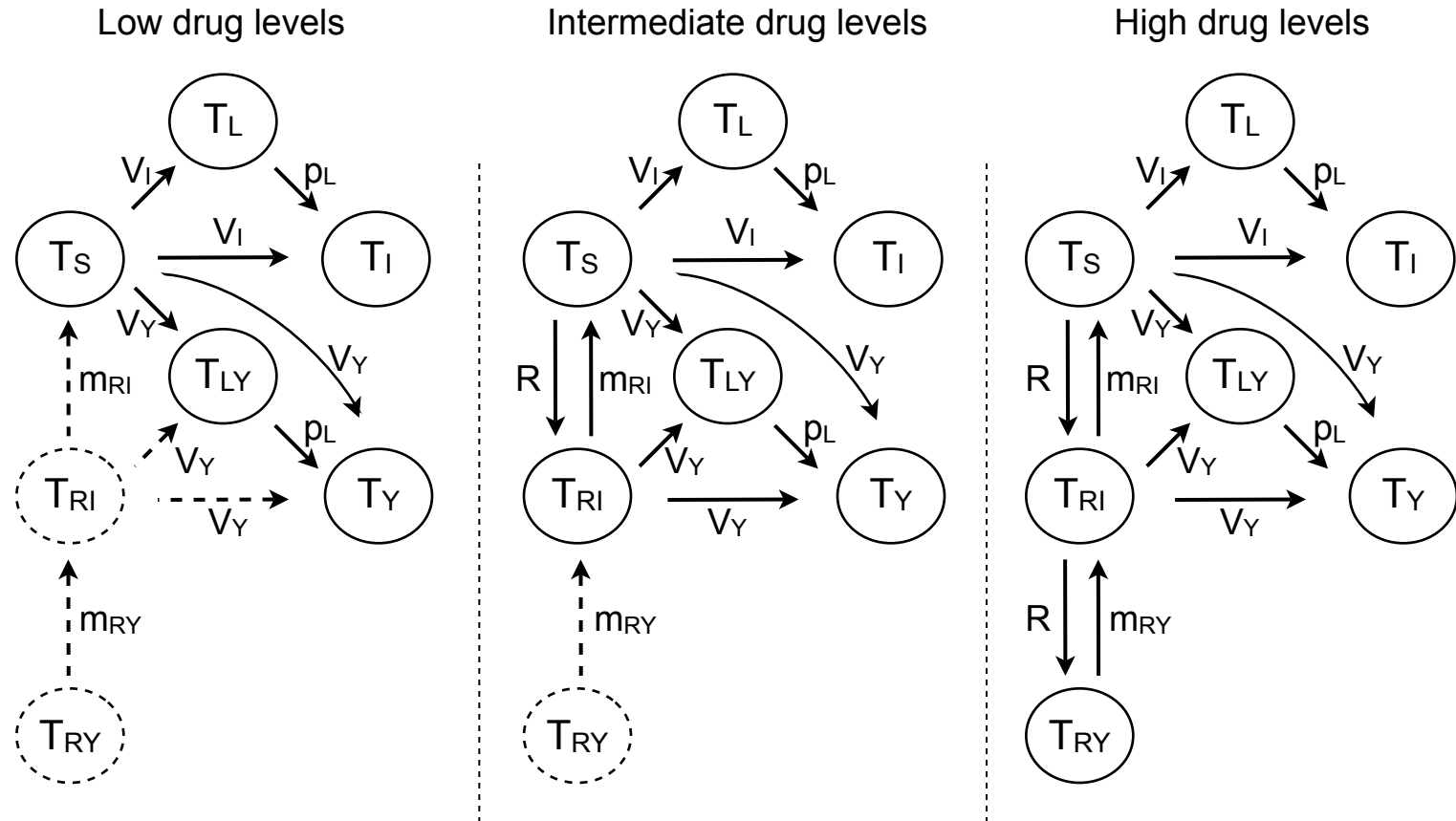
# The model itself changes

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- Within each regime, the model itself will be different
- We have three models, connected by the drug behaviour
- Thus, as the drug levels change, so too does each model.



# The Model - Flow Chart



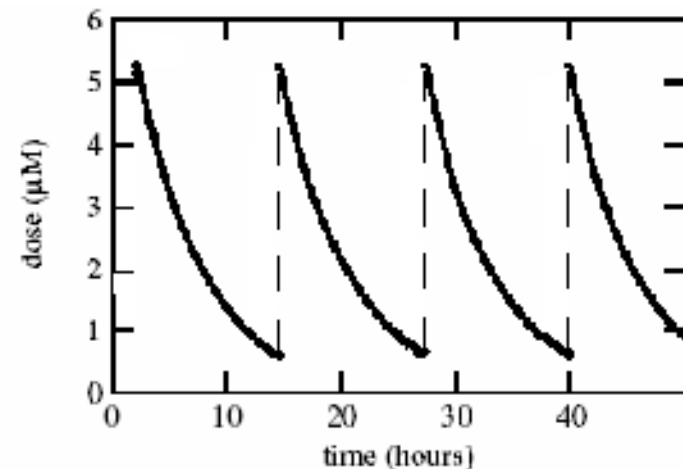
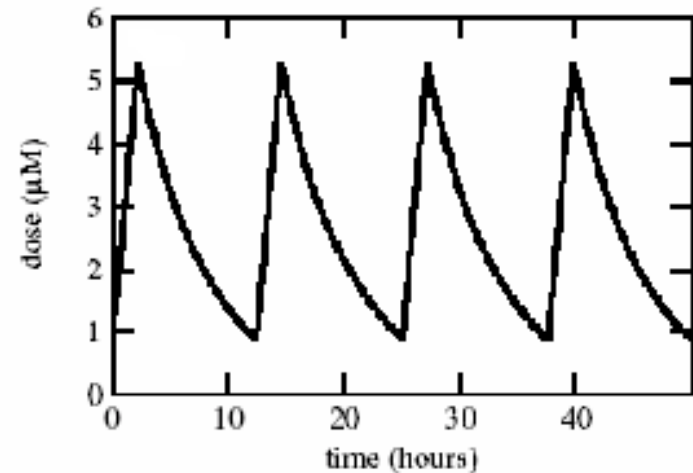
$V_I$ =wild-type virus       $V_Y$ =mutant virus       $V_{NI}$ =noninfectious       $T_S$ =Susceptible T cells  
 $T_I$ =Infected (wild type)       $T_Y$ =Infected (mutant)       $R$ =drug       $T_L$ =Latently infected (wild type)  
 $T_{RY}$ =Highly inhibited       $T_{RI}$ =Intermediate inhibited       $m_j$ =waning rates       $T_{LY}$ =Latently infected (mutant)



# Impulsive Differential Equations

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- Assume drug effects are instantaneous
- That is, the time-to-peak is assumed to be negligible
- Impulsive differential equations provide an adequate mathematical model of evolutionary processes that suddenly change their state.



# Putting it together

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- The model thus consists of a system of ODEs (virus and T cells) together with an ODE and a difference equation (drugs).



# The model in Region 1 (low drugs)

$$\begin{aligned} \frac{dV_I}{dt} &= n_I \omega T_I - d_V V_I - r_I T_S V_I \\ \frac{dV_Y}{dt} &= n_I \omega T_Y - d_V V_Y - r_Y T_S V_Y - r_Y T_{RI} V_Y \\ \frac{dV_{NI}}{dt} &= n_I (1 - \omega) (T_I + T_Y) - d_V V_{NI} \\ \frac{dT_S}{dt} &= \lambda - r_I T_S V_I - r_Y T_S V_Y - d_S T_S + m_{RI} T_{RI} \\ \frac{dT_I}{dt} &= (1 - \psi) r_I T_S V_I - d_I T_I + p_L T_{LI} \\ \frac{dT_{LI}}{dt} &= \psi r_I T_S V_I - d_S T_{LI} - p_L T_{LI} \\ \frac{dT_Y}{dt} &= (1 - \psi) r_Y T_S V_Y - d_I T_Y + (1 - \psi) r_Y T_{RI} V_Y + p_L T_{LY} \\ \frac{dT_{LY}}{dt} &= \psi r_Y T_S V_Y - d_S T_{LY} - p_L T_{LY} + \psi r_Y T_{RI} V_Y \\ \frac{dT_{RI}}{dt} &= -r_Y T_{RI} V_Y - (d_S + m_{RI}) T_{RI} + m_{RY} T_{RY} \\ \frac{dT_{RY}}{dt} &= -(d_S + m_{RY}) T_{RY} \end{aligned}$$

$V_I$ =wild-type virus  
 $V_Y$ =mutant virus  
 $V_{NI}$ =noninfectious  
 $T_S$ =Susceptible T cells  
 $T_I$ =Infected (wild type)  
 $T_Y$ =Infected (mutant)  
 $T_{LI}, T_{LY}$ =Latently infected  
 $T_{RI}$ =Intermediately inhibited  
 $T_{RY}$ =Highly inhibited  
 $R$ =drug  
 $\lambda$ =lymphic source  
 $d_j$ =clearance rates  
 $m_j$ =waning rates  
 $n_i \omega$ =# infectious virions  
 $r_i$ =wild type infection rate  
 $r_Y$ =mutant infection rate  
 $\psi$ =latently infected proportion

# The model in Region 2 (intermediate drugs)

$$\begin{aligned} \frac{dV_I}{dt} &= n_I \omega T_I - d_V V_I - r_I T_S V_I \\ \frac{dV_Y}{dt} &= n_I \omega T_Y - d_V V_Y - r_Y T_S V_Y - r_Y T_{RI} V_Y \\ \frac{dV_{NI}}{dt} &= n_I (1 - \omega) (T_I + T_Y) - d_V V_{NI} \\ \frac{dT_S}{dt} &= \lambda - r_I T_S V_I - r_Y T_S V_Y - d_S T_S + m_{RI} T_{RI} - r_P T_S R \\ \frac{dT_I}{dt} &= (1 - \psi) r_I T_S V_I - d_I T_I + p_L T_{LI} \\ \frac{dT_{LI}}{dt} &= \psi r_I T_S V_I - d_S T_{LI} - p_L T_{LI} \\ \frac{dT_Y}{dt} &= (1 - \psi) r_Y T_S V_Y - d_I T_Y + (1 - \psi) r_Y T_{RI} V_Y + p_L T_{LY} \\ \frac{dT_{LY}}{dt} &= \psi r_Y T_S V_Y - d_S T_{LY} - p_L T_{LY} + \psi r_Y T_{RI} V_Y \\ \frac{dT_{RI}}{dt} &= -r_Y T_{RI} V_Y - (d_S + m_{RI}) T_{RI} + m_{RY} T_{RY} + r_P T_S R \\ \frac{dT_{RY}}{dt} &= -(d_S + m_{RY}) T_{RY} \end{aligned}$$

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 $m_j$ =waning rates  
 $n_i \omega$ =# infectious virions  
 $r_i$ =wild type infection rate  
 $r_Y$ =mutant infection rate  
 $\psi$ =latently infected proportion  
 $r_P$ =drug uptake rate

# The model in Region 3 (high drugs)

$$\begin{aligned} \frac{dV_I}{dt} &= n_I \omega T_I - d_V V_I - r_I T_S V_I \\ \frac{dV_Y}{dt} &= n_I \omega T_Y - d_V V_Y - r_Y T_S V_Y - r_Y T_{RI} V_Y \\ \frac{dV_{NI}}{dt} &= n_I (1 - \omega) (T_I + T_Y) - d_V V_{NI} \\ \frac{dT_S}{dt} &= \lambda - r_I T_S V_I - r_Y T_S V_Y - d_S T_S + m_{RI} T_{RI} - r_R T_S R \\ \frac{dT_I}{dt} &= (1 - \psi) r_I T_S V_I - d_I T_I + p_L T_{LI} \\ \frac{dT_{LI}}{dt} &= \psi r_I T_S V_I - d_S T_{LI} - p_L T_{LI} \\ \frac{dT_Y}{dt} &= (1 - \psi) r_Y T_S V_Y - d_I T_Y + (1 - \psi) r_Y T_{RI} V_Y + p_L T_{LY} \\ \frac{dT_{LY}}{dt} &= \psi r_Y T_S V_Y - d_S T_{LY} - p_L T_{LY} + \psi r_Y T_{RI} V_Y \\ \frac{dT_{RI}}{dt} &= -r_Y T_{RI} V_Y - (d_S + m_{RI}) T_{RI} + m_{RY} T_{RY} + r_R T_S R - r_Q T_{RI} R \\ \frac{dT_{RY}}{dt} &= -(d_S + m_{RY}) T_{RY} + r_Q T_{RI} R \end{aligned}$$

$V_I$ =wild-type virus  
 $V_Y$ =mutant virus  
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 $m_j$ =waning rates  
 $n_i \omega$ =# infectious virions  
 $r_i$ =wild type infection rate  
 $r_Y$ =mutant infection rate  
 $\psi$ =latently infected proportion  
 $r_R, r_Q$ =drug uptake rates

...with the (impulsive) dynamics of the drugs:

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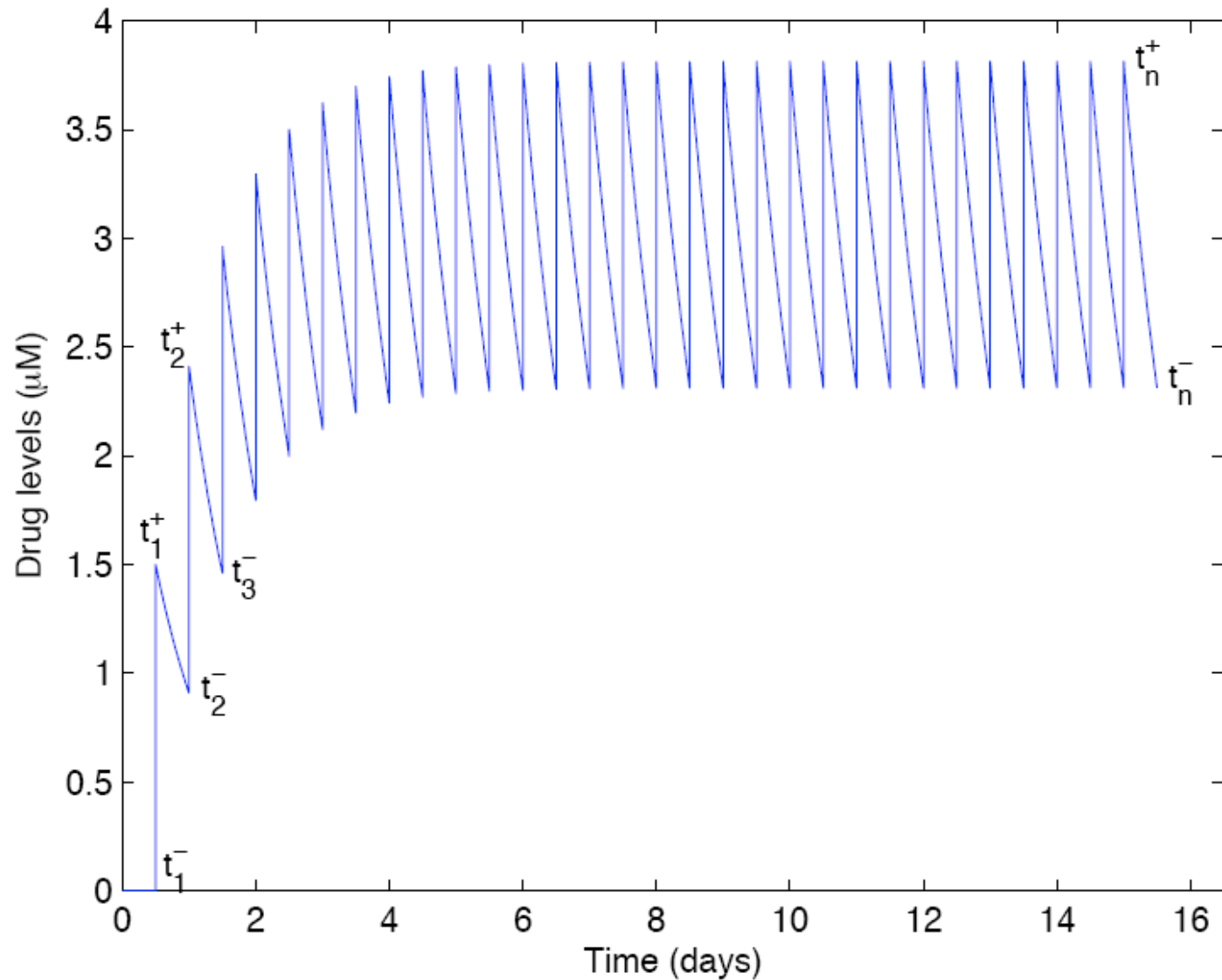
$$\frac{dR}{dt} = -d_R R \quad t \neq t_k$$

$$\Delta R = \begin{cases} R^i & \text{if a dose is to be taken} \\ 0 & \text{if no dose is to be taken.} \end{cases}$$



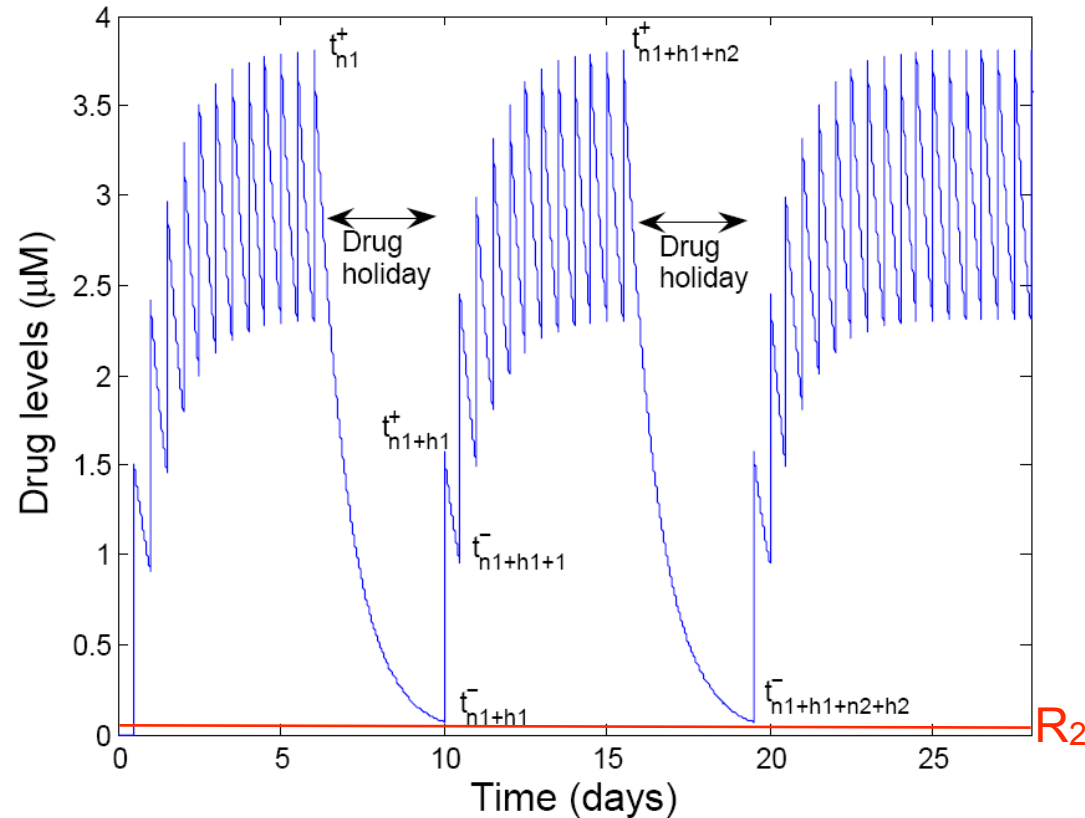
*R*=drug  
*d<sub>R</sub>*=decay rate  
*R<sup>i</sup>*=dosage  
*t<sub>k</sub>*=impulse time

# Perfect adherence



# Imperfect Adherence to Drug Therapy

- As long as the drug concentration level does not drop below  $R_2$ , there is sufficient amount of drug to control both viral strains
- Drug holidays can be taken, and the number of doses that can be missed in order to stay over  $R_2$  can be calculated.





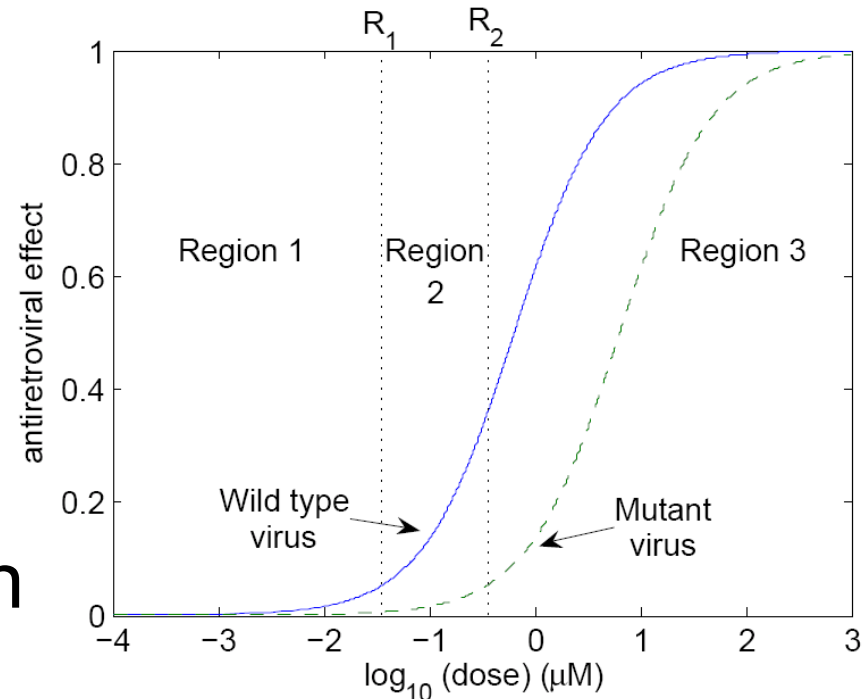
# How to determine $R_1$ ?

- Define  $R_1$  by

$$\frac{R_1}{R_1 + IC_{50}} = \frac{R_2}{R_2 + nIC_{50}}$$

where  $n$  is the degree of  $n$ -fold resistance conferred by the mutation

- eg for 10-fold resistance, you need 10 times as much drug to have the same effect
- Thus  $R_1 = R_2/n$ .



$R_1 = \text{Region 1 threshold}$   
 $R_2 = \text{Region 2 threshold}$

# How to determine $R_2$ ?

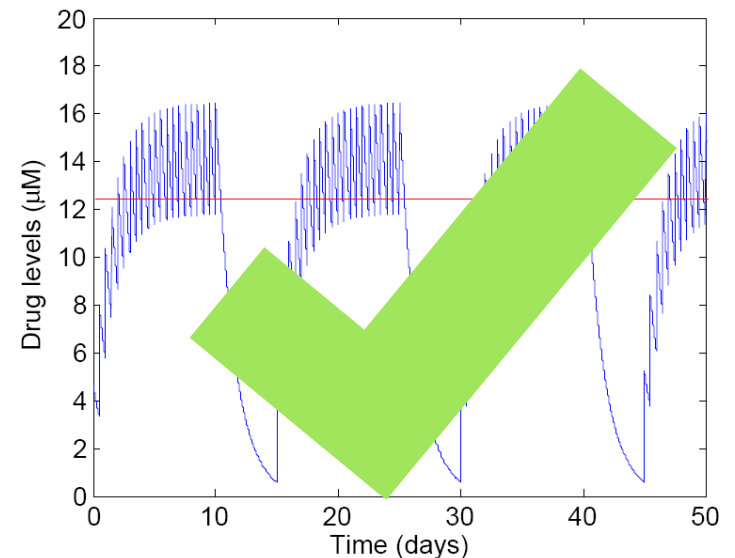
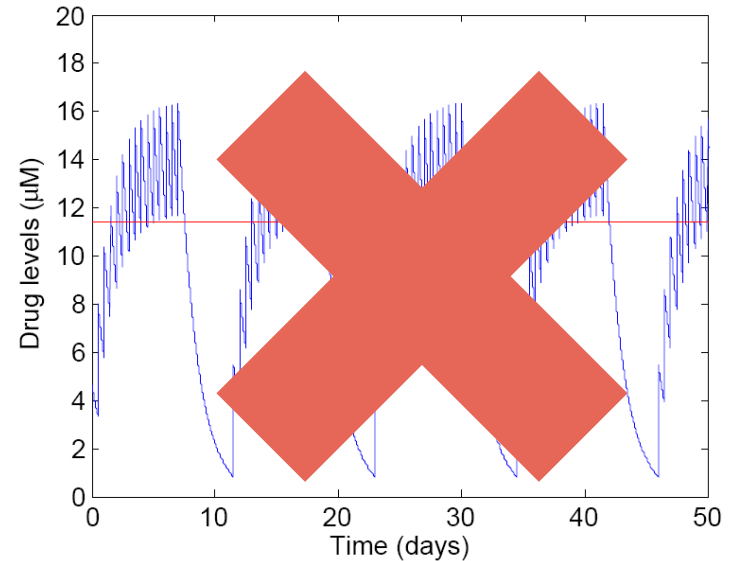
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- $V_Y(0) > 0$  and  $V_Y'(0) < 0$ , so the resistance viral load is initially decreasing
- The resistance viral load either reaches a minimum at time  $t^*$  or it decreases indefinitely (in which case we can define  $t^*$  arbitrarily)
- Define  $R_2 = R(t^*)$
- Thus  $R_2$  is reached when resistance stops decreasing and starts increasing
- This guarantees that resistance has not emerged at time  $t^*$ .

$V_Y$  = mutant virus  
 $R_2$  = Region 2 threshold

# An additional condition

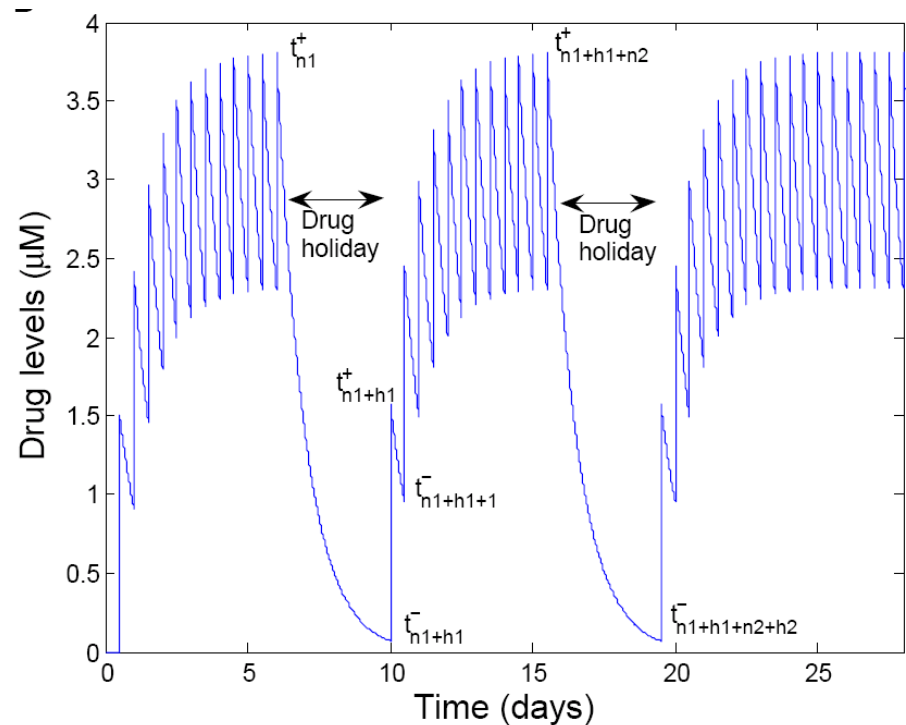
- The mean drug concentration must be larger than the trough value when no drug holiday is taken
- This ensures that drugs are maintained at sufficiently high levels over the length of the entire induction phase.



# Imperfect adherence

We need to determine:

- $n_1$ , the number of doses to be taken initially
- $h_1$ , the number of doses that can be safely missed during the drug holiday
- $n_2$ , the number of doses needed to return to high drug levels
- $k$ , the number of doses that must be taken at the end so that the induction phase ends at high levels.



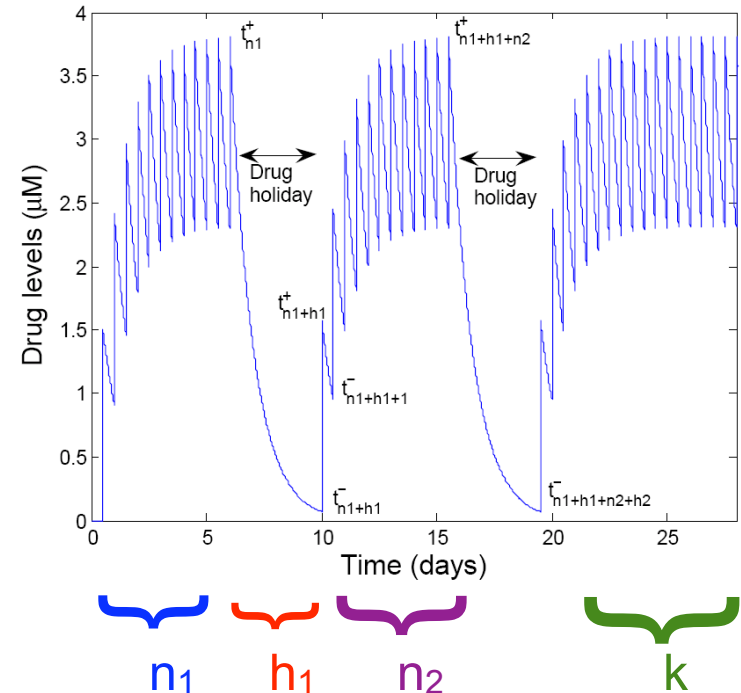
# Taking the first $n_1$ doses

- To be within  $\varepsilon_1$  of the periodic orbit, we require

$$n_1 > \frac{1}{d_r \tau} \ln \left[ \frac{R^i}{\varepsilon_1 (1 - e^{-d_r \tau})} \right] - 1$$

where  $\varepsilon_1$  is chosen to ensure the mean drug concentration is sufficiently high over the entire induction period.

$d_r$ =drug waning rate  
 $\tau$ =period  $R^i$ =dosage



# Missing $h_1$ doses

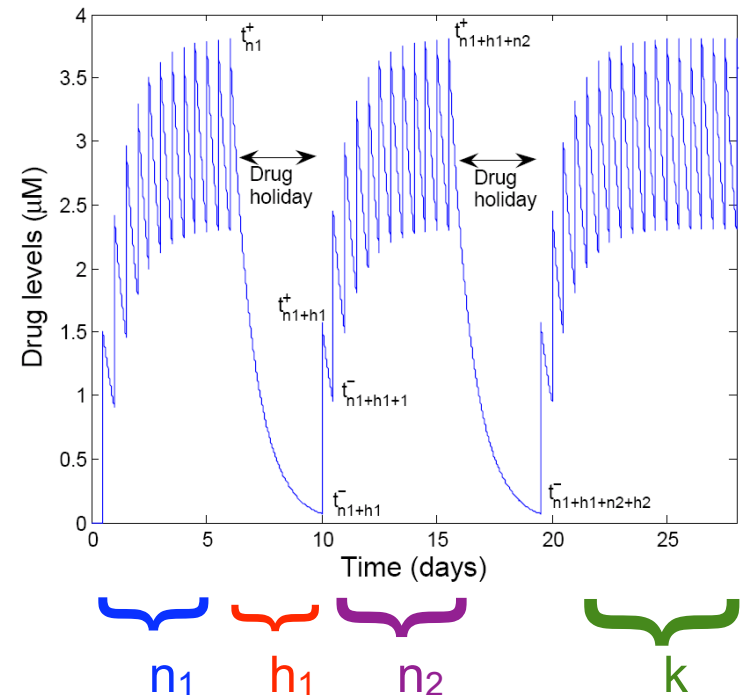
- To avoid Region 2 after  $h_1$  doses have been missed, we require

$$h_1 < \frac{1}{d_r \tau} \ln \left[ \frac{R^i}{R_2} \left( \frac{1 - e^{-(n_1+1)d_r \tau}}{1 - e^{-d_r \tau}} \right) \right]$$

- The number of doses that must be taken following a drug holiday satisfies

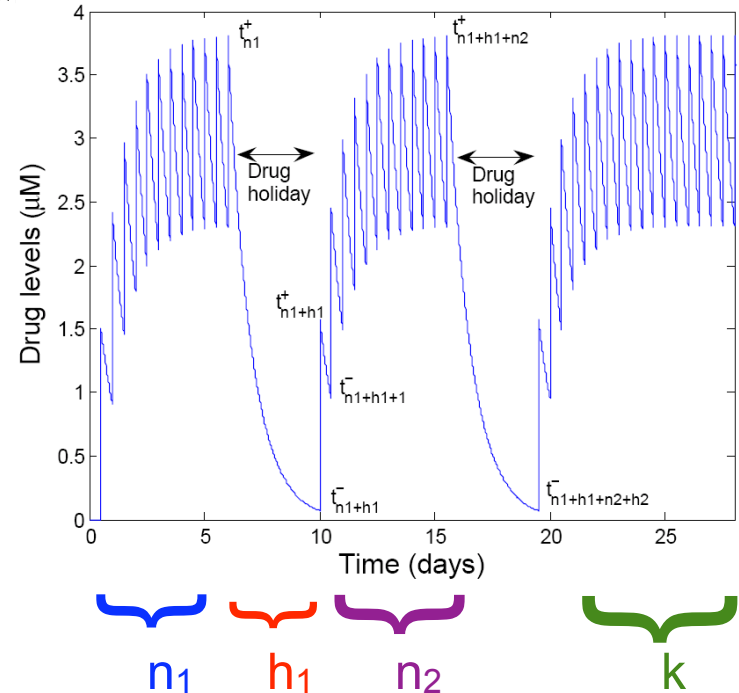
$$n_2 > \frac{1}{d_r \tau} \ln \left[ \frac{R^i e^{-d_r \tau} - R_2 (1 - e^{-d_r \tau})}{\epsilon_2 (1 - e^{-d_r \tau})} \right].$$

$d_r$ =drug waning rate  $\tau$ =period  
 $R_2$ =Region 2 threshold  $R^i$ =dosage



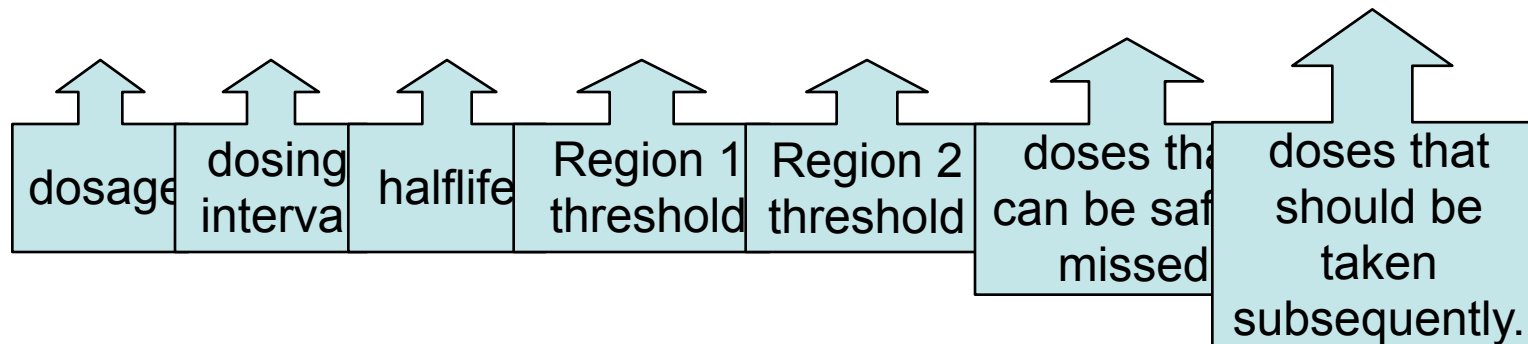
# Straightforward formulas

- If the tolerance is the same, then  $h_1=h_2=\dots$   
(ie all drug holidays have the same length)
- Furthermore,  $n_2=n_3=\dots=k$   
(ie all dosing intervals after a drug holiday are equal, including the final one)
- The formulas for  $h_i$  and  $n_j$  are straightforward and can be understood by policymakers.



# Results

Drug (units)	$R^i$ ( $\mu M$ )	$\tau$ (days)	$T_{1/2}$ (hours)	$R_1$ ( $\mu M$ )	$R_2$ ( $\mu M$ )	missable days (max)	subsequent days (min)
Abacavir (ABC)	12	1/2	15	$10^{-1.0269}$	$10^{-0.0269}$	3	7
Didanosine (ddI)	4.65	1/2	25	$10^{-1.2218}$	$10^{-0.2218}$	5	7.5
Emtricitabine (FTC)	7.2	1	39	$10^{-0.9788}$	$10^{0.0212}$	6	17
Lamivudine (3TC)	6	1/2	20	$10^{-1.1249}$	$10^{-0.1249}$	3.5	8.5
Stavudine (d4T)	2.144	1/2	7.5	$10^{-1.6383}$	$10^{-0.6383}$	1	2.5
Tenofovir (TDF)	1.184	1	60	$10^{-1.5229}$	$10^{-0.5229}$	10	24
Zidovudine (ZDV)	4.24	1/3	7	$10^{-1.6021}$	$10^{-0.6021}$	1.33	2.67
Delavirdine (DLV)	26.6	1/3	5.8	$10^{-1.4559}$	$10^{-0.4559}$	1.67	2.67
Efavirenz (EFV)	12.9	1	45	$10^{-0.8356}$	$10^{0.1644}$	9	22
Nevirapine (NVP)	7.5	1/2	27	$10^{-1.0088}$	$10^{-0.0088}$	5	12.5



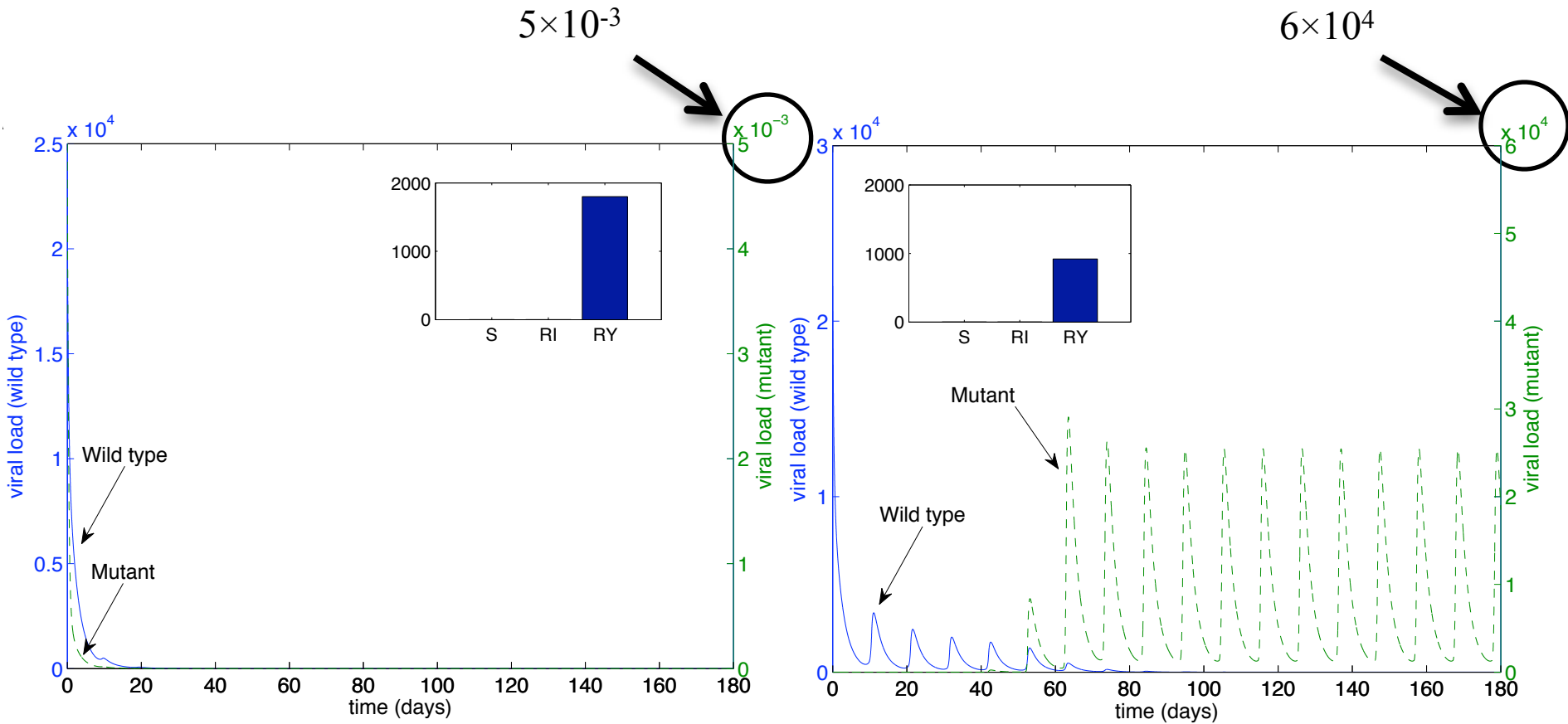


# Example

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- A patient is taking the triple-drug cocktail FTC (emtricitabine), ddl (didanosine) and EFV (efavirenz), can have
  - FTC: 6 days off, 17 subsequently on
  - ddl: 5 days off, 7.5 subsequently on ← *First drug to reach  $R_2$*
  - EFV: 9 days off, 22 subsequently on
- Therefore, during an 180 day induction period, a patient can have 13 holidays, each of which are 5 days long, followed by 7.5 days of strict therapy
- $[7.5+(5+7.5)\times 13+7.5=177.5<180]$ .

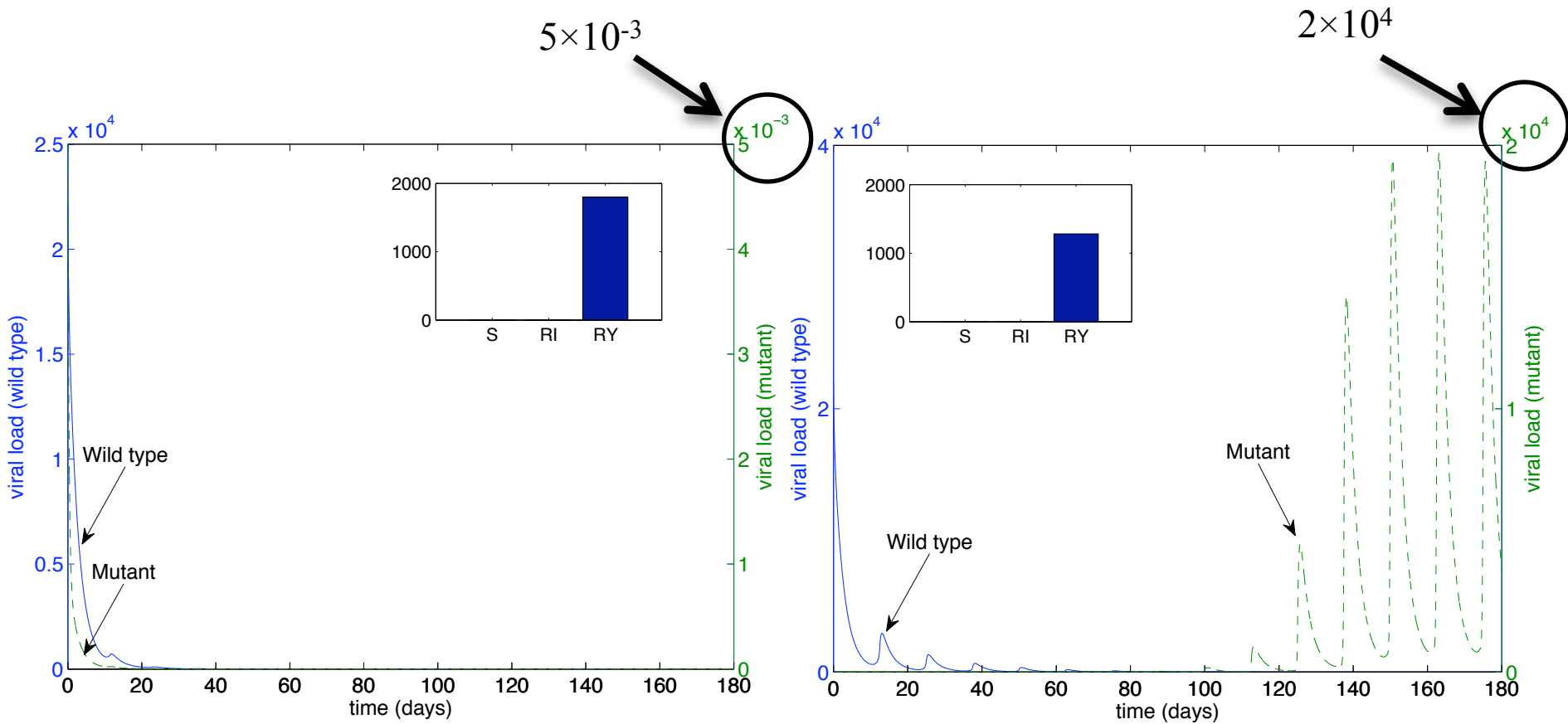
# Abacavir (ABC)



Taking the prescribed drug holidays

Missing one extra dose during each holiday

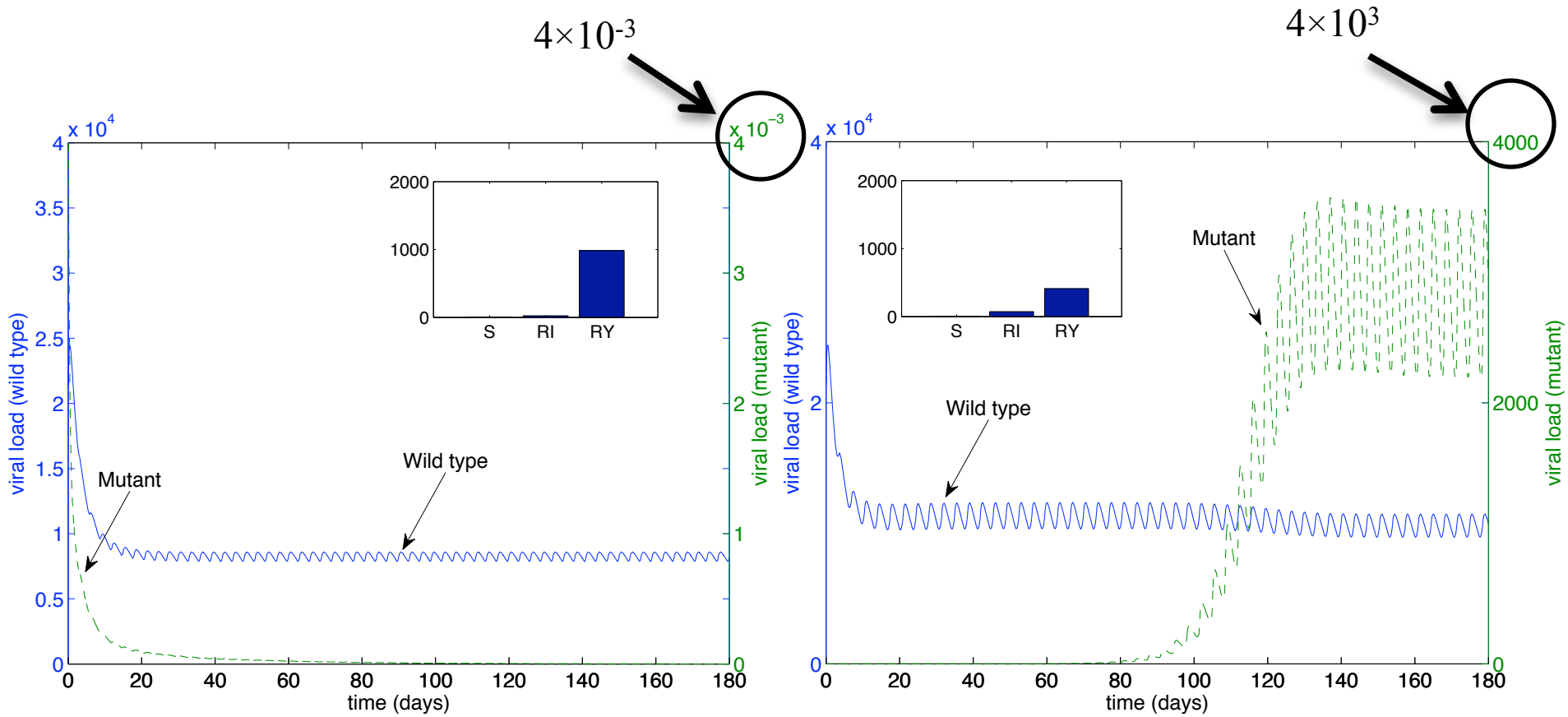
# Lamivudine (3TC)



Taking the prescribed drug holidays

Missing one extra dose during each holiday

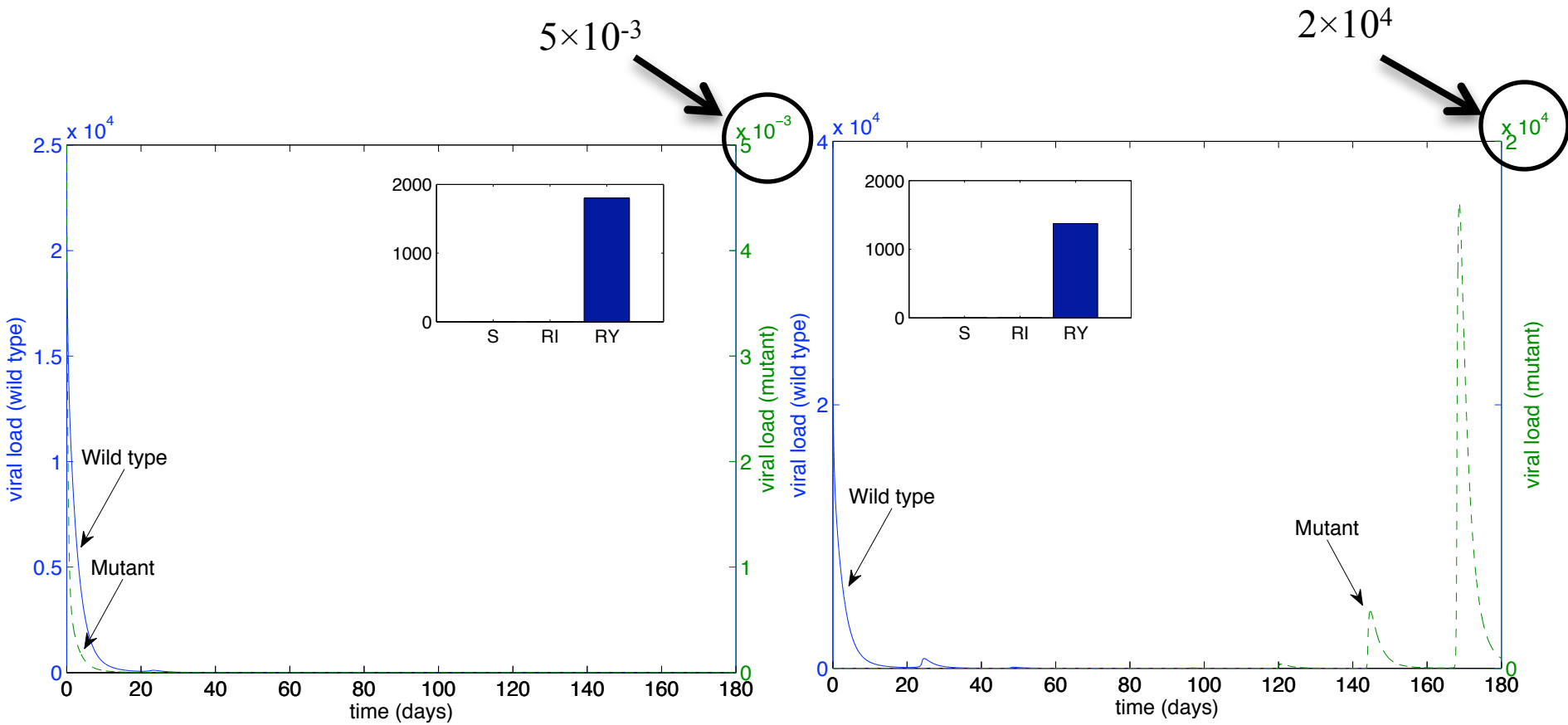
# Stavudine (d4T)



Taking the prescribed drug holidays

Missing one extra dose during each holiday

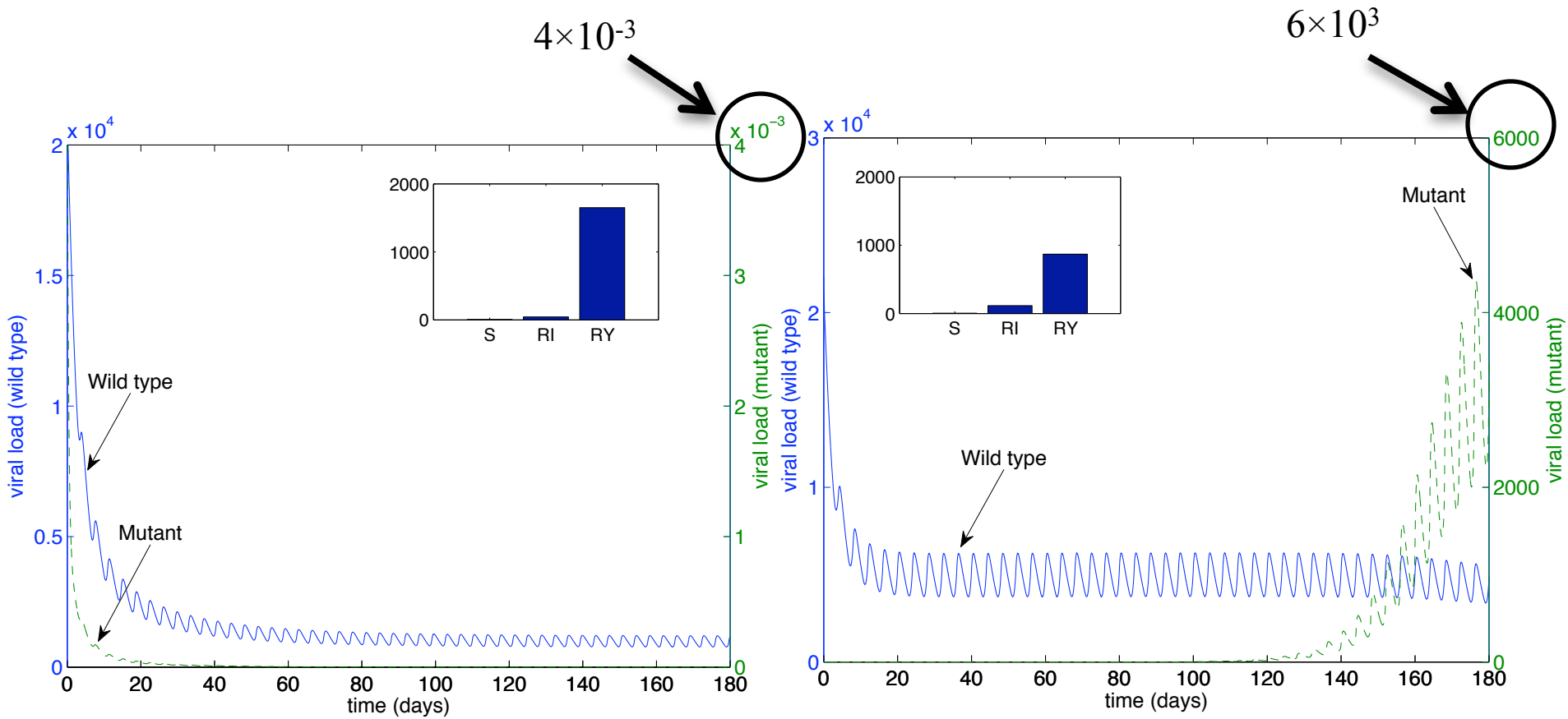
# Emtricitabine (FTC)



Taking the prescribed drug holidays

Missing one extra dose during each holiday

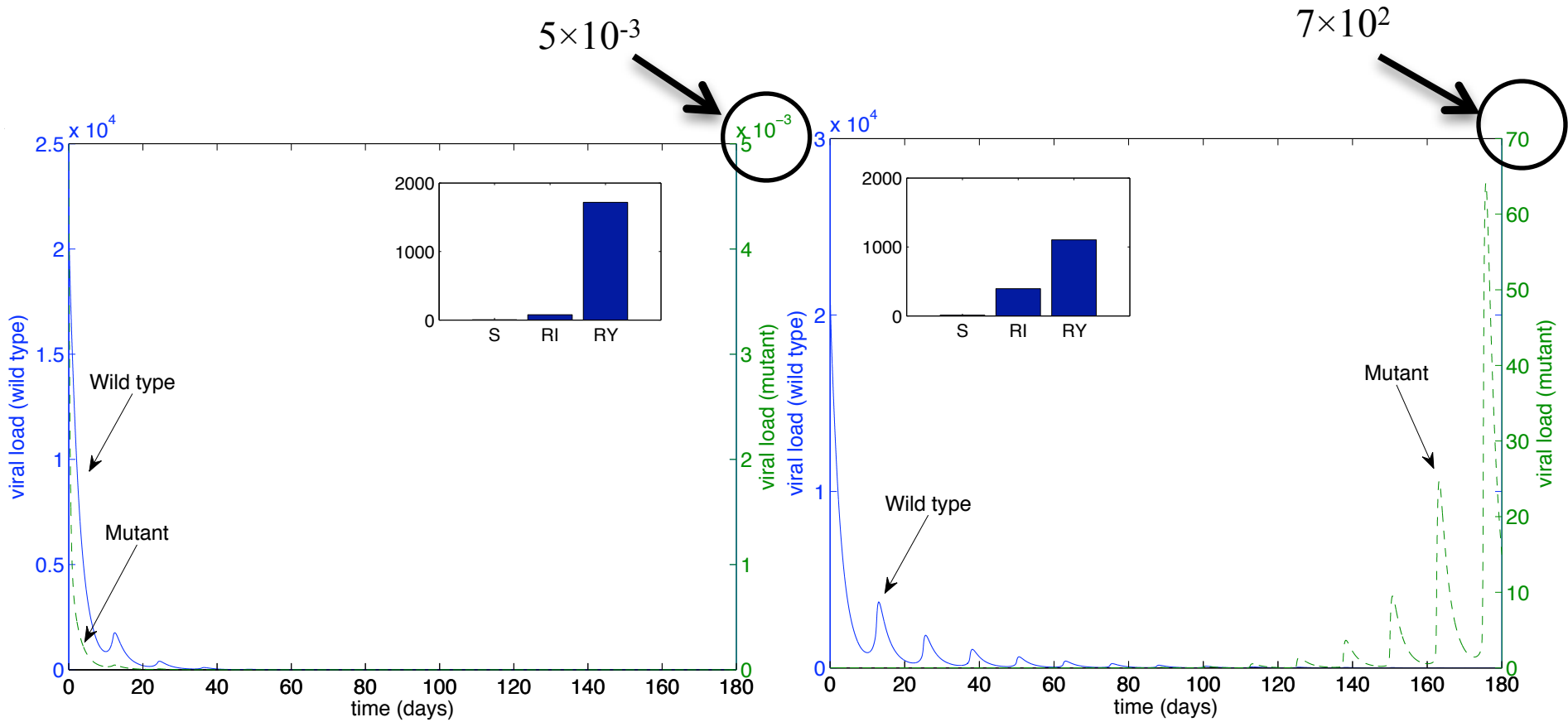
# Zidovudine (ZDV)



Taking the prescribed drug holidays

Missing one extra dose during each holiday

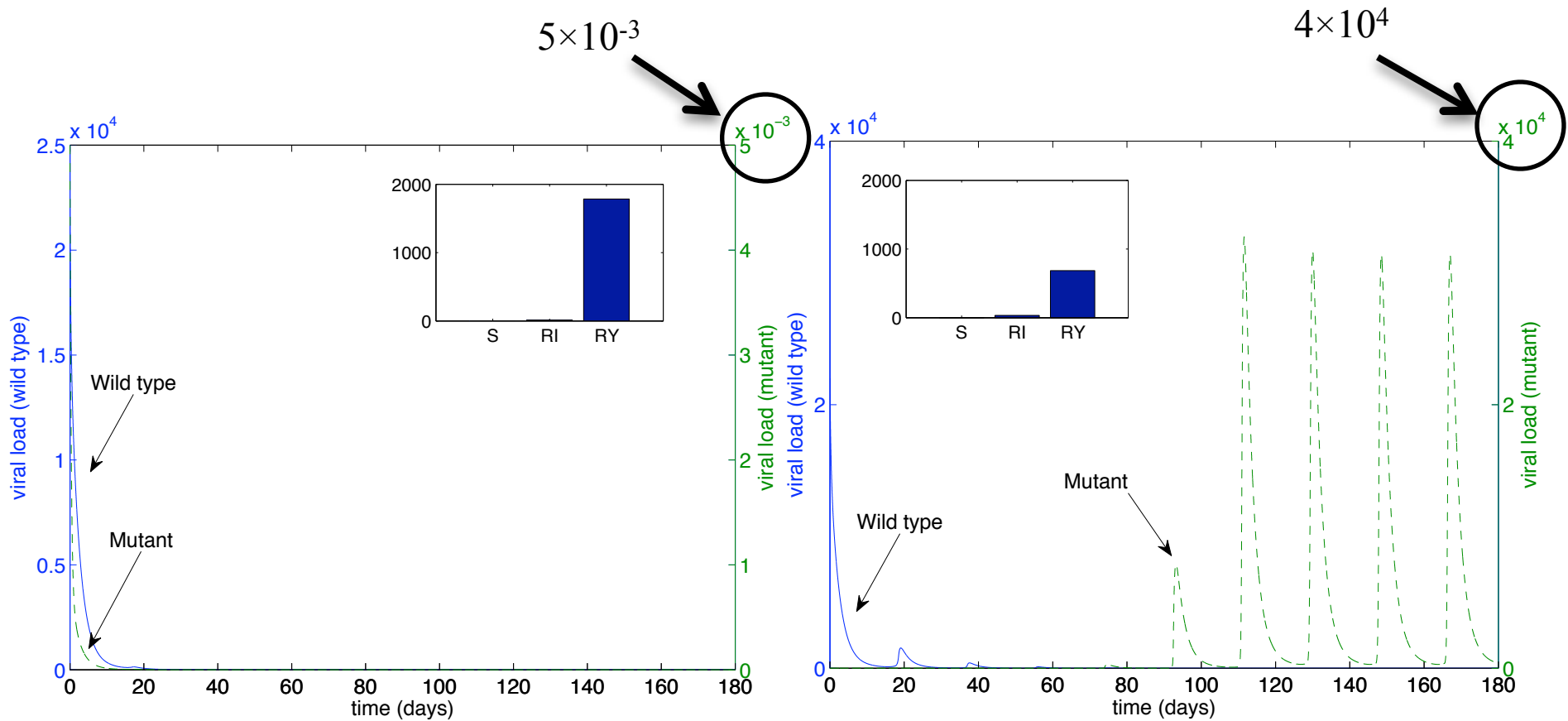
# Didanosine (ddI)



Taking the prescribed drug holidays

Missing one extra dose during each holiday

# Nevirapine (NVP)



Taking the prescribed drug holidays

Missing three extra doses during each holiday



# Drug Holidays

For a 180 day induction phase, you can have:

FDA-approved combination			Number of drug holidays	Length of each holiday (days)	Minimum subsequent therapy (days)
ABC*	3TC	NVP	16	3	7
ABC*	3TC	EFV	16	3	7
TDF	3TC*	EFV	14	3.5	8.5
ddI	3TC*	EFV	14	3.5	8.5
d4T*	3TC	EFV	50	1	2.5
d4T*	3TC	NVP	50	1	2.5
ddI*	FTC	EFV	13	5	7.5
TDF	FTC*	EFV	7	6	17
TDF	FTC	NVP*	9	5	12.5
ZDV*	3TC	ABC	44	1.33	2.66
ZDV*	3TC	EFV	44	1.33	2.66
ZDV*	3TC	NVP	44	1.33	2.66
ZDV*	3TC	TDF	44	1.33	2.66
ZDV*	DLV	3TC	44	1.33	2.66
ZDV*	DLV	ddI	44	1.33	2.66

\* denotes the drug which reaches  $R_2$  first.

# Comparison with clinical results

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- Long interruptions:
  - SMART trial (2006) - 16 month interruptions
  - DART trial (2008) - 12 week interruptions
- No benefit observed
- SMART trial was halted prematurely due to significant mortality and morbidity
- Observed a 2-fold risk of AIDS or death for treatment interruptions greater than 3 months
- Thus, long treatment interruptions do not appear viable.



# Short interruptions

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- 2-6 week breaks observed no benefit...  
...but no increase in viral resistance either
- <7 day interruptions resulted in only 5% of participants increasing HIV counts
- Thus, short breaks appear to be tolerable.



# FOTO: Five On, Two Off

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- A 2007 pilot study
- Guided in part by our earlier modelling
- Virologic suppression in 89.6% of patients
- Excellent adherence and strong preference for this adherence regimen
- FOTO only failed for ZDV, d4T and DLV, which is predicted by our results (max 1.33, 1 and 1.67 days, respectively)
- All others maintained 100% virologic suppression
- Results strongly correlate with our predictions.

# Summary

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- Inductive therapy is a regimen with an endpoint
- This presents special challenges to adherence
- Treatment interruptions need to be considered on a drug-specific basis
- For all PI-sparing FDA-approved combinations, we can determine the length and number of drug holidays that can be taken, as well as the number of doses that must be taken subsequently
- FOTO therapy is extremely promising and in line with our predictions.

# Conclusions

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- Induction therapy with partial adherence is tolerable
- However, the outcome depends on the specific drug cocktail
- If treatment interruptions occur, they must be short and followed by a strict period of dose taking
- FOTO is acceptable for all PI-sparing cocktails, except those containing ZDV, d4T or DLV, which can only tolerate extremely short drug holidays.