

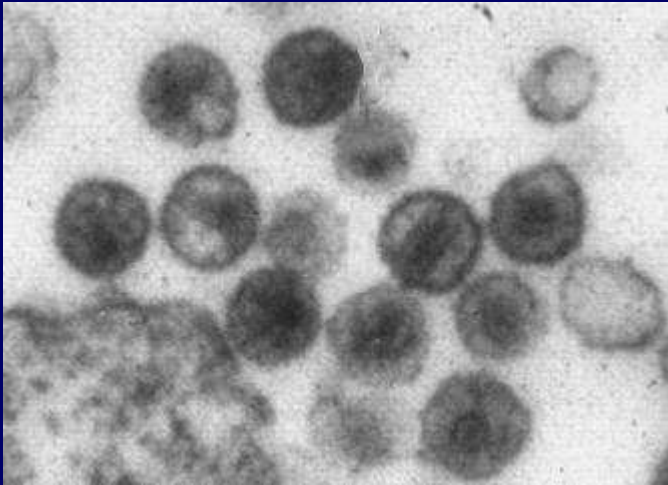
The Dynamics of Virus Infection and Treatment

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What is HIV infection?

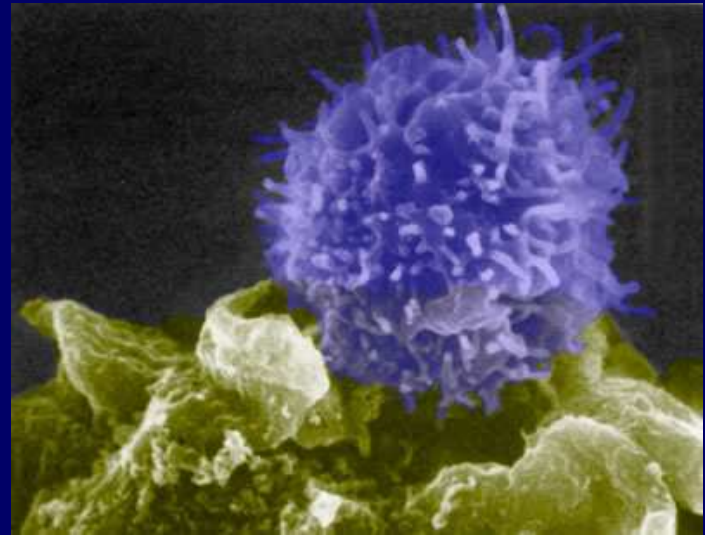
The virus



A retrovirus

Infects immune cells bearing:
CD4 & CCR5/CXCR4

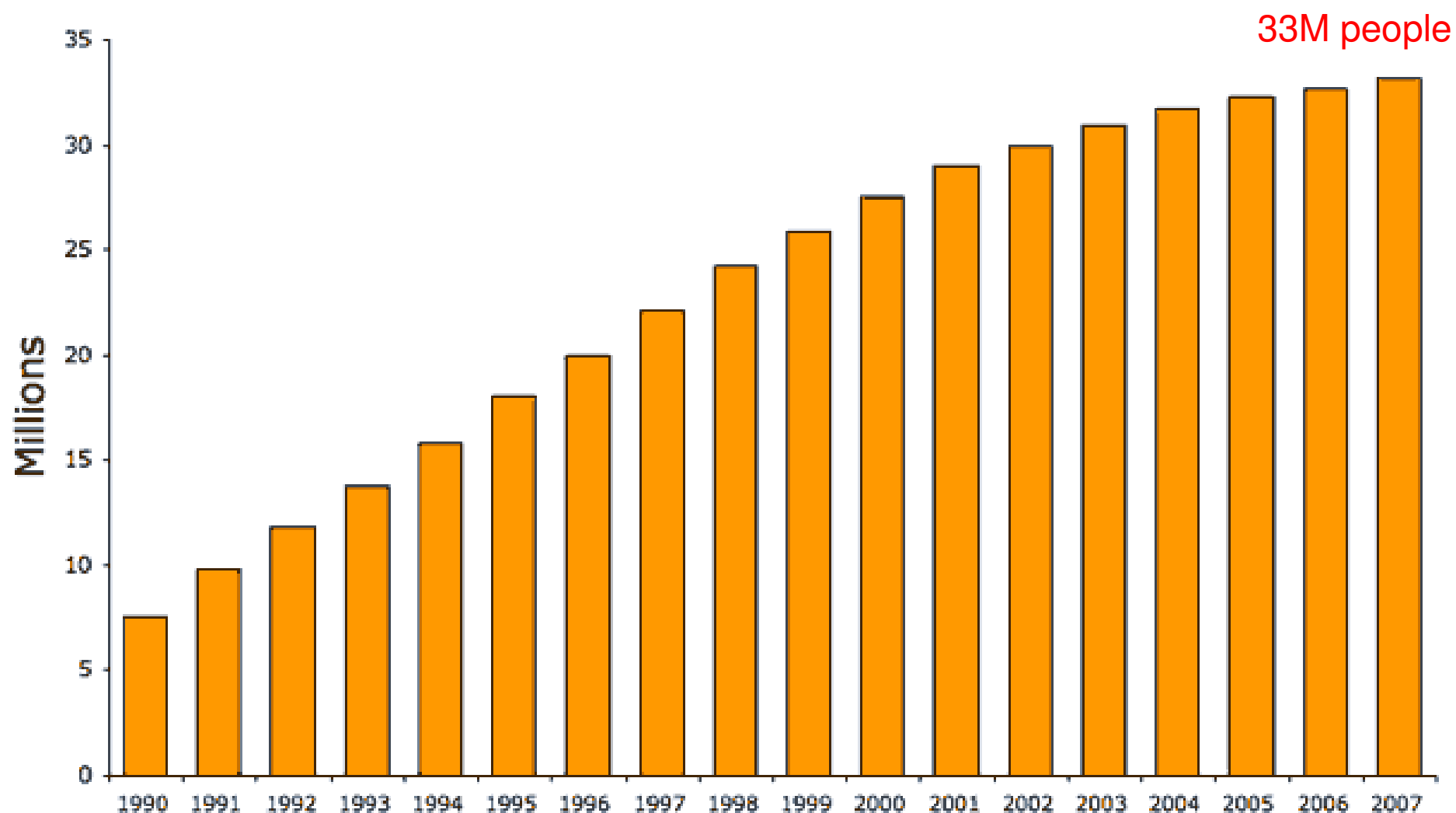
The host



CD4+ T-cells (or helper T cells)

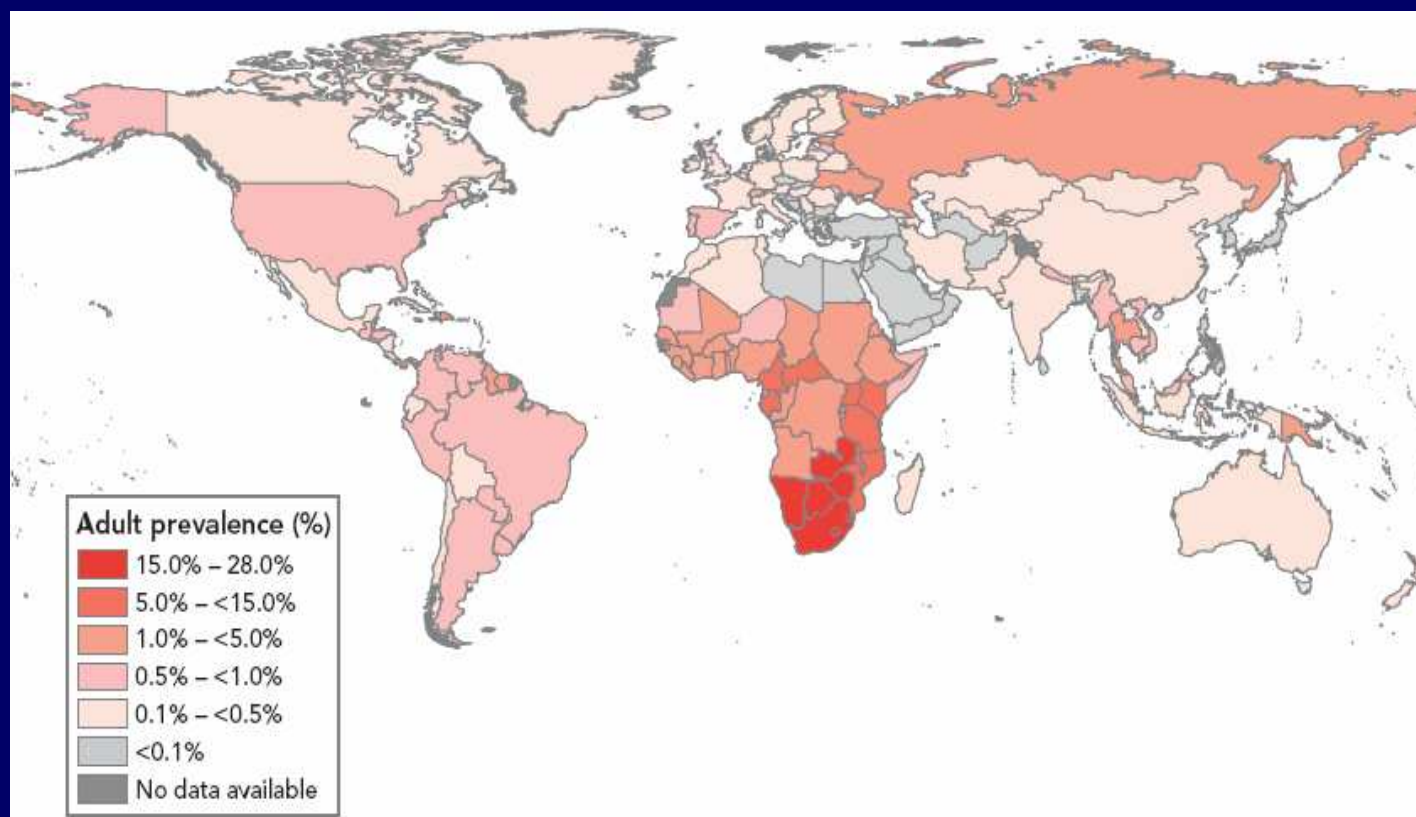
Macrophages and dendritic cells

Number of people infected

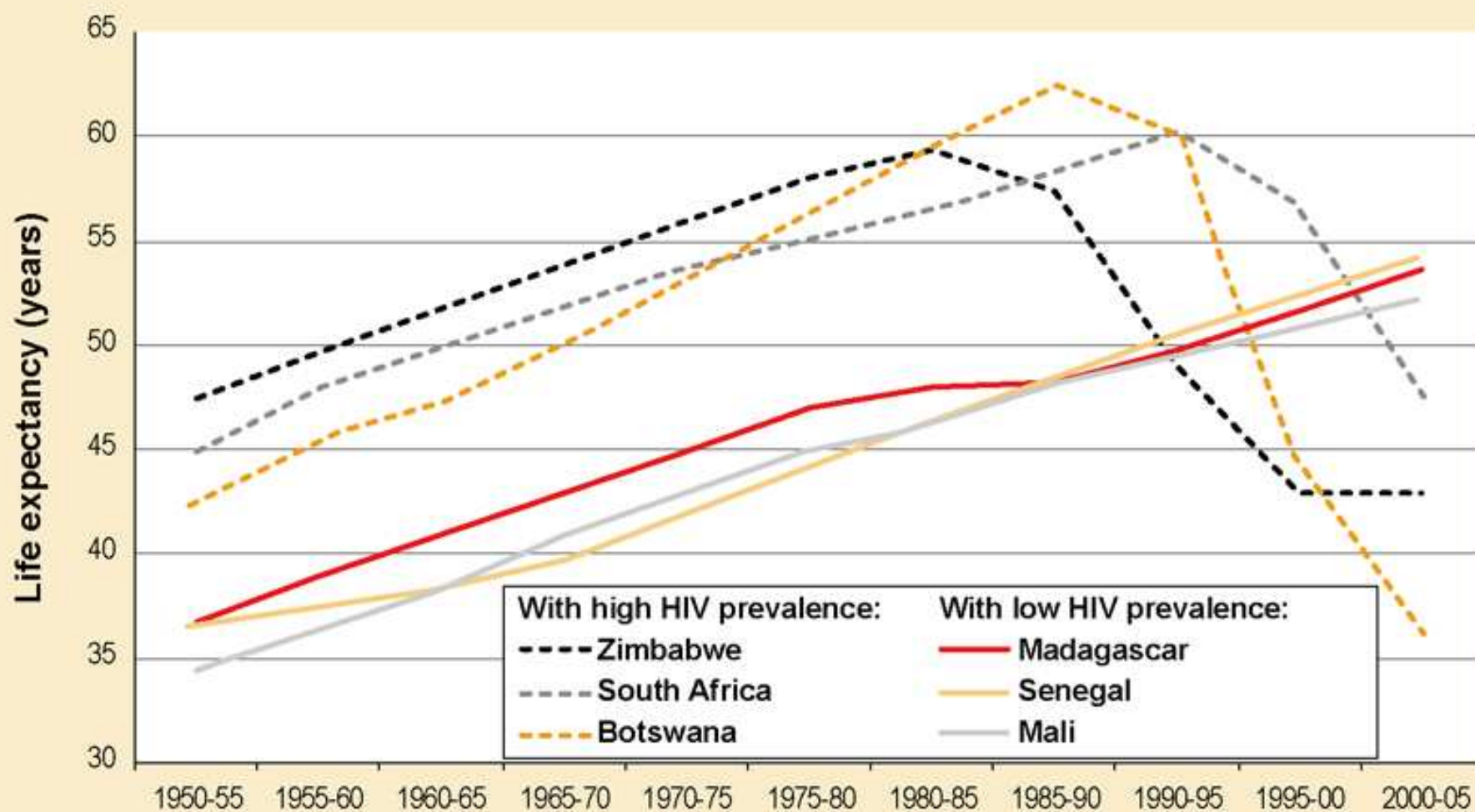


A global view of HIV infection

33 million people [30–36 million] living with HIV, 2007

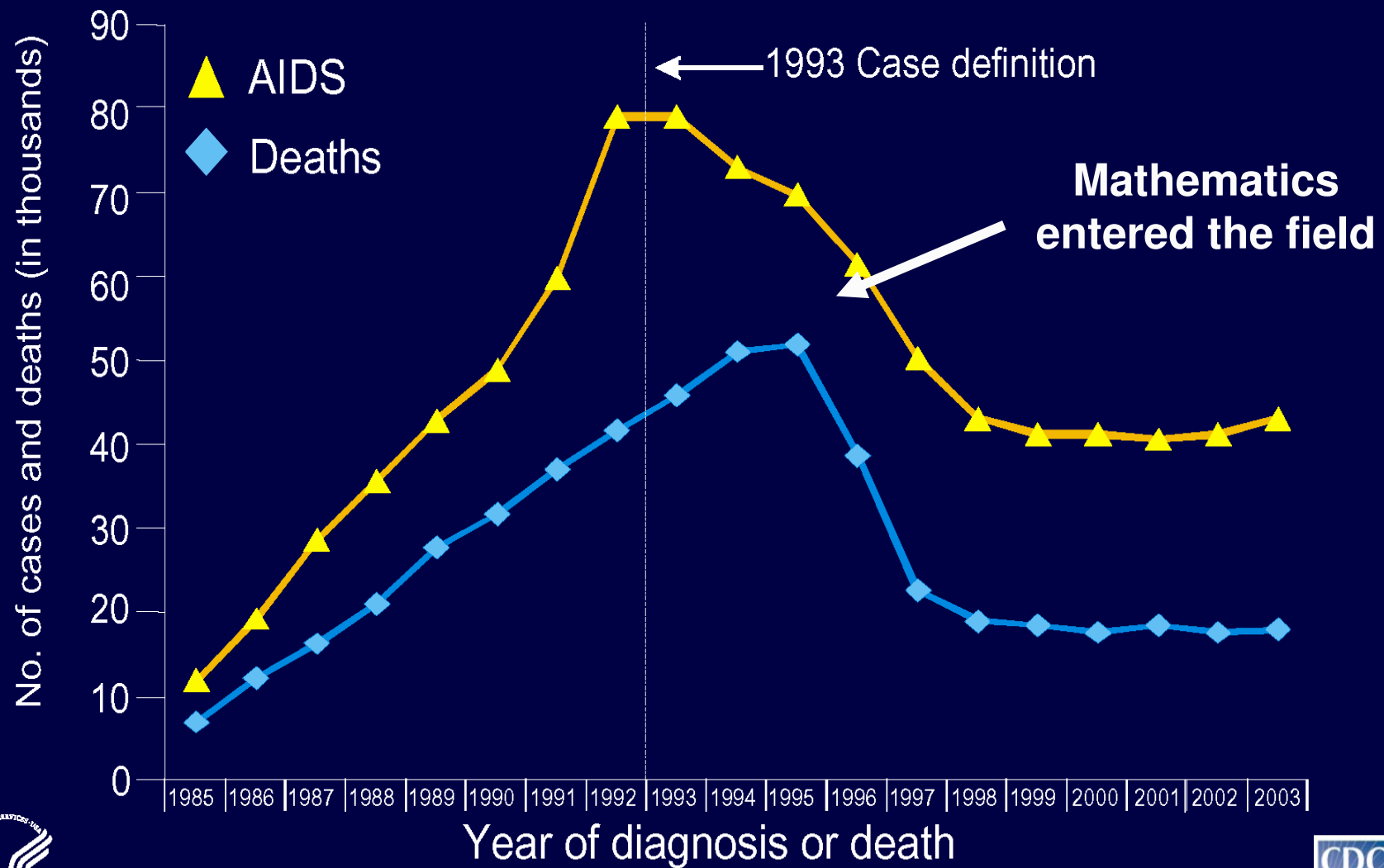


Life expectancy in African countries



Source: UN Department of Economic and Social Affairs (2001) *World Population Prospects, the 2000 Revision*

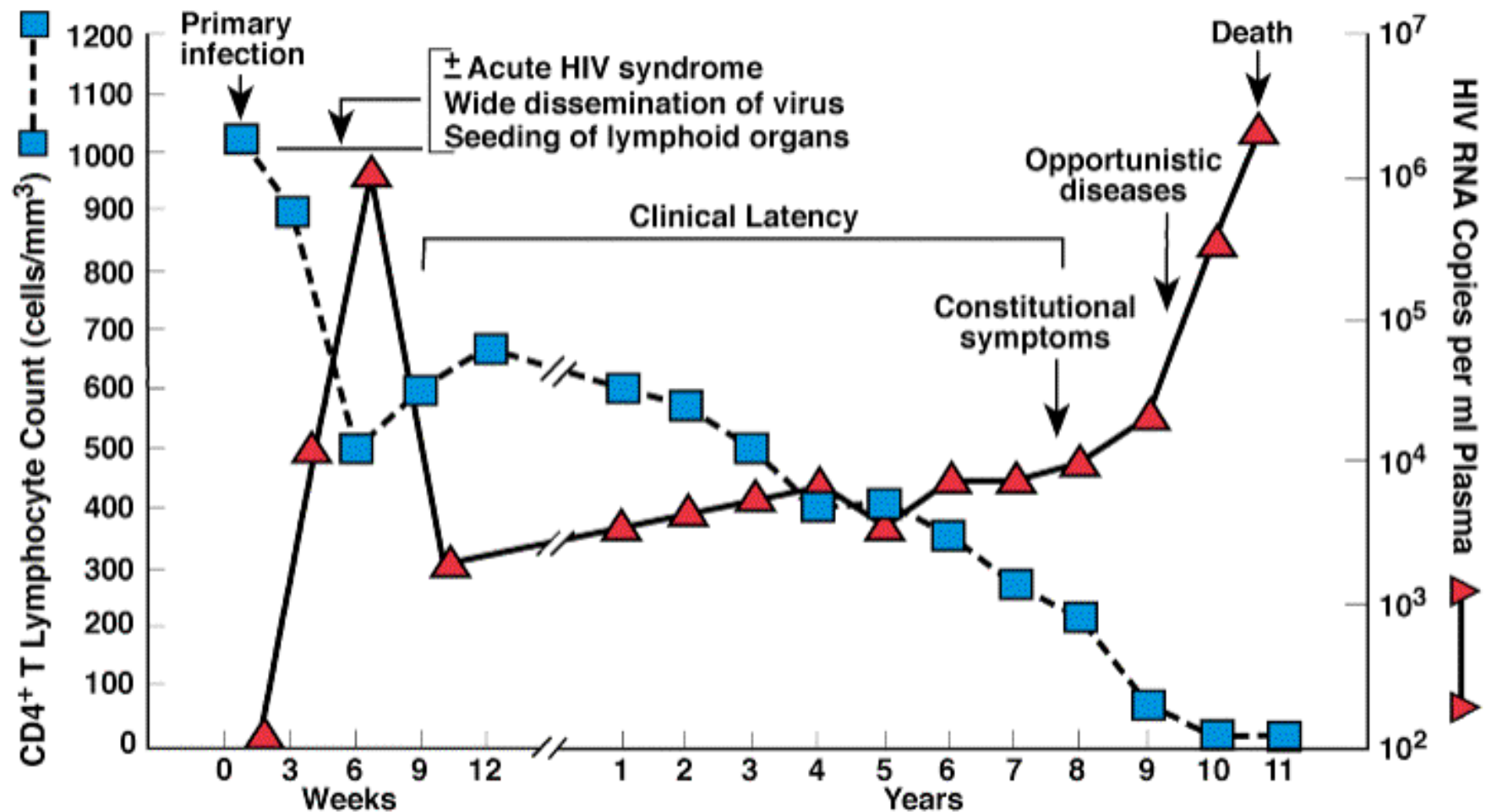
Estimated Number of AIDS Cases and Deaths among Adults and Adolescents with AIDS, 1985–2003—United States



Note. Adjusted for reporting delays.



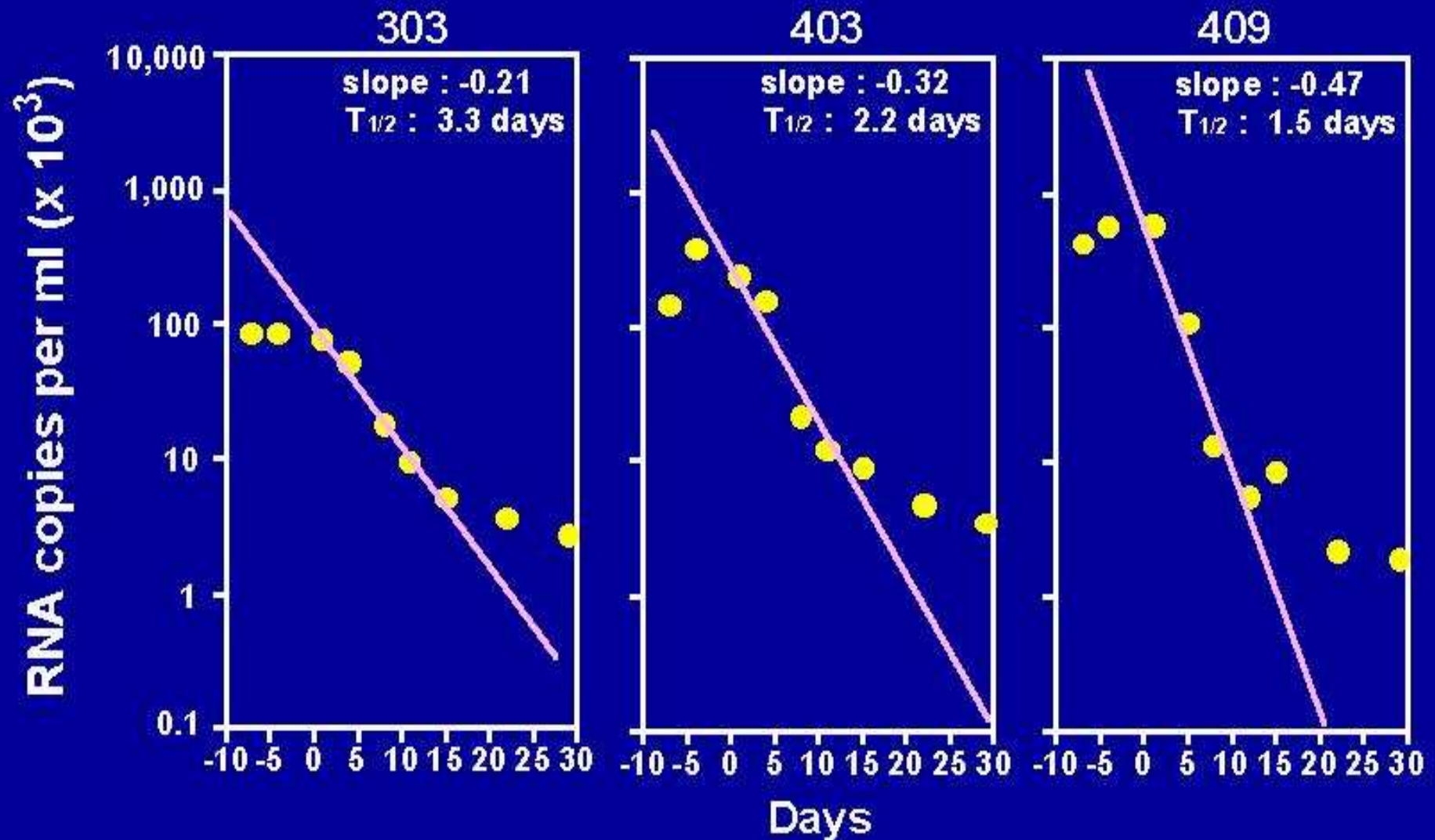
Typical Course of HIV Infection



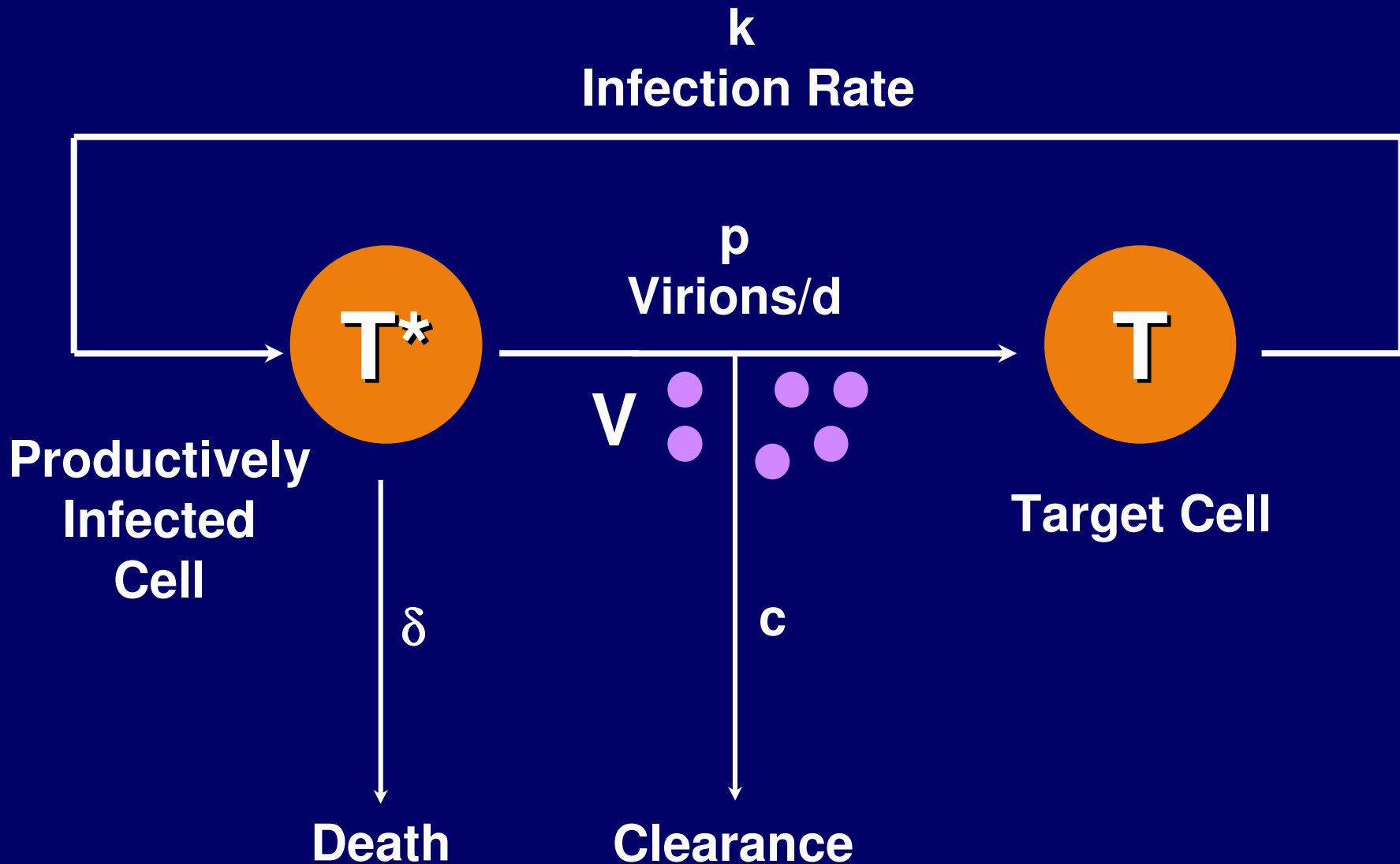
Modified From: Fauci, A.S., et al, *Ann. Intern. Med.*, 124:654, 1996

No treatment

HIV-1 protease inhibitor (ritonavir) given at t=0



Model of HIV Infection



Model of HIV Infection

$$\frac{dT(t)}{dt} = \lambda - dT - kTV$$

$$\frac{dT^*(t)}{dt} = kTV - \delta T^*$$

$$\frac{dV(t)}{dt} = N\delta T^* - cV$$

$$T(0) = T_0$$

$$T^*(0) = 0$$

$$V(0) = V_0$$

Variables

T Target Cell Density

T^* Infected Target Cell Density

V Virus Concentration

Parameters

λ Supply of target cells

d Net loss rate of target cells

k Infectivity rate constant

δ Infected cell death rate

$N\delta = p$ Virion production rate

c Virion clearance rate constant

Model derived by trying to explain effects of antiretroviral drugs; Here $T = \text{constant} = T_0$

$$\frac{dT^*(t)}{dt} = (1 - \epsilon_{RT}) k V_I T_0 - \delta T^*$$

$$\frac{dV_I(t)}{dt} = (1 - \epsilon_{PI}) N \delta T^* - c V_I$$

$$\frac{dV_{NI}(t)}{dt} = \epsilon_{PI} N \delta T^* - c V_{NI}$$

Drug efficacy

ϵ_{RT} ϵ_{PI}

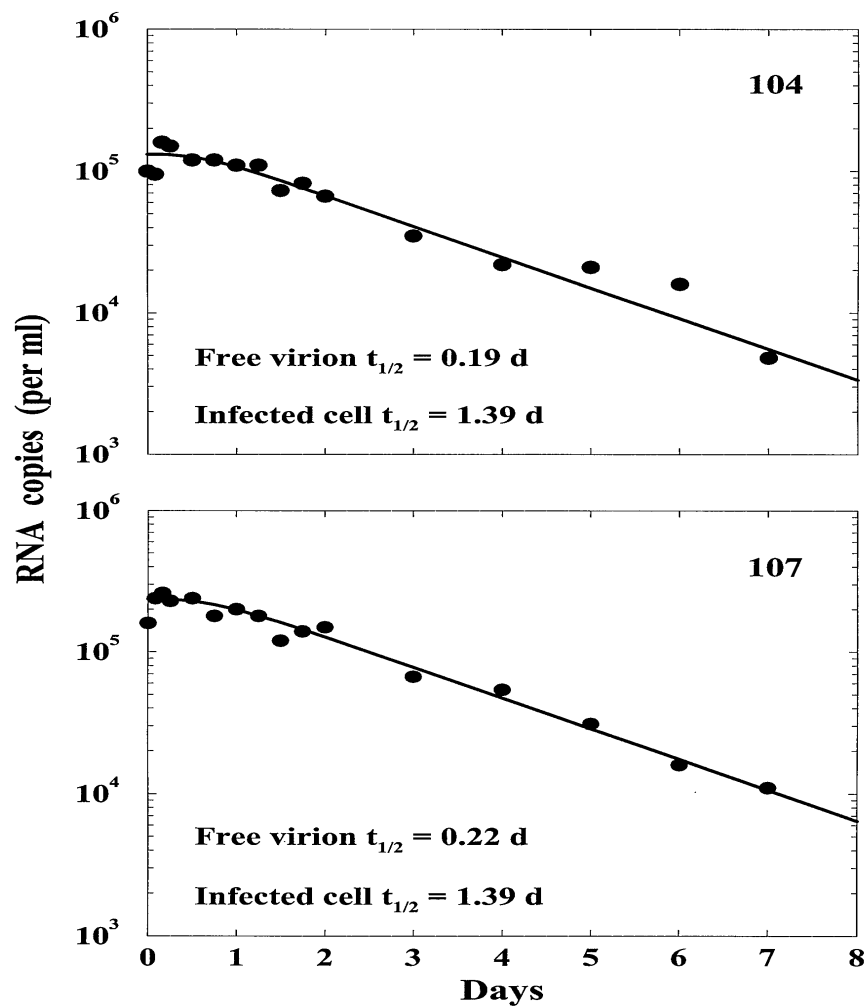
Subscripts:

“I”: infectious

“NI”: non-infectious

From *HIV-Dynamics in Vivo: ...*,
Perelson, *et al*, Science, 1996

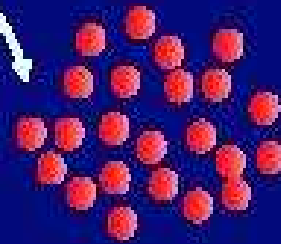
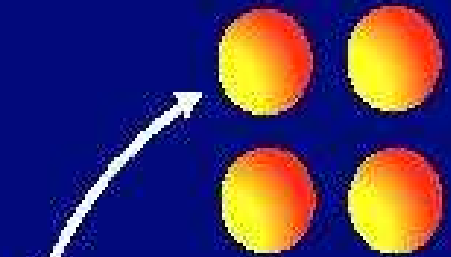
HIV-1: First Phase Kinetics



Perelson et al.
Science 271, 1582
1996

productively infected
CD4+ lymphocytes

$t_{1/2} < 1.5 \text{ d}$

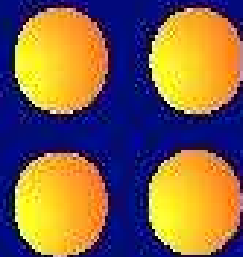


HIV-1

$t_{1/2} < 30 \text{ min} - 1 \text{ hr}$

10^{10} to 10^{12} virions/d
from
 10^7 to 10^9 T cells

uninfected, activated
CD4+ lymphocytes



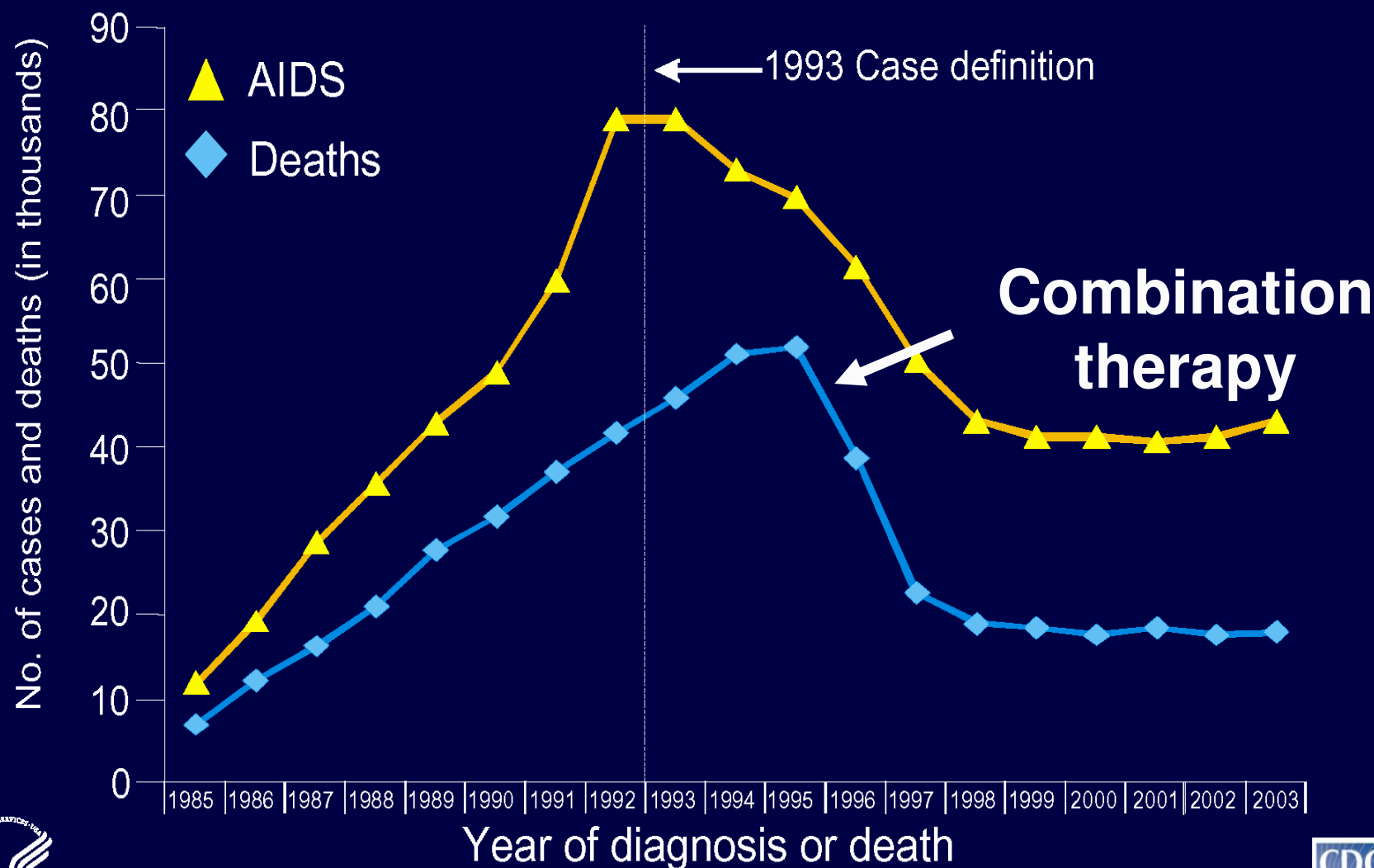
Implications

- 1 HIV infection is not a slow process.
- 1 Virus replicates rapidly and is cleared rapidly – can compute to maintain set point level $> 10^{10}$ virions produced/day.
- 1 Most cells infected by HIV are killed rapidly.
- 1 Rapid replication implies HIV can mutate and become drug resistant.
- 1 Calculations showed the need for triple combination therapy to overcome resistance.

Rate of generation of HIV-1 mutants

Base Changes	Probability of mutant	Number created/day	Number of possible mutants	Fraction of all possible mutants created/day
0	0.74	7.4×10^7	1	
1	0.22	2.2×10^7	3.0×10^4	1
2	0.033	3.3×10^6	4.5×10^8	7.4×10^{-3}
3	0.0033	3.3×10^5	4.5×10^{12}	7.4×10^{-8}

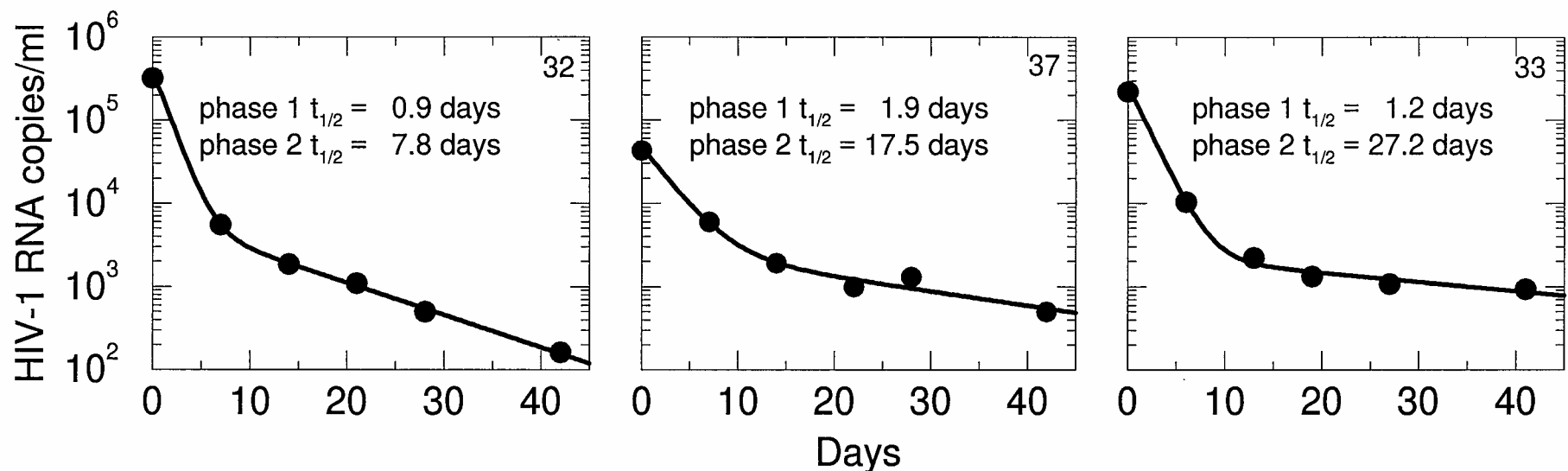
Estimated Number of AIDS Cases and Deaths among Adults and Adolescents with AIDS, 1985–2003—United States



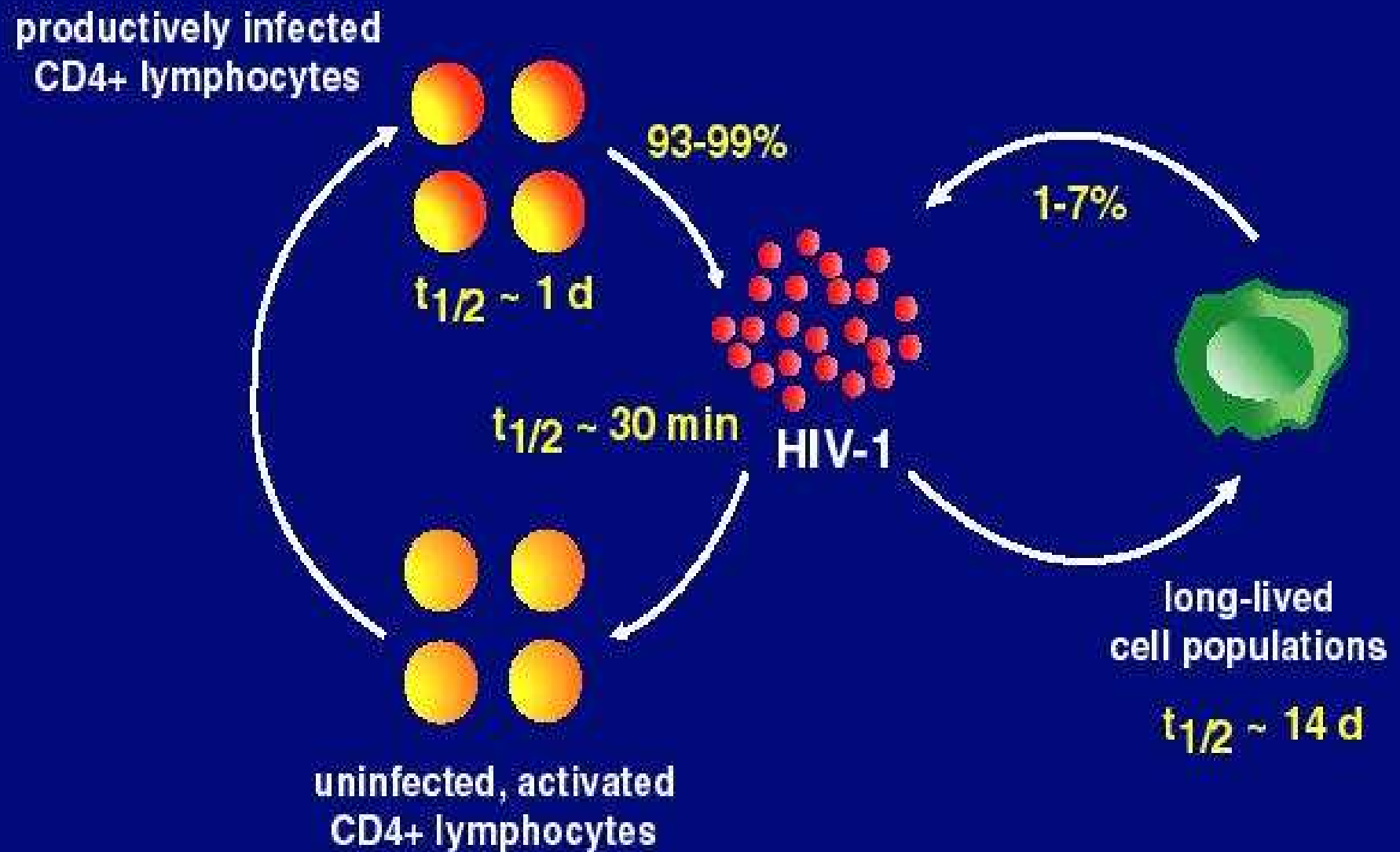
Note. Adjusted for reporting delays.

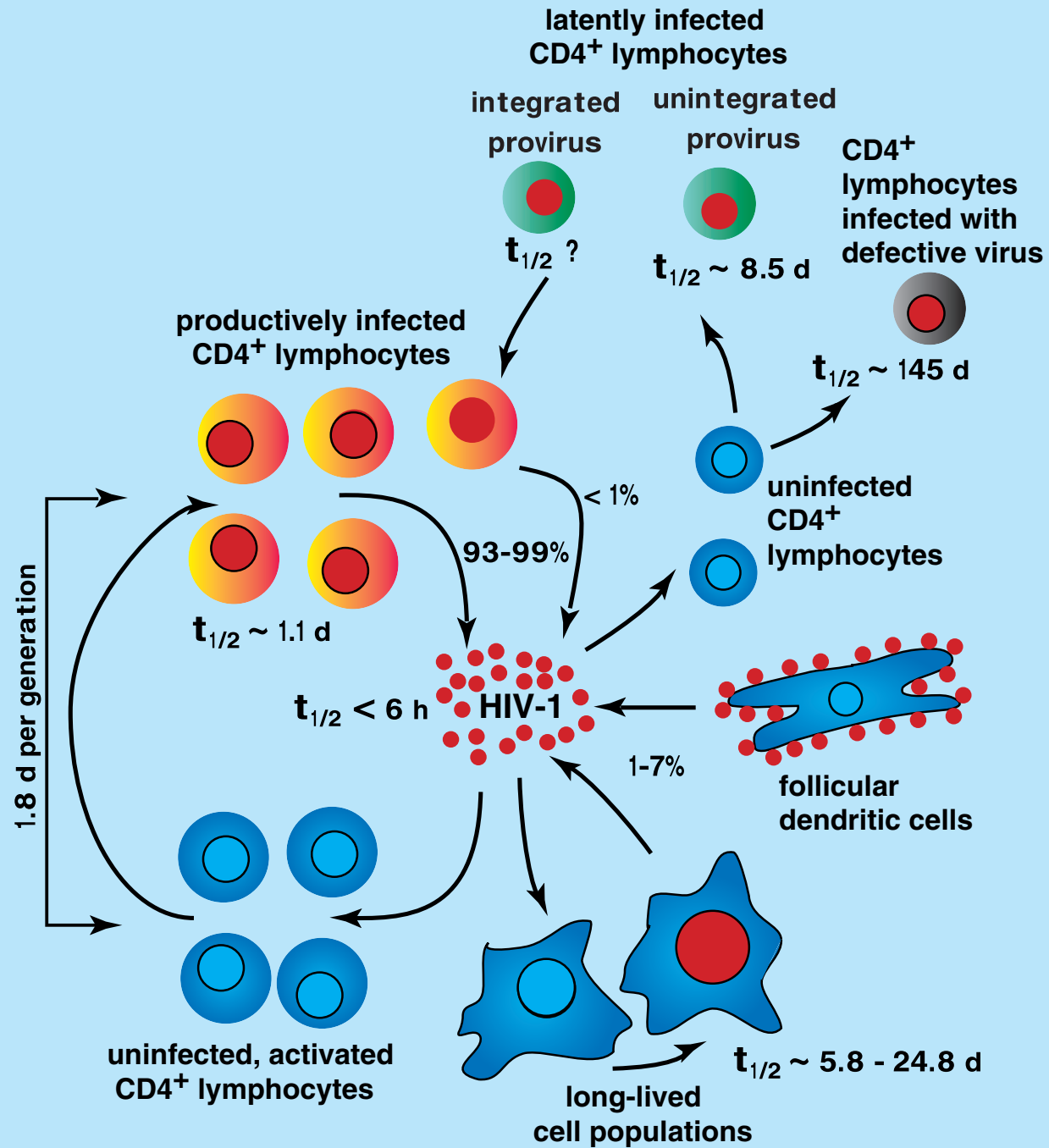


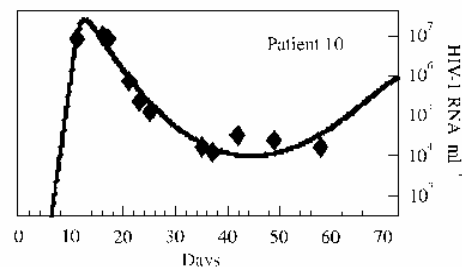
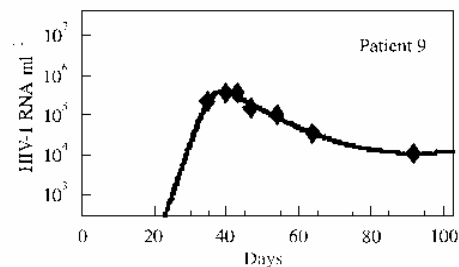
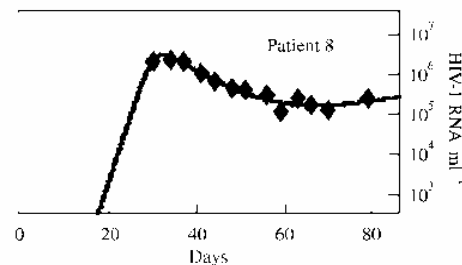
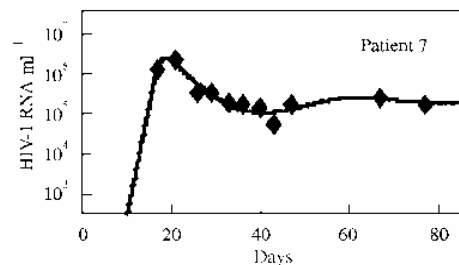
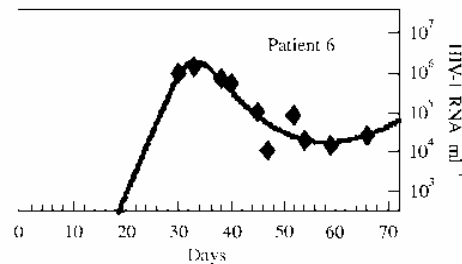
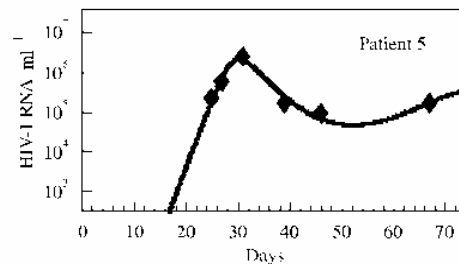
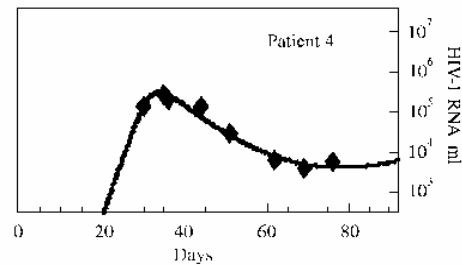
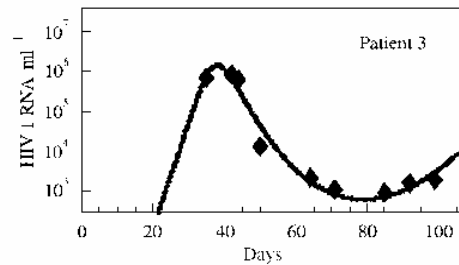
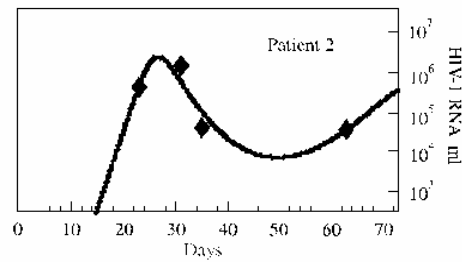
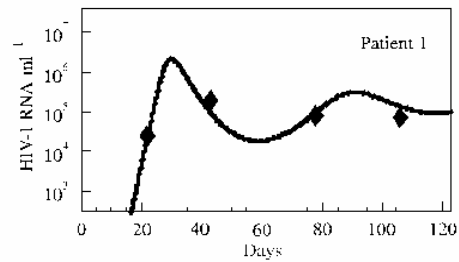
HIV-1: Two Phase Kinetics (Combination Therapy)



Perelson et al. Nature 387, 186 (1997)







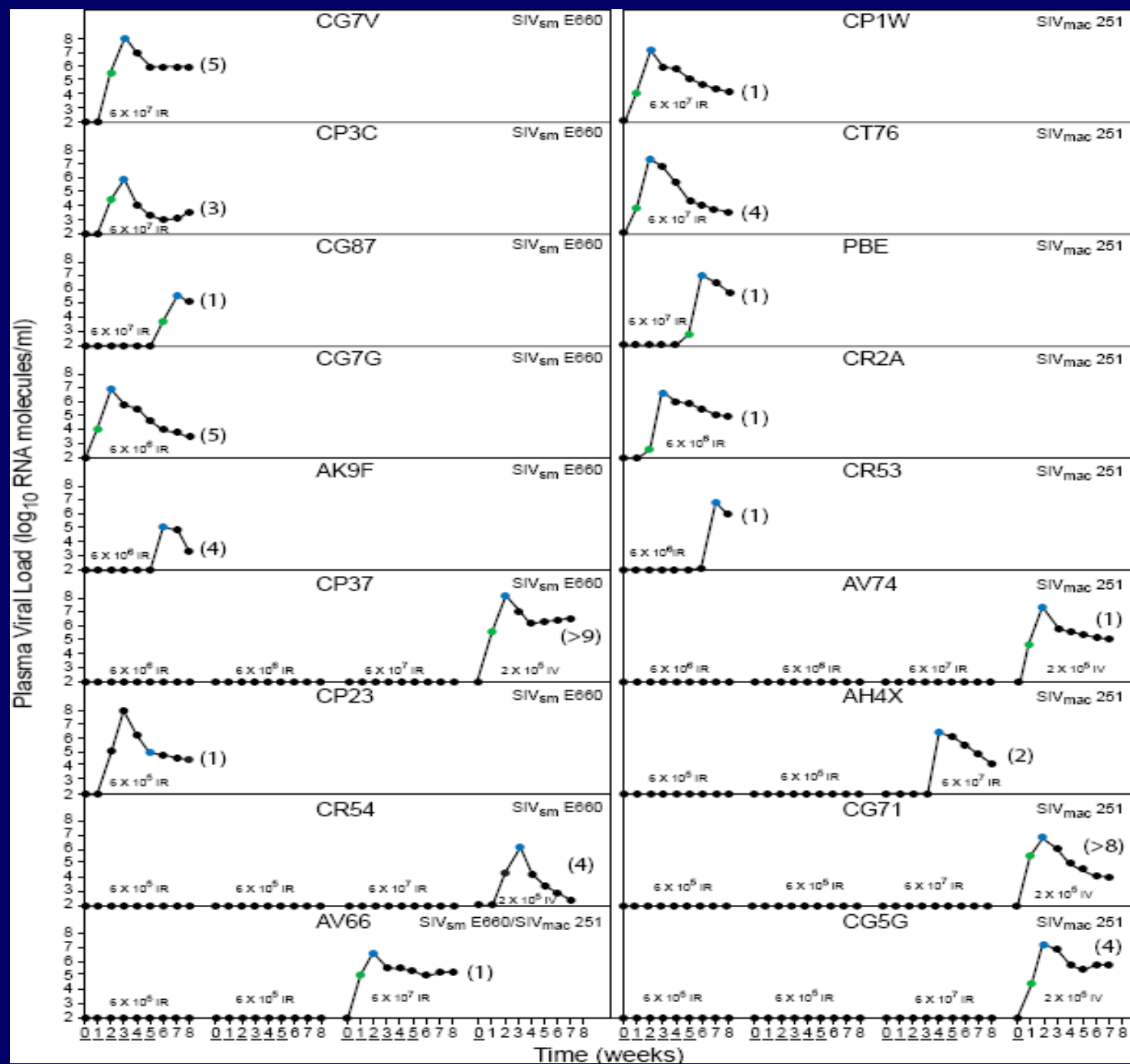
Model also fits
primary infection data.

Stafford et al.
J Theoret Biol.
203: 285 (2000)

Note virus is not
visible at early times =
eclipse phase
1 – 3 weeks in humans

Requirements for Infection

- 1 Prob. HIV transmitted/ sex act \sim .001 - .01
- 1 Inject low doses of SIV into monkeys – many times no noticeable infection results
- 1 Suggests early events are stochastic and not all encounters with virus lead to infection.

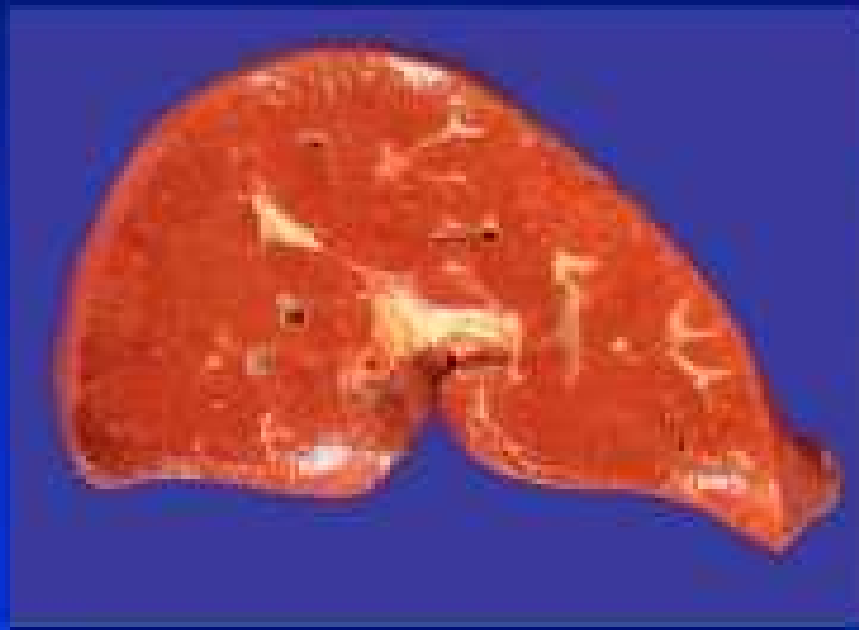


Hepatitis C Virus Infection

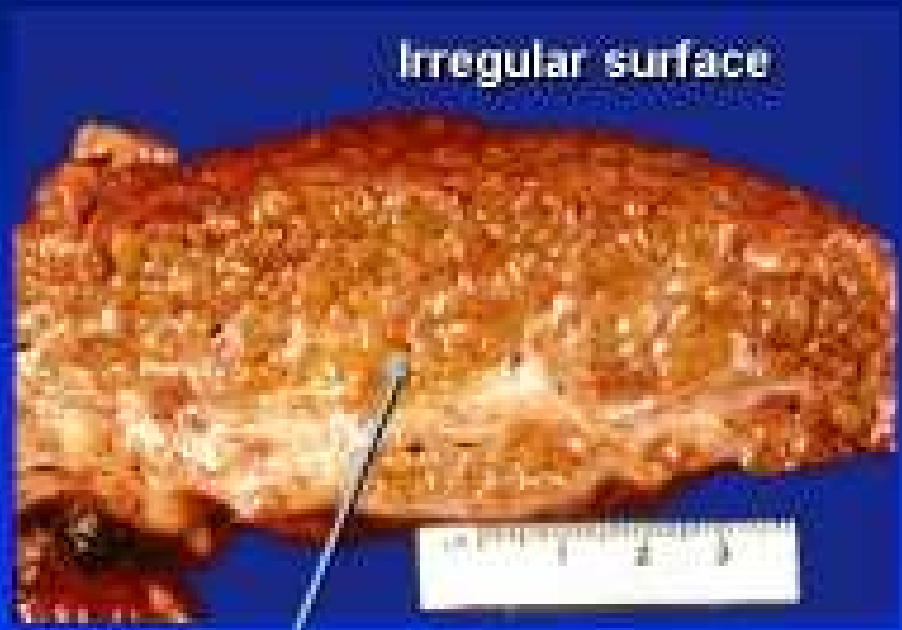
- 1 HCV is a positive strand RNA virus that infects the liver
- 1 It can lead to cirrhosis and liver cancer with a varying time course, from a few years (fulminant hepatitis) to > 30 years
- 1 ~ 4 million infected in the US
- 1 Can be treated with interferon (IFN), but ~50% of people fail to respond to best available therapy.
- 1 No vaccine available.

Cirrhosis

Normal



Cirrhosis



Irregular surface

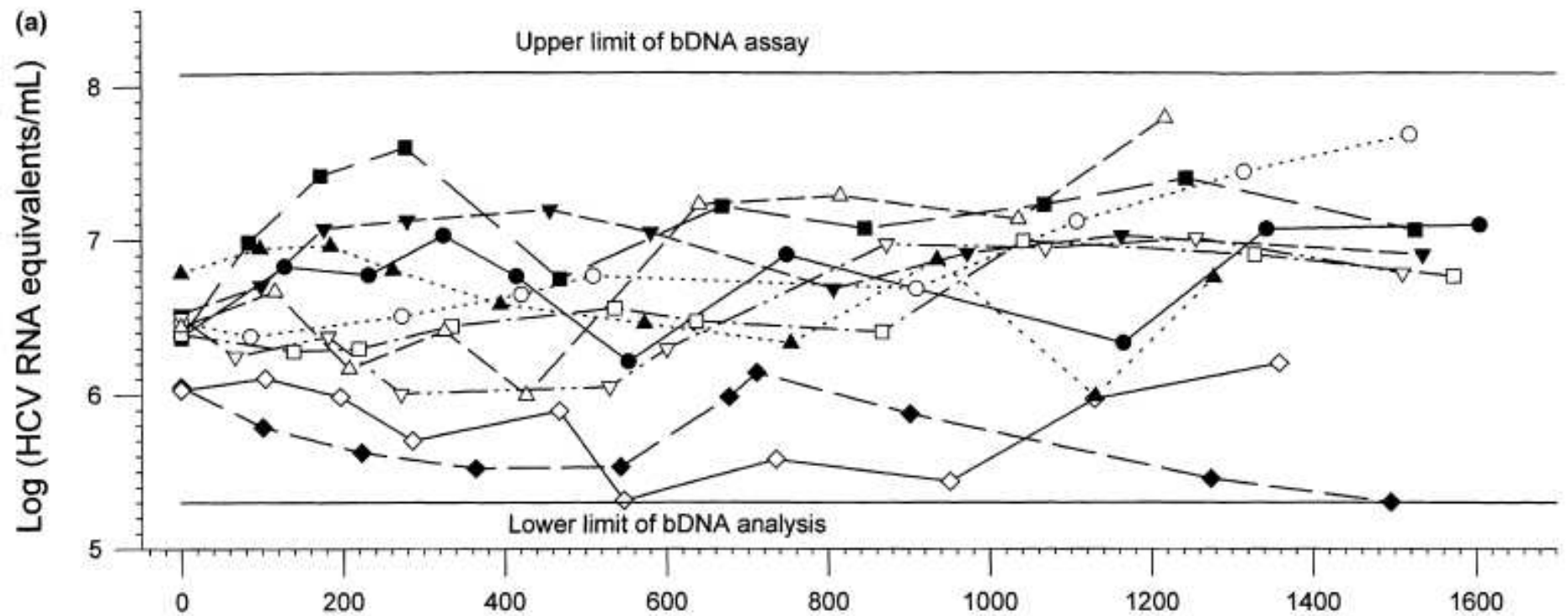
Nodules



HCV treatment

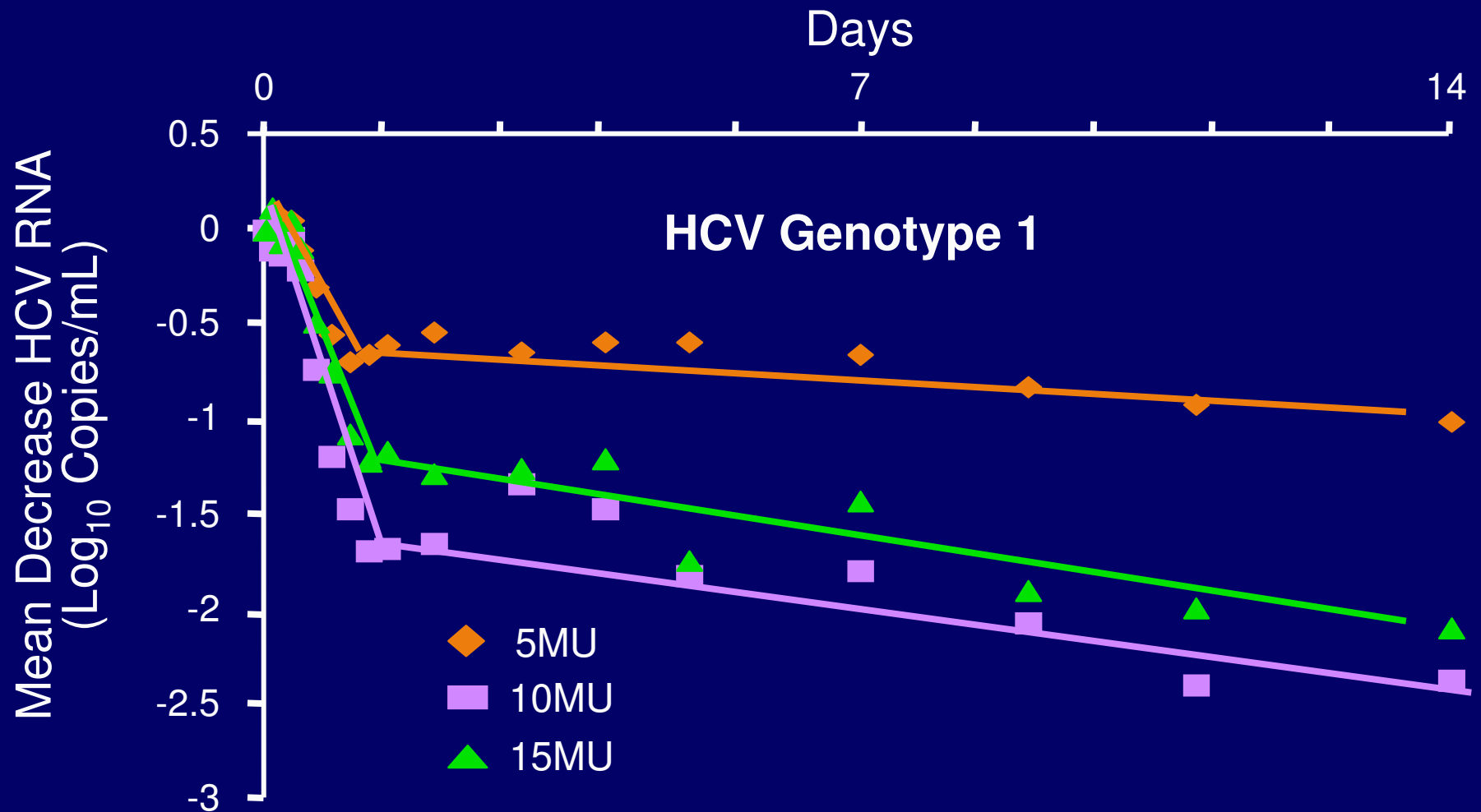
- 1 Current standard therapy, consisting of pegylated interferon and ribavirin, is effective in only ~50% of treated HCV patients; serious adverse effects in some patients
- 1 A number of small molecule drugs, including **protease inhibitors** and **polymerase inhibitors**, are being developed and evaluated in clinical trials.

HCV RNA stable in chronic infection



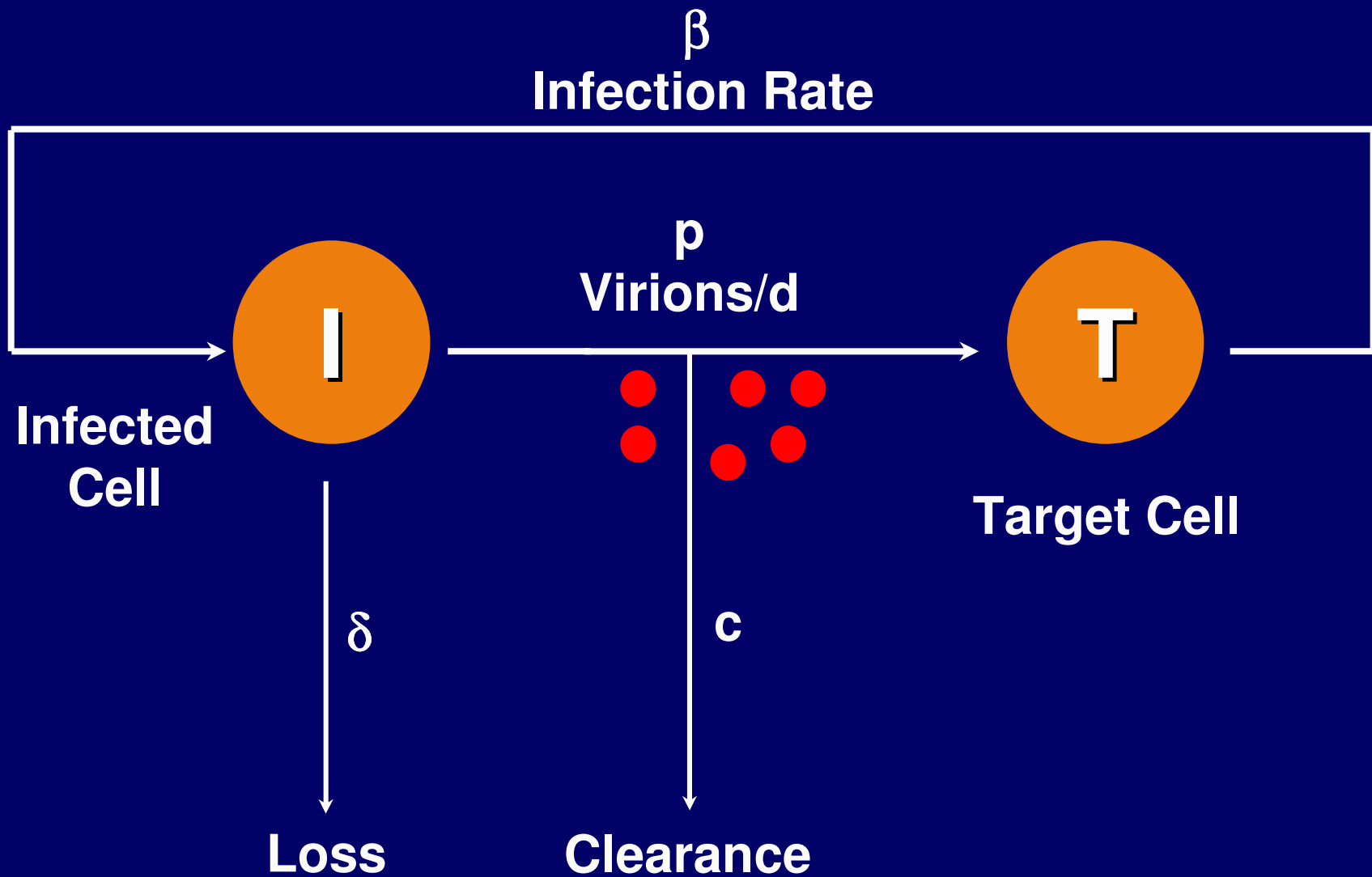
Yeo et al. J. Viral Hepat. 8:256 2001

Mean Decrease in HCV RNA Levels Over First 14 Days of Daily IFN- α Treatment

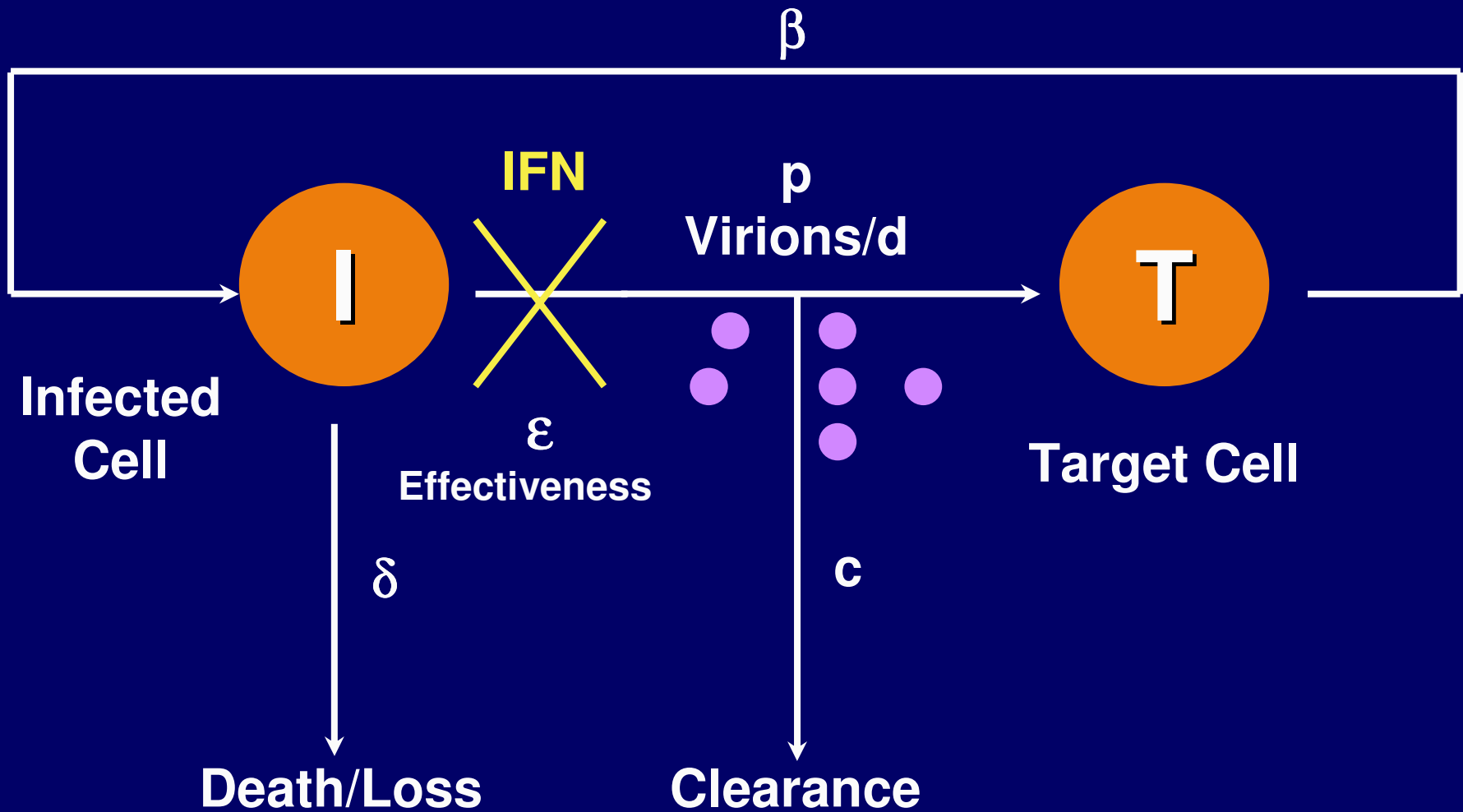


Lam N. DDW. 1998 (abstract L0346).

Model of HCV Infection



IFN Partially Blocks Production



IFN Effectiveness in Blocking Production

1 Let ε = *effectiveness* of IFN in blocking production of virus

- $\varepsilon = 1$ is 100% effectiveness
- $\varepsilon = 0$ is 0% effectiveness

1
$$dV/dt = (1 - \varepsilon)pI - cV$$

Early Kinetic Analysis

- 1 Before therapy, assume steady state so that $pl_0 = cV_0$. Also, assume at short times, $l = \text{constant} = l_0$, so that

$$dV/dt = (1-\varepsilon)pl - cV = (1-\varepsilon)cV_0 - cV, \quad V(0) = V_0$$

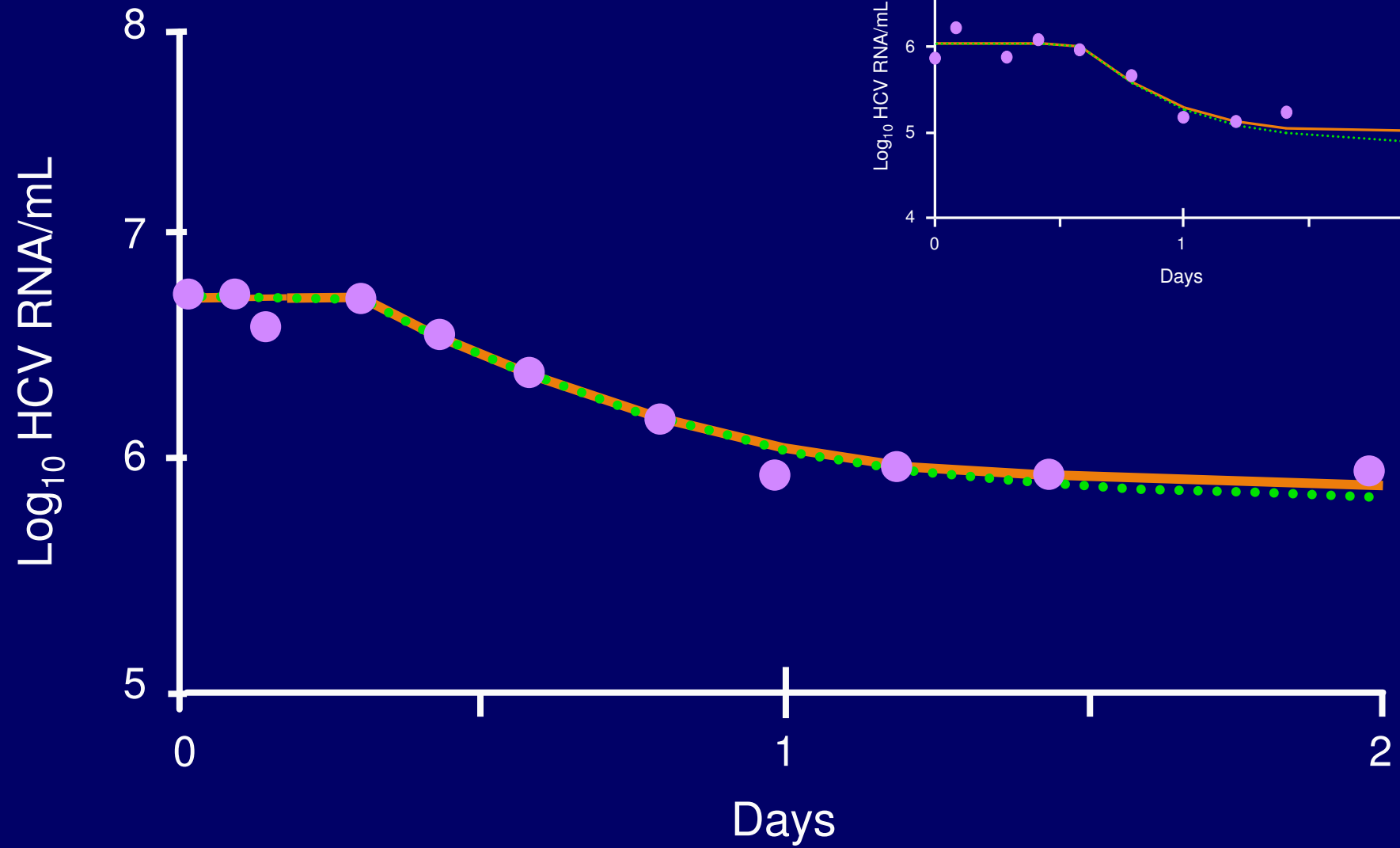
- 1 Model predicts that after therapy is initiated, the viral load will initially change according to:

$$V(t) = V_0[1 - \varepsilon + \varepsilon \exp(-ct)]$$

- 1 This equation can be fit to data and **c** and **ε** estimated.
- 1 This suggests drug effectiveness can be determined within the first few days of treatment!

10MU

15MU

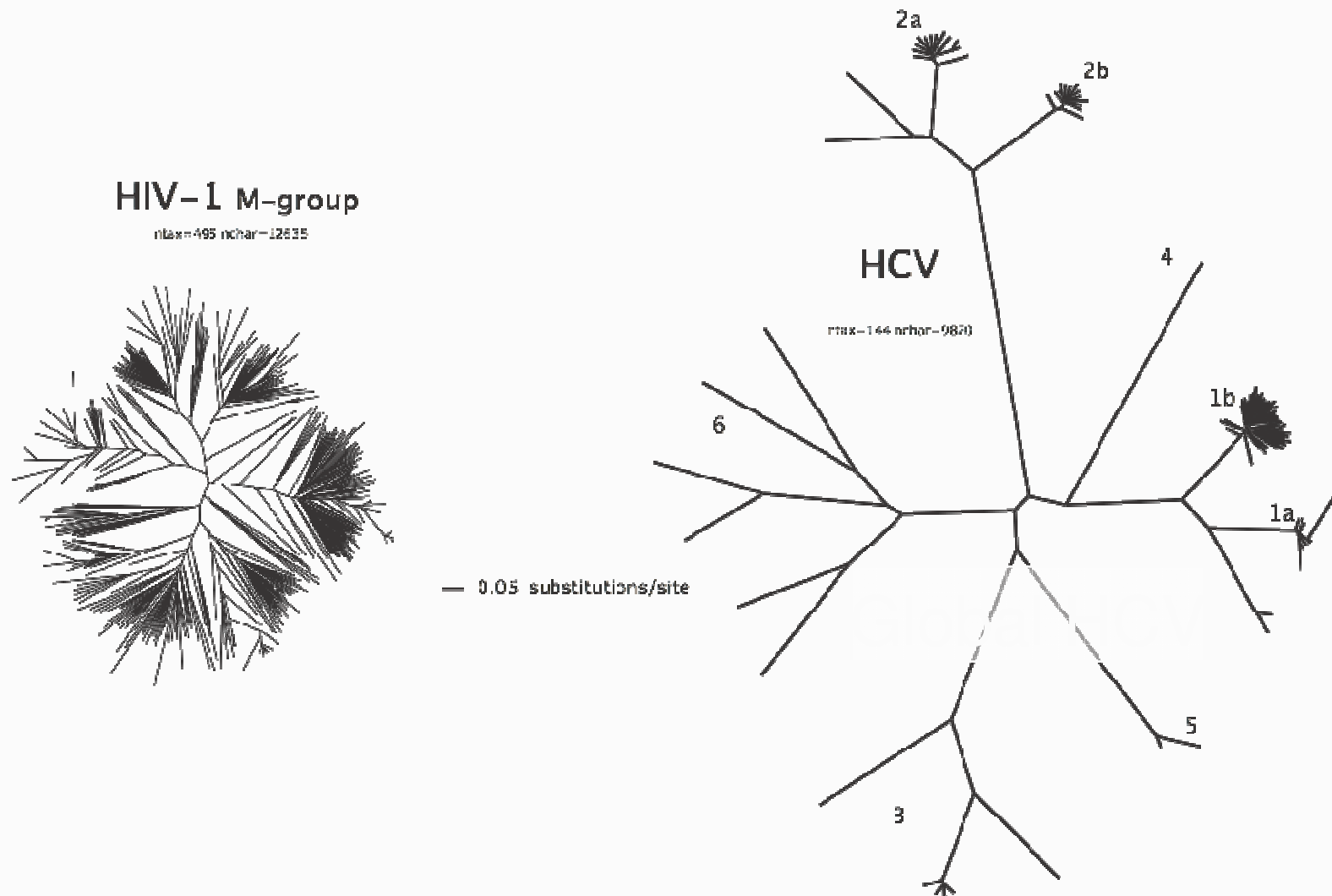


Viral Kinetics of HCV Genotype 1

	Drug Efficacy	Viral Clearance Constant (1/d)	Half-life of Virions (Hours)	Production & Clearance Rates (10¹² Virions/d)
5MU	81 ± 4%	6.2 ± 0.8	2.7	0.4 ± 0.2
10MU	95 ± 4%	6.3 ± 2.4	2.6	2.3 ± 4
15MU	96 ± 4%	6.1 ± 1.9	2.7	0.6 ± 0.8

$t_{1/2}$ estimates independently validated;
Ramratnam et al. Lancet 1999

Inter-subtype distances are greater for HCV than for HIV-1



Standard Model of HCV Dynamics

Equations

$$\frac{dT}{dt} = \lambda - dT - \beta VT$$

$$\frac{dI}{dt} = \beta VT - \delta I$$

$$\frac{dV}{dt} = (1 - \varepsilon)pI - cV$$

Variables

T Target Cell Density
 I Infected Cell Density
 V Virus Concentration

Parameters

λ Supply of target cells
 d Net loss rate of target cells
 β Infectivity rate constant
 δ Infected cell death rate
 ε Drug efficacy
 p Virion production rate
 c Virion clearance rate constant

Initial Conditions

$T(0) = T_0$ $V(0) = V_0$
 $I(0) = I_0$

Solution: Change in Viral Load

- 1 Assuming $T = T_0 = \text{constant}$, and pretreatment steady state $\beta T_0 = c\delta/p$

$$V(t) = \frac{1}{2}V_0 \left[\left(1 - \frac{c + \delta - 2\varepsilon c}{\theta}\right) e^{-\lambda_1(t-t_0)} + \left(1 + \frac{c + \delta - 2\varepsilon c}{\theta}\right) e^{-\lambda_2(t-t_0)} \right]$$

where

$$\lambda_1 = \frac{1}{2}(c + \delta + \theta)$$

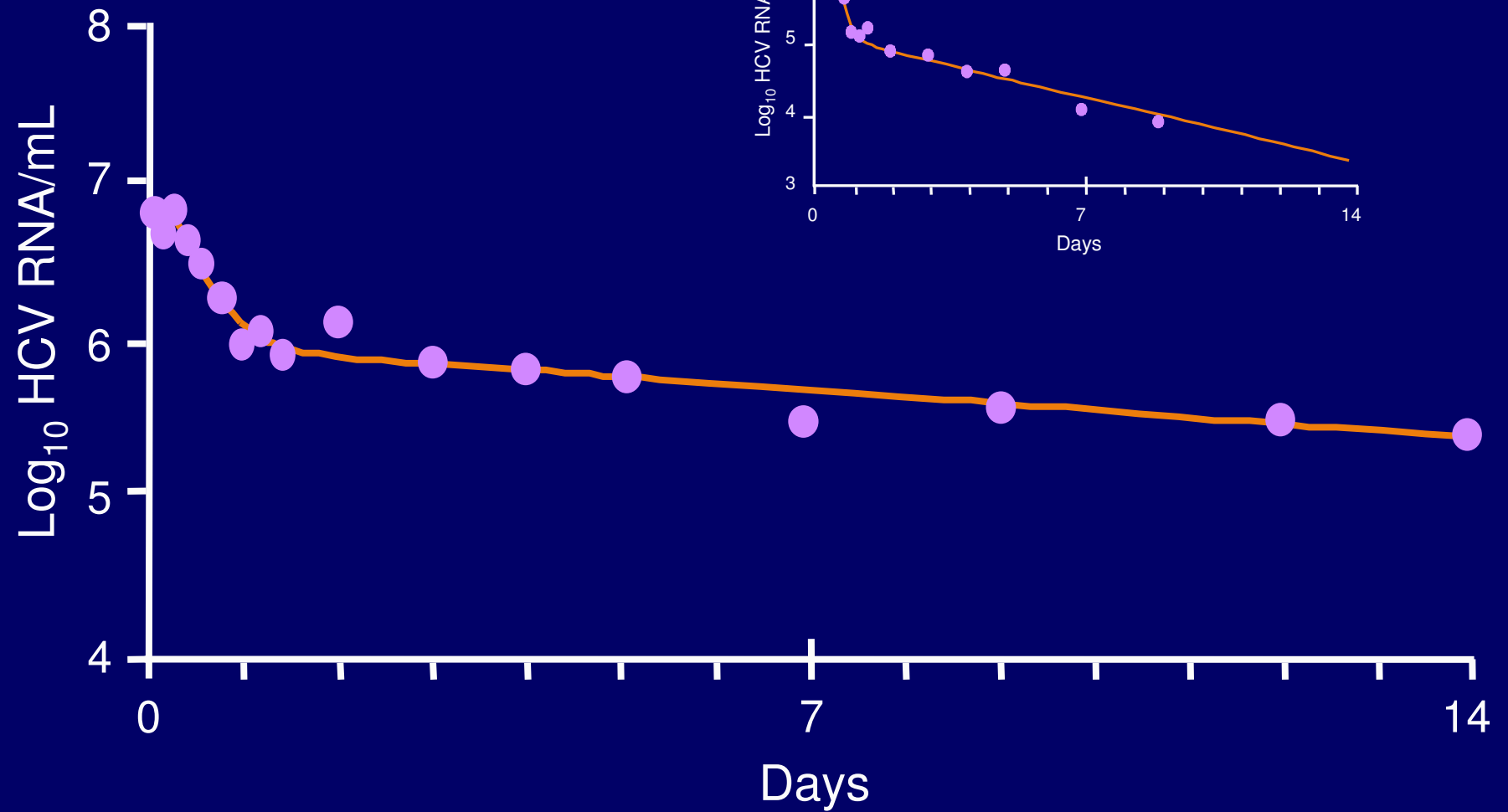
$$\lambda_2 = \frac{1}{2}(c + \delta - \theta)$$

$$\theta = \sqrt{(c - \delta)^2 + 4(1 - \varepsilon)c\delta}$$

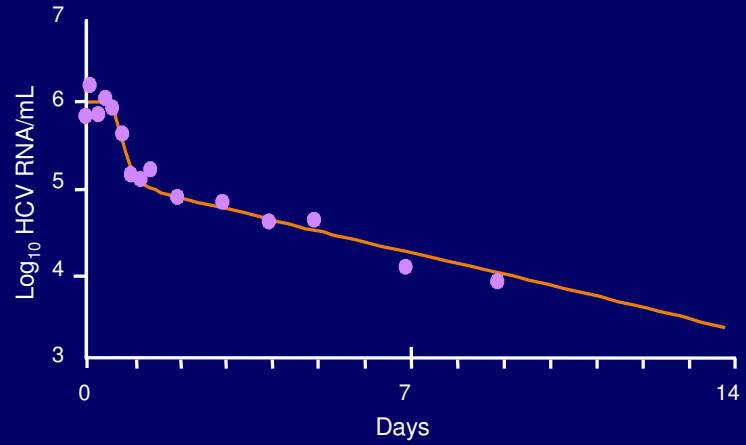
t_0 = delay between treatment commencement and onset of effect

- 1 When $c \gg \delta$, $\lambda_1 \approx c$ and $\lambda_2 \approx \varepsilon\delta$

10MU



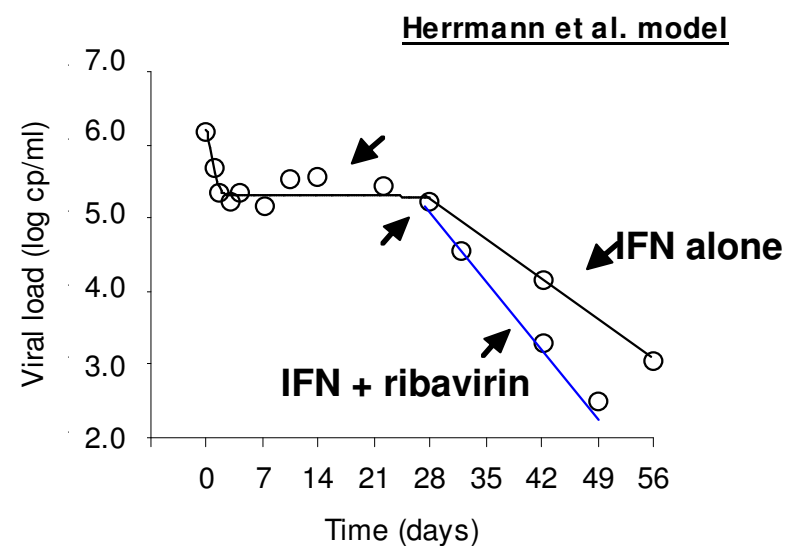
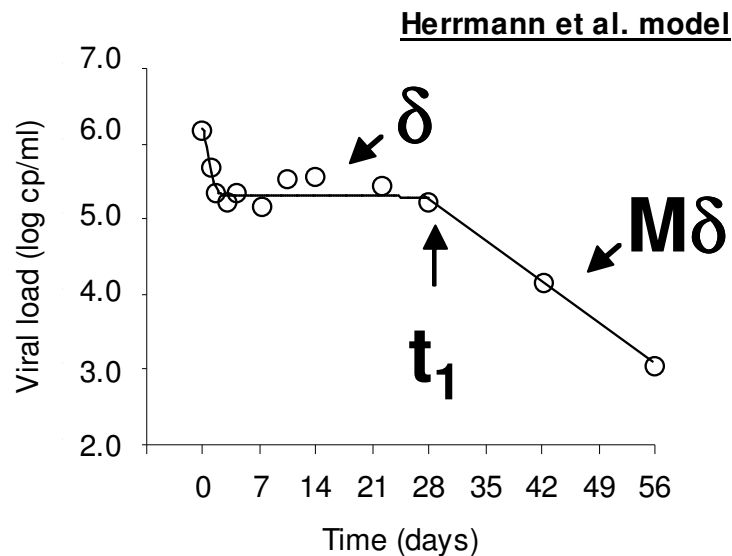
15MU



Viral Kinetics of HCV Genotype 1

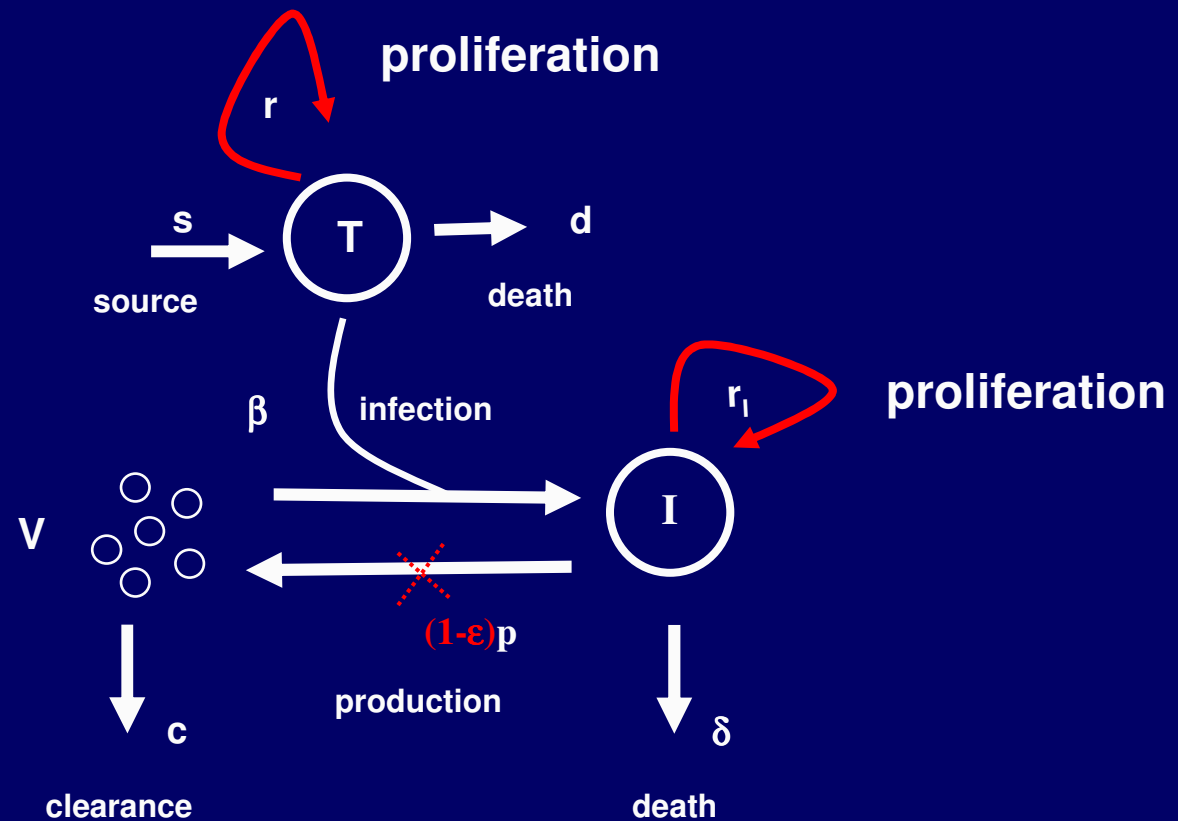
	Drug Efficacy	Second Phase Decay Constant, δ (1/d)	Half-life of Infected Cells (Days)
5MU	$81 \pm 4\%$	0.09 ± 0.14	2.2–69.3
10MU	$95 \pm 4\%$	0.10 ± 0.05	4.3–17.3
15MU	$96 \pm 4\%$	0.24 ± 0.15	1.7–6.3

Triphasic Decay



Herrmann et al., Hepatol. 37: 1351 (2003) suggest the pretreatment infected cell loss rate δ is increased to a treatment-enhanced infected cell loss rate $M\delta$ at a time t_1 . RBV increases M .

Extended Model: Proliferation



Dahari et al., Hepatology 2007; JTB 2007

Model with proliferation

$$\frac{dT}{dt} = s + rT \left(1 - \frac{T + I}{T_{max}} \right) - dT - (1 - \eta)\beta VT,$$

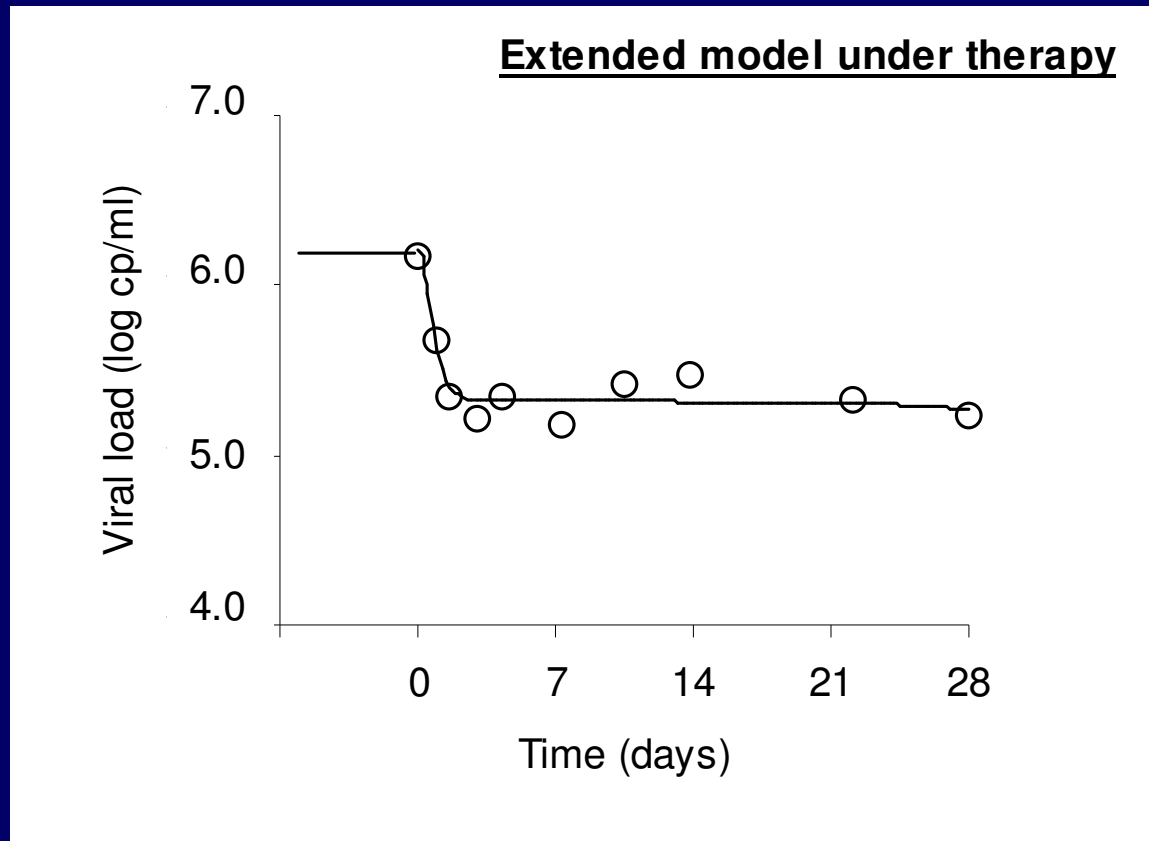
$$\frac{dI}{dt} = (1 - \eta)\beta VT + rI \left(1 - \frac{T + I}{T_{max}} \right) - \delta I,$$

$$\frac{dV}{dt} = (1 - \varepsilon_p)pI - cV,$$

Critical Drug Efficacy

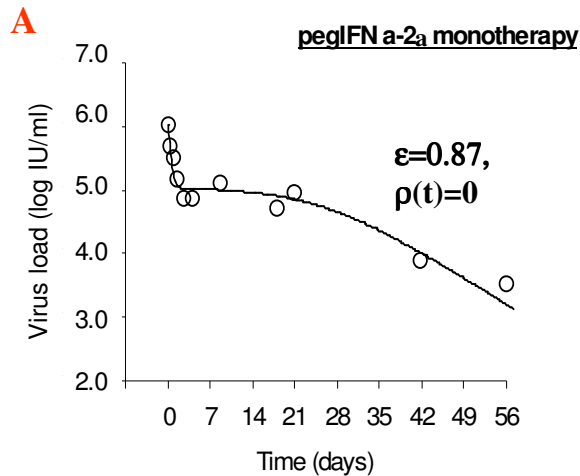
- 1 Drug effectiveness ε :
 - $1-\varepsilon = (1-\varepsilon_p)(1-\eta)$.
- 1 There exists a drug effectiveness, called the critical effectiveness, ε_c , at which the infected steady state amount of virus goes to zero.
- 1 Thus, with $\varepsilon > \varepsilon_c$ model predicts elimination of virus.

$\varepsilon < \varepsilon_c$ Flat 2nd phase

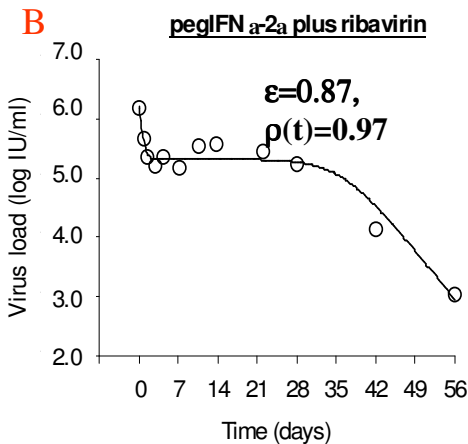


Can have flat 2nd phase with $\delta > 0$ since infected cells replaced by replication and new infection

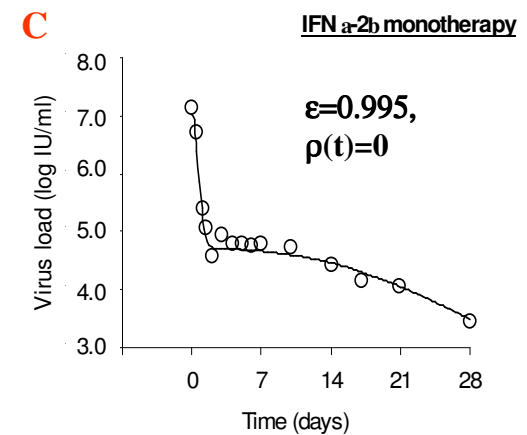
Extended model: Fits to data



Herrmann et al.
(Hepatology 2003)



Herrmann et al.
(Hepatology 2003)

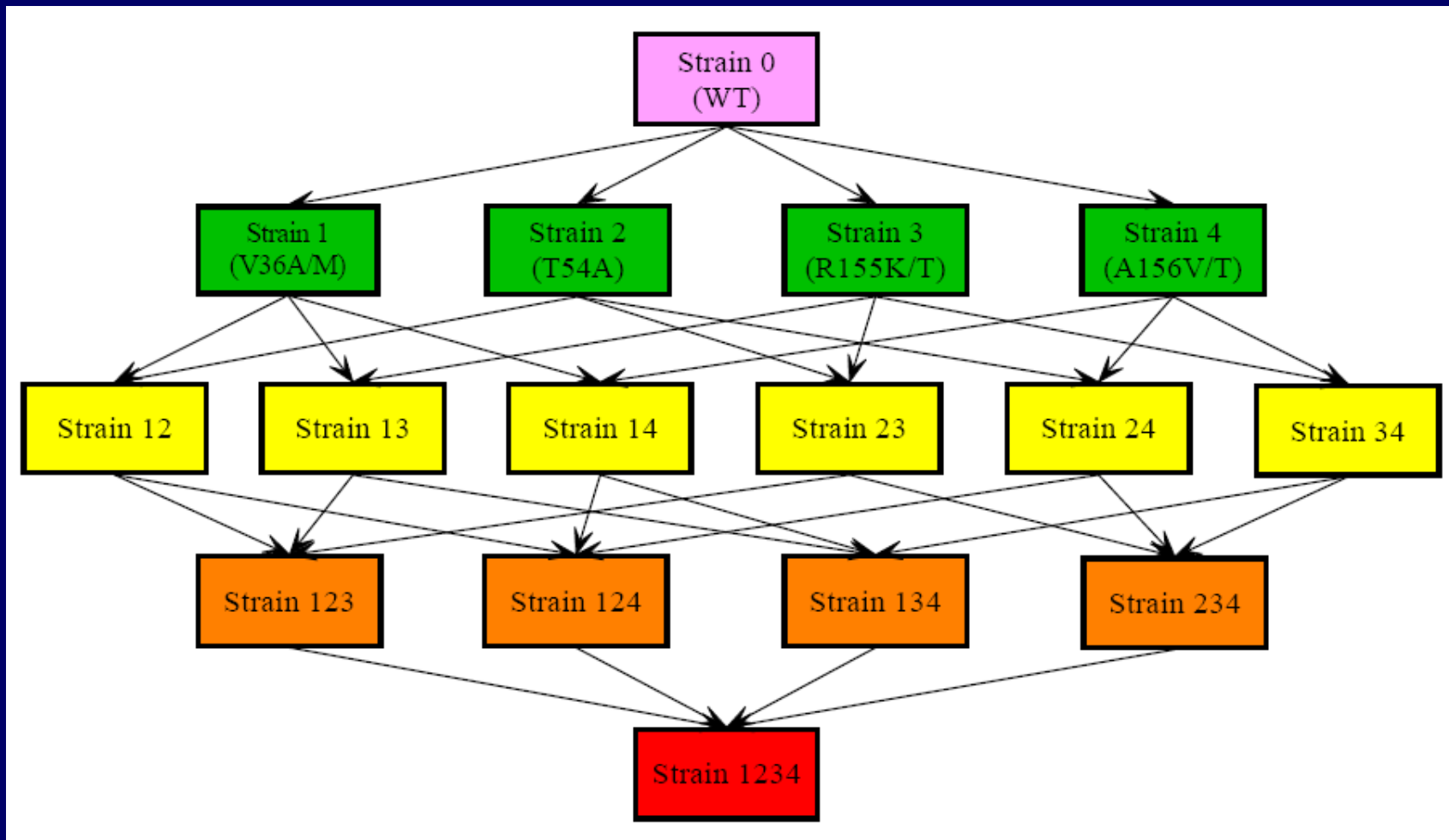


Bekkering et al.
(BMC Gastro 2001)

We fit the extended model to data from patients treated with pegylated interferon α -2a alone (**A**) or in combination with ribavirin (**B**), and with daily therapy with interferon α -2b alone (**C**).

Can explain triphasic response and enhancement of final phase slope without invoking immunomodulation

Mutations between multiple strains



Conclusions

- 1 Mathematics can have a large impact in medicine. I have given you two examples (HIV and HCV). Within the infectious disease community this type of work is being accepted and valued. There are enormous opportunities in this area, not only in infectious disease, but also in cancer, metabolic diseases, immunology,.....

Collaborators (HIV)

- 1 John Pearson, LANL
- 1 Paul Krapivsky, Boston Univ

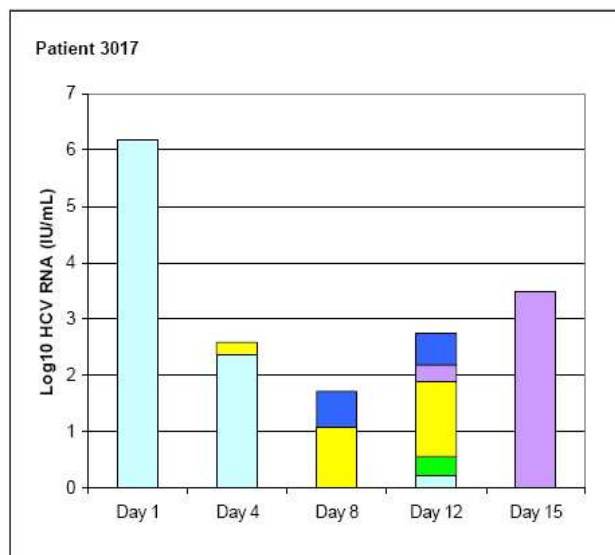
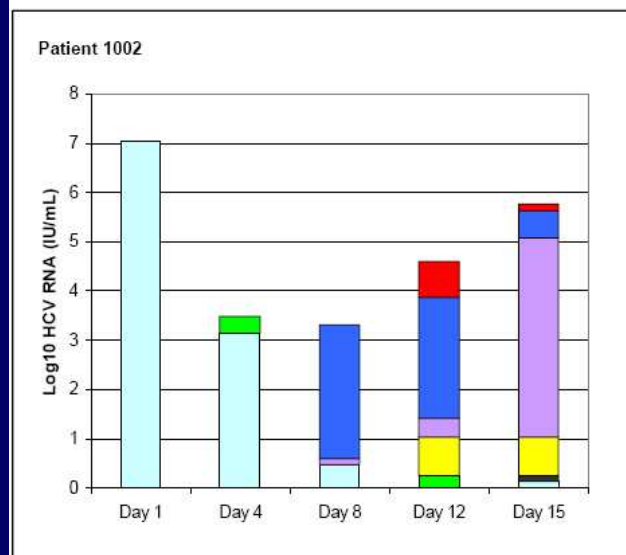
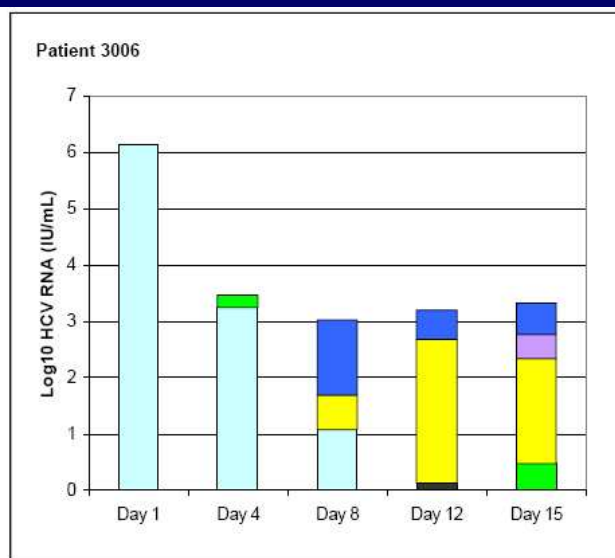
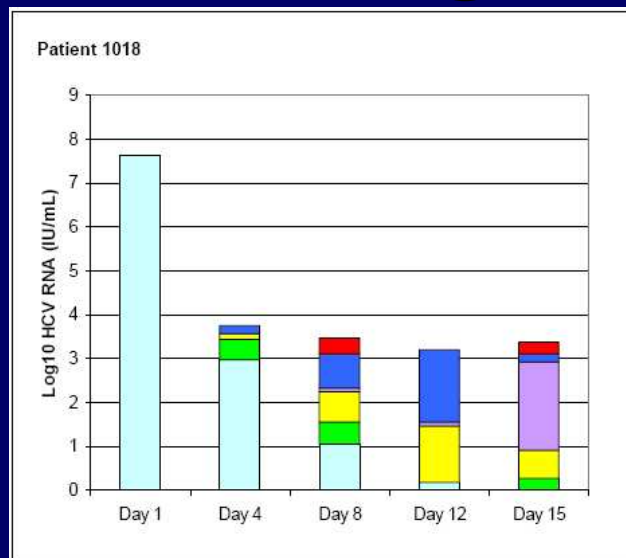
Collaborators (HCV)

- 1 Avidan Neumann, Bar-Ilan Univ
- 1 Harel Dahari, Libin Rong, Emi Shudo
and Ruy M. Ribeiro, Los Alamos
- 1 Tim Reluga, Penn State

A new drug against HCV

- 1 Telaprevir (Vertex Pharmaceuticals):
a new HCV protease inhibitor
- 1 Demonstrates substantial antiviral activity
even as monotherapy
- 1 Drug resistant variants are 5%-20% of the
total virus population **as early as day 3** after
treatment initiation (Kieffer, et al. Hepatology,
2007)
- 1 Such rapid appearance of drug resistance **has
not been seen** with monotherapy for either
HIV or hepatitis B virus (HBV) infection

Rapid emergence of resistance



■ Wild-type
 ■ T54A
 ■ V36A/M
 ■ R155K/T
 ■ 36/155
 ■ A156V/T
 ■ 36/156

Two-strain model

$$\frac{dT(t)}{dt} = s - dT - \beta_s V_s T - \beta_r V_r T$$

$$\frac{dI_s(t)}{dt} = \beta_s V_s T - \delta I_s$$

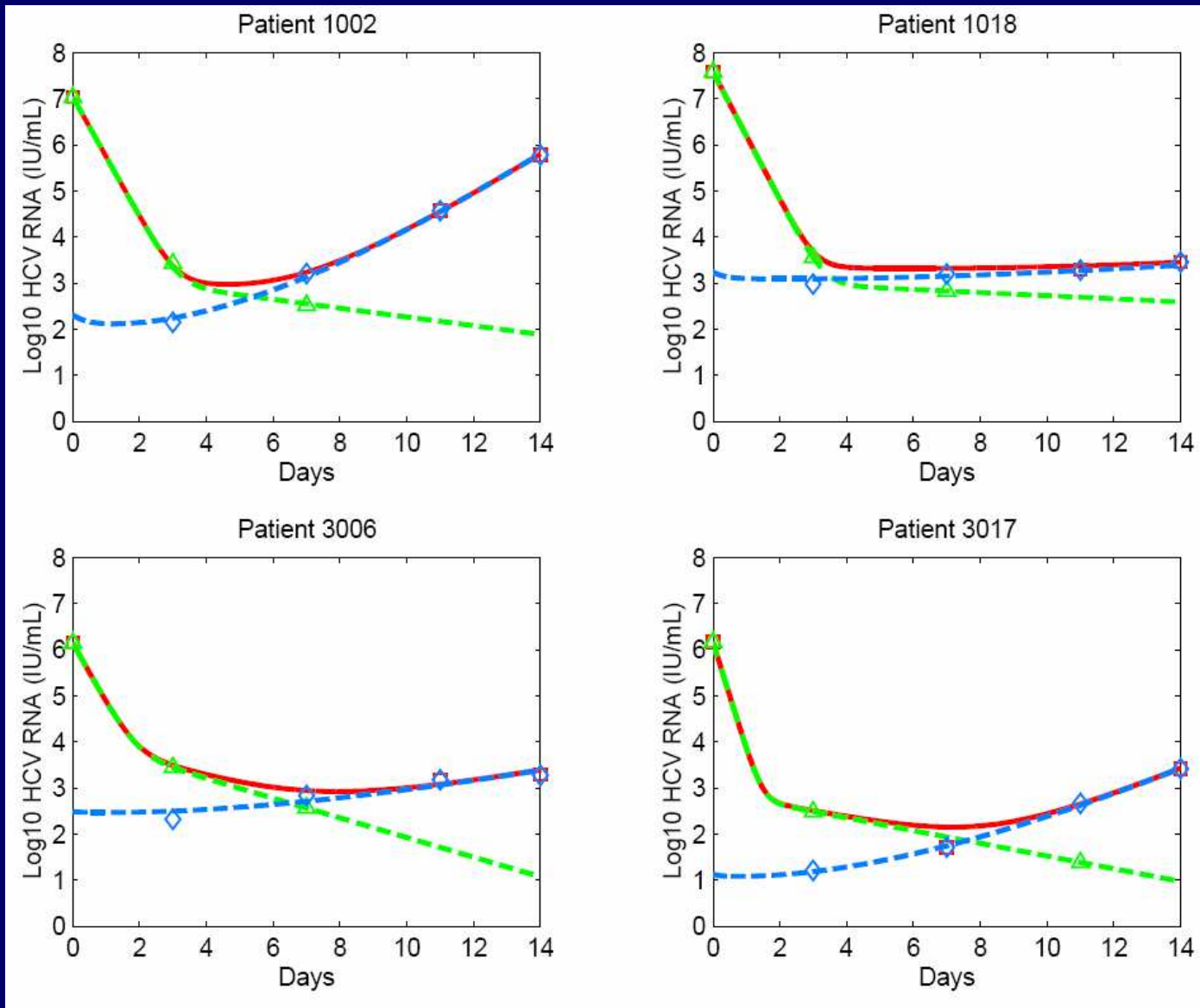
$$\frac{dI_r(t)}{dt} = \beta_r V_r T - \delta I_r$$

$$\frac{dV_s(t)}{dt} = (1 - \mu)(1 - \epsilon_s) p_s I_s - c V_s$$

$$\frac{dV_r(t)}{dt} = \mu(1 - \epsilon_s) p_s I_s + (1 - \epsilon_r) p_r I_r - c V_r$$

s=drug sensitive, r=drug resistant

Model fits



Parameter estimates

Patients	δ (day ⁻¹)	μ (10 ⁻⁶)	ε_s	ε_r	β_s (10 ⁻⁸ mL day ⁻¹ virion ⁻¹)	p_s (virions cell ⁻¹ day ⁻¹)	f_p	c (day ⁻¹)
1018	0.08	12.75	0.99997	0.001	0.77	17.1	0.72	3.2
3006	0.50	3.03	0.99431	0.023	7.06	5.6	0.98	3.0
1002	0.22	6.70	0.99988	0.015	13.64	10.4	0.64	3.0
3017	0.32	0.88	0.99952	0.013	25.85	4.2	0.90	5.4
Average ± SD	0.28 ± 0.18	5.84 ± 5.20	0.99842 ± 0.0027	0.013 ± 0.009	11.83 ± 10.72	9.33 ± 5.82	0.81 ± 0.16	3.65 ± 1.17



**Drug efficacy of telaprevir
against resistant virus**

Percentage people HIV infected

