

CONTROL OF INFLUENZA A VIRUS INFECTION BY VARYING DEATH RATES OF INFECTED CELLS: ANALYSIS OF A SPATIAL MODEL

Murray E. Alexander

*NRC Institute for Biodiagnostics, and Adjunct Professor (Physics), University of
Winnipeg*

Beni M. Sahai

*Cadham Provincial Laboratory; and Adjunct Professor (Biology), University of
Winnipeg*

Acknowledgements:

Natural Sciences & Engineering Research Council of Canada

Canadian Institutes of Health Research

Background

- Influenza A viral infection of respiratory epithelium triggers innate immune response
 - secretion from infected epithelial cells of type-1 interferons, $\text{INF-}\alpha/\beta$
 - inflammatory & chemotactic cytokines from alveolar macrophages and (mobile) neutrophils
 - dendritic cells (following phagocytosis of newly-synthesized virus particles)
- Leads to
 - activation of **NK (natural killer) cells**
 - viral antigen-bearing **macrophages** and **dendritic cells**
 - Macrophages, DCs lead, via clonal expansion, to **influenza A-specific cytotoxic T lymphocytes (CTLs)**.
- Activated NK cells kill newly-infected epithelial cells
- Anti-influenza CTLs destroy virus-producing epithelial cells

Level 2
model

Innate immunity

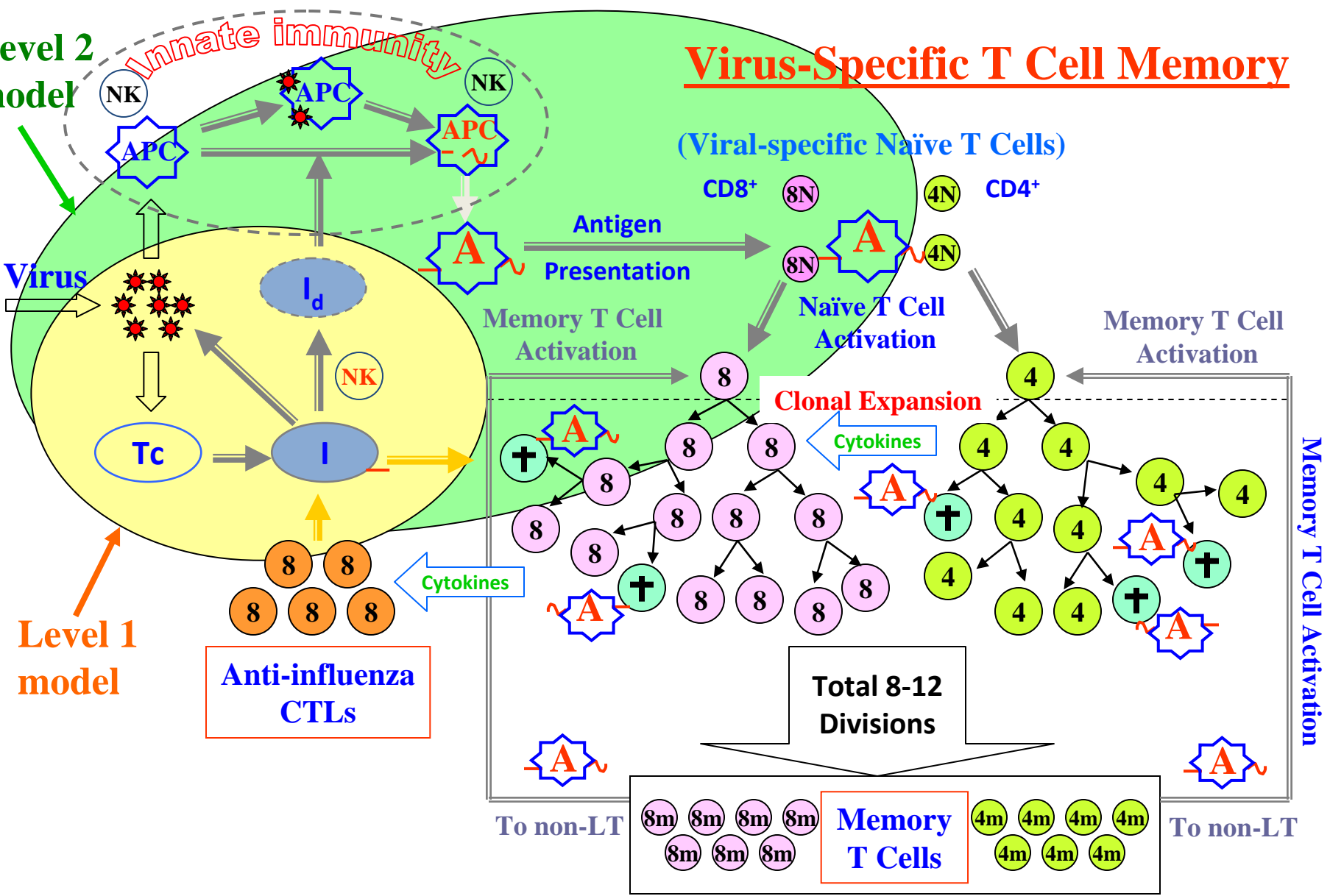
Virus-Specific T Cell Memory

(Viral-specific Naïve T Cells)

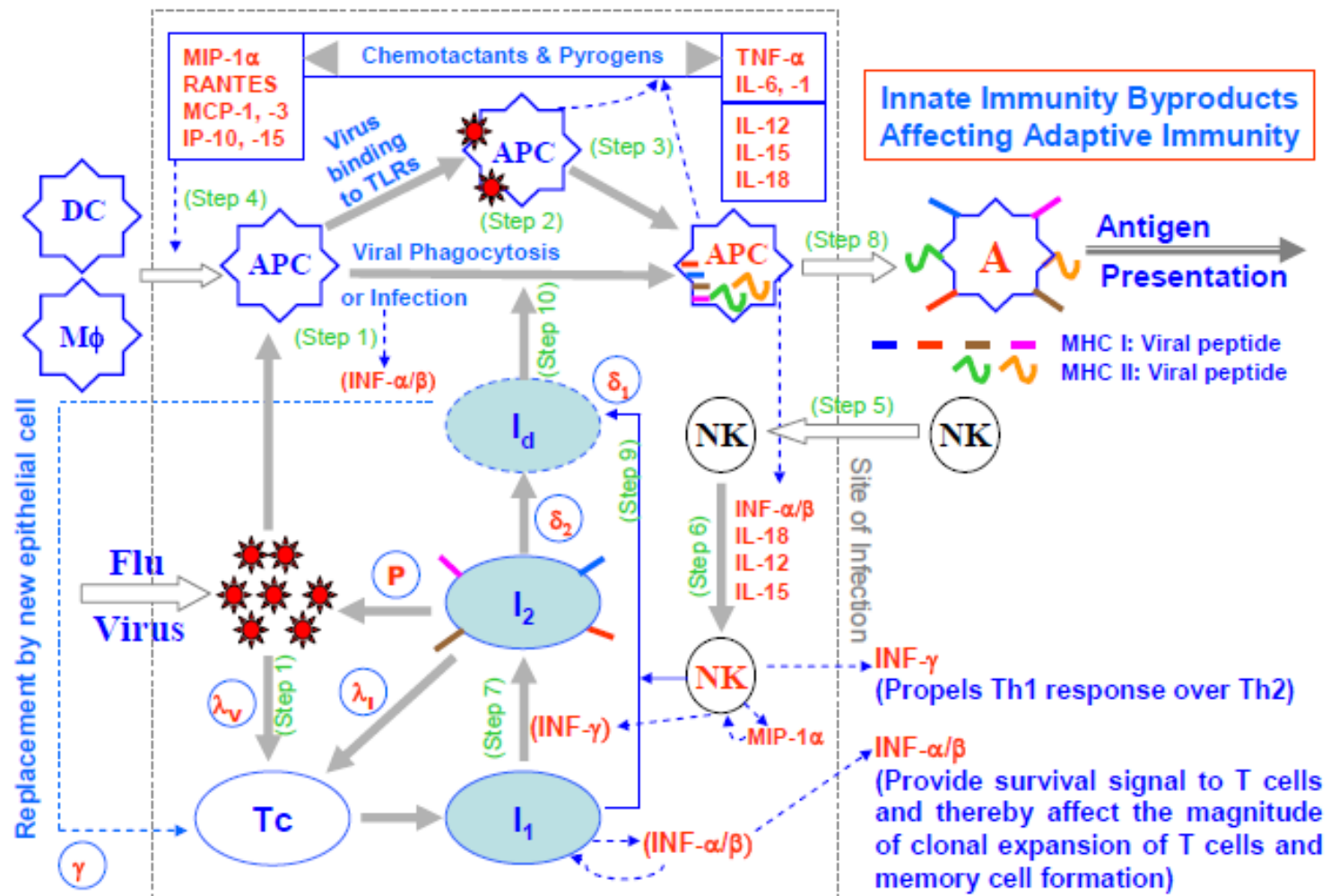
Virus

Level 1
model

Anti-influenza
CTLs



Innate Immunity to Influenza Infection



Level 1 model

$$\frac{dT_{R,x}}{dt} = -\lambda_V T_{R,x} V_x - \lambda_I T_{R,x} \sum_{e \in N_1} I_{2,x'} + \frac{\gamma}{K} \sum_{e \in N_1} I_{2,x+e} \quad \text{Regeneration} \quad (1)$$

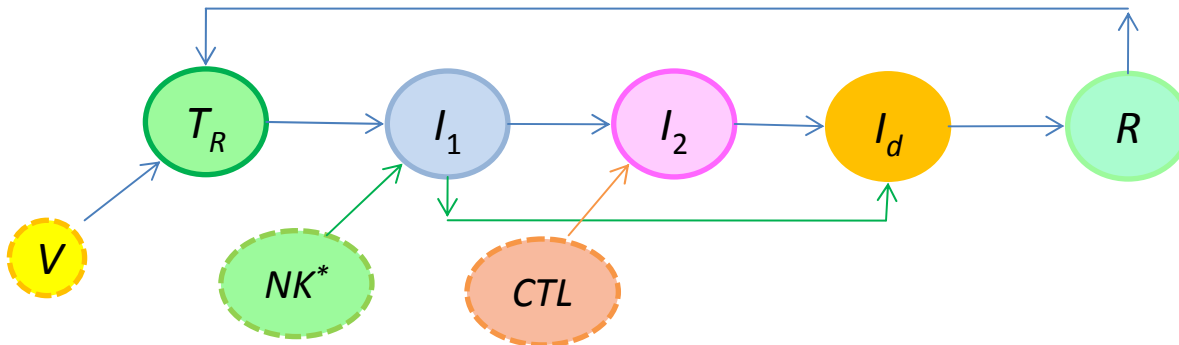
$$\frac{dI_{1,x}}{dt} = \lambda_V T_{R,x} V_x + \lambda_I T_{R,x} \sum_{e \in N_1} I_{2,x+e} - kI_{1,x} - \delta_2 [NK^*]_x I_{1,x} \quad (2)$$

$$\frac{dI_{2,x}}{dt} = kI_{1,x} - \delta_1 I_{2,x} - \delta_3 [CTL] I_{2,x} \quad (3)$$

$$\frac{dV_x}{dt} = pI_{2,x} - cV_x - \beta_{APC} [APC]_x V_x - \beta_{NEU} [NEU]_x V_x - \lambda_V (I_{1,x} + I_{2,x} + T_{R,x}) V_x + D_V \nabla^2 V_x \quad (4)$$

$$\frac{dI_{d,x}}{dt} = \delta_2 [NK^*]_x I_{1,x} + \delta_1 I_{2,x} - (\beta_{d,1} [APC]_x + \beta_{d,2} [NEU]_x) I_{d,x} + \delta_3 [CTL] I_{2,x} \quad (5)$$

$$\frac{dR_x}{dt} = (\beta_{d,1} [APC]_x + \beta_{d,2} [NEU]_x) I_{d,x} - \gamma R_x \quad \text{Phagocytosis} \quad (6)$$



Epithelial cell repair & regeneration

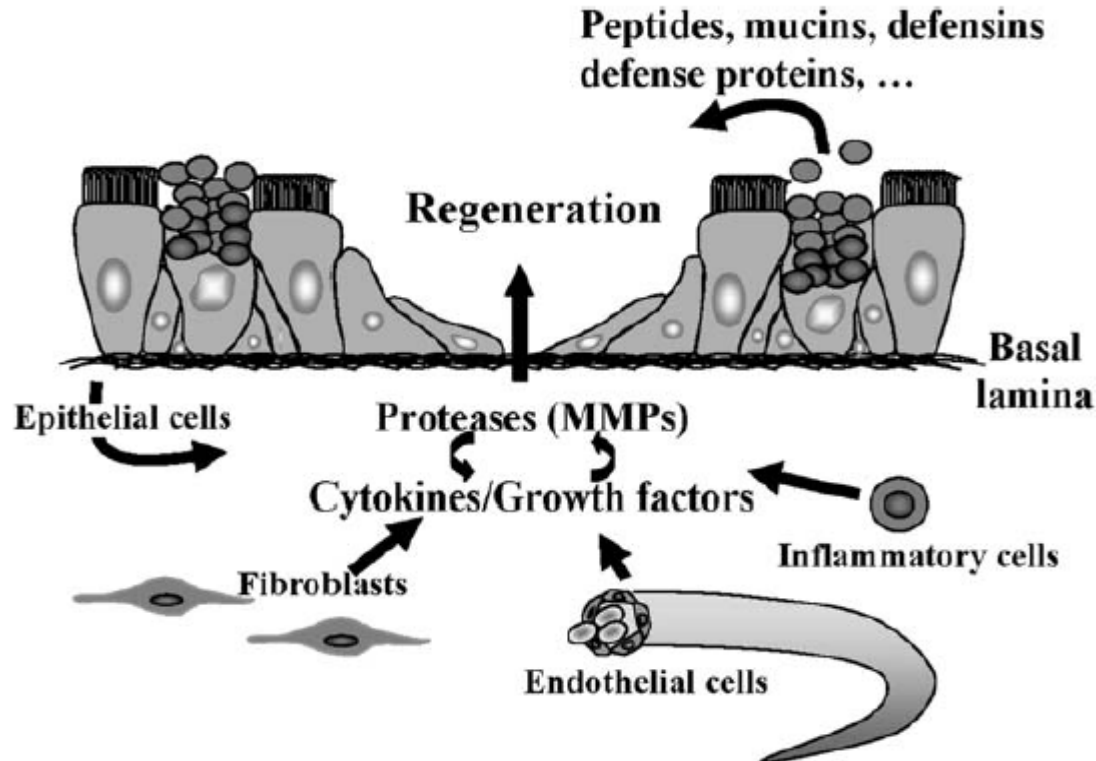


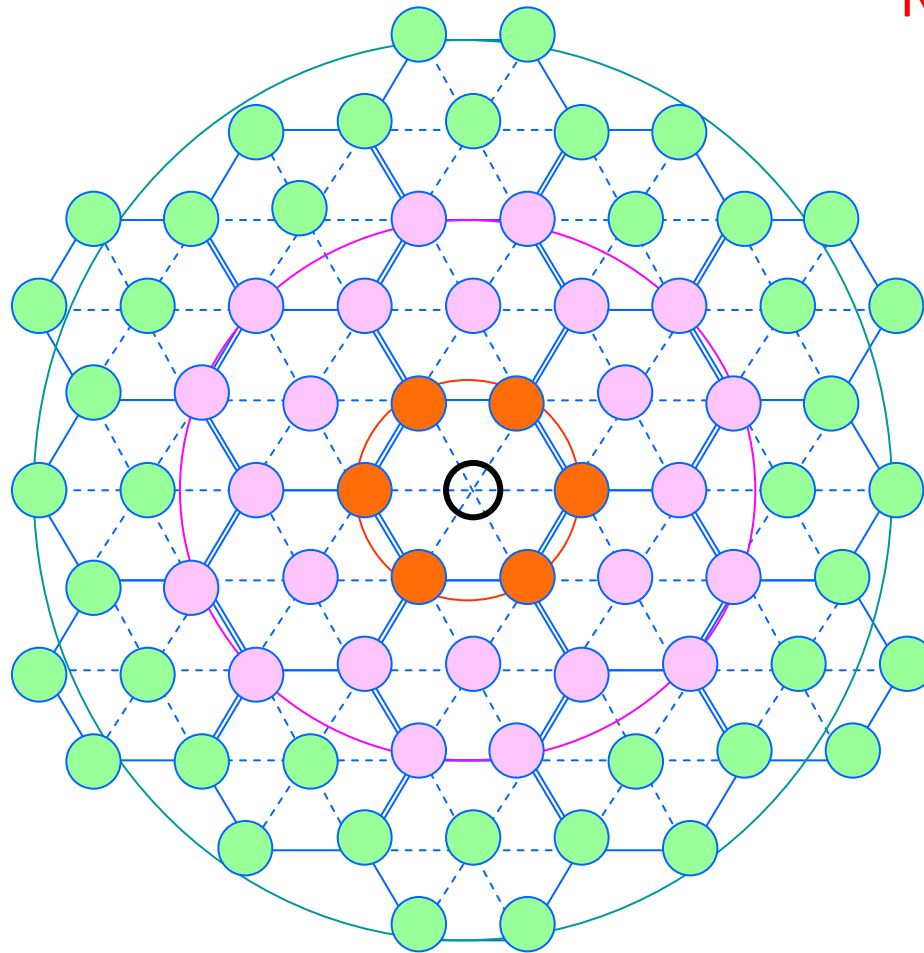
Figure 1. Cellular and molecular factors involved in the repair and regeneration of the airway epithelium. These factors, which closely interact during the different steps of airway epithelial regeneration after injury, are modulated by the components of the extracellular matrix; the matrix metalloproteinases (MMPs), cytokines, and growth factors released by the epithelial cells; and by the mesenchymal cells (fibroblasts, inflammatory cells, and chondrocytes).

From: *Proc Am Thorac Soc* 3(2006):726–733

HEXAGONAL LATTICE OF EPITHELIAL CELLS

Neighbourhoods:

N_1 N_2 N_3 ...



Linear Stability of Infection-Free Equilibrium (IFE)

$$\Delta_2 = \delta_2[NK *]; \quad \Delta_3 = \delta_1 + \delta_3[CTL];$$

- Define $\rho_1 = \beta_{APC}[APC] + \beta_{NEU}[NEU];$
 $\rho_2 = \beta_{d,1}[APC] + \beta_{d,2}[NEU];$
 $L_V = \lambda_V \bar{T}_R; \quad L_I = \lambda_I \bar{T}; \quad \Lambda = c + \rho_1 + L_V$
(constants)

- In infection-free equilibrium, the $\{T_{R,x}\}$ constitute a **uniform hexagonal lattice**.
- Introduce a small perturbation

$$\delta \mathbf{X}_x = (\delta T_{R,x}, \delta I_{1,x}, \delta I_{2,x}, \delta V_x, \delta I_{d,x}, \delta R_x)$$

about I.F.E. $\mathbf{X}_0 = (\bar{T}_R, 0, 0, 0, 0, 0)$ and linearize

LINEARIZATION ABOUT I.F.E.

$$\delta \dot{T}_{R,\mathbf{x}} = -L_V \delta V_{\mathbf{x}} - L_I \sum_{\mathbf{e} \in N_1} \delta I_{2,\mathbf{x}+\mathbf{e}} + \frac{\gamma}{K} \sum_{\mathbf{e} \in N_1} \delta R_{\mathbf{x}+\mathbf{e}}$$

$$\delta \dot{I}_{1,\mathbf{x}} = L_V \delta V_{\mathbf{x}} + L_I \sum_{\mathbf{e} \in N_1} \delta I_{2,\mathbf{x}+\mathbf{e}} - (k + \Delta_2) \delta I_{1,\mathbf{x}}$$

$$\delta \dot{I}_{2,\mathbf{x}} = k \delta I_{1,\mathbf{x}} - \Delta_3 \delta I_{2,\mathbf{x}}$$

$$\delta \dot{V}_{\mathbf{x}} = p \delta I_{2,\mathbf{x}} - (\Lambda + D_V) \delta V_{\mathbf{x}} + \frac{D_V}{K} \sum_{\mathbf{e} \in N_1} \delta V_{\mathbf{x}+\mathbf{e}}$$

$$\delta \dot{I}_{d,\mathbf{x}} = \Delta_2 \delta I_{1,\mathbf{x}} + \Delta_3 \delta I_{2,\mathbf{x}} - \rho_2 \delta I_{d,\mathbf{x}}$$

$$\delta \dot{R}_{\mathbf{x}} = \rho_2 \delta I_{d,\mathbf{x}} - \gamma \delta R_{\mathbf{x}}$$

where $K = \#\{N_1\} = 6$

Let $\delta\mathbf{X} = \delta\mathbf{X}_c \exp(\lambda t + i\mathbf{k} \cdot \mathbf{x}) \quad \longrightarrow$

$$\lambda \delta\mathbf{X}_c = J_{\mathbf{k}} \delta\mathbf{X}_c$$

$$J_{\mathbf{k}} = \begin{bmatrix} 0 & 0 & -L_I \varphi(\mathbf{k}) & -L_V & 0 & \gamma \frac{\varphi(\mathbf{k})}{K} \\ 0 & -(k + \Delta_2) & L_I \varphi(\mathbf{k}) & L_V & 0 & 0 \\ 0 & k & -\Delta_3 & 0 & 0 & 0 \\ 0 & 0 & p & -\Lambda - D_V \left(1 - \frac{\varphi(\mathbf{k})}{K}\right) & 0 & 0 \\ 0 & \Delta_2 & \Delta_3 & 0 & -\rho_2 & 0 \\ 0 & 0 & 0 & 0 & \rho_2 & -\gamma \end{bmatrix}$$

where
$$\varphi(\mathbf{k}) = 2 \left(\cos\left(\frac{2\pi}{N_1} k_1\right) + \cos\left(\frac{2\pi}{N_2} k_2\right) + \cos\left(2\pi \left(\frac{k_1}{N_1} + \frac{k_2}{N_2}\right)\right) \right)$$

$$\mathbf{k} = 2\pi \left(\frac{k_1}{N_1} \hat{\mathbf{e}}_1^* + \frac{k_2}{N_2} \hat{\mathbf{e}}_2^* \right) \quad \hat{\mathbf{e}}_1^*, \hat{\mathbf{e}}_2^* \longrightarrow \text{Basis vectors of reciprocal lattice}$$

Characteristic equation

$$\lambda(\lambda + \gamma)(\lambda + \rho_2)C(\lambda; \mathbf{k}) = 0$$

$C(\lambda; \mathbf{k})$ is a cubic polynomial in λ :

$$C(\lambda; \mathbf{k}) = \lambda^3 + a_2\lambda^2 + a_1\lambda + a_0$$

where

$$a_0 = (\Delta_3(k + \Delta_2) - kL_I\varphi) \left(\Lambda + D_V \left(1 - \frac{\varphi(\mathbf{k})}{K} \right) \right) - kpL_V$$

$$a_1 = \Delta_3 \left(k + \Delta_2 + \Lambda + D_V \left(1 - \frac{\varphi(\mathbf{k})}{K} \right) \right) + \left(\Lambda + D_V \left(1 - \frac{\varphi(\mathbf{k})}{K} \right) \right) (k + \Delta_2) - kL_I\varphi$$

$$a_2 = \Delta_3 + k + \Delta_2 + \Lambda + D_V \left(1 - \frac{\varphi(\mathbf{k})}{K} \right)$$

$$\begin{aligned} \Delta_2 &= \delta_2[NK *]; & \Delta_3 &= \delta_1 + \delta_3[CTL]; & L_V &= \lambda_V \bar{T}_R; & L_I &= \lambda_I \bar{T}; & \Lambda &= c + \rho_1 + L_V \\ \rho_1 &= \beta_{APC}[APC] + \beta_{NEU}[NEU]; & \rho_2 &= \beta_{d,1}[APC] + \beta_{d,2}[NEU]; \end{aligned}$$

Routh-Hurwitz criteria

$$(i) \quad a_2 > 0; \quad (ii) \quad a_1 a_2 - a_0 > 0; \quad (iii) \quad a_0 (a_1 a_2 - a_0) > 0$$

Conclusion

Since $\Lambda + D_V \left(1 - \frac{\varphi}{K} \right) > 0 \quad \forall \mathbf{k}$ it follows that,

*for sufficiently large Δ_3 (representing killing of cells by CTLs), the Routh-Hurwitz criteria are satisfied and **the system is stable against spread of infection, independent of the diffusion rate D_V .***

Continuous form of diffusion term

- Replace discrete approximation: $\nabla^2 V_{\mathbf{x}} \approx \frac{1}{K} \sum_{\mathbf{e} \in N_1} V_{\mathbf{x}+\mathbf{e}} - V_{\mathbf{x}}$

by its continuous form (also an approximation to diffusion!)

In hexagonal frame:

$$\mathbf{r} = \hat{\mathbf{e}}_1 \xi + \hat{\mathbf{e}}_2 \eta; \quad \nabla = \hat{\mathbf{e}}_1^* \frac{\partial}{\partial \xi} + \hat{\mathbf{e}}_2^* \frac{\partial}{\partial \eta}$$

and $\nabla^2 \exp(\lambda t + i\mathbf{k} \cdot \mathbf{x}) = -|\mathbf{k}|^2 \exp(\lambda t + i\mathbf{k} \cdot \mathbf{x})$

where $|\mathbf{k}|^2 = \left| \kappa_1 \hat{\mathbf{e}}_1^* + \kappa_2 \hat{\mathbf{e}}_2^* \right|^2 = \frac{4}{3} (\kappa_1^2 + \kappa_2^2 \pm \kappa_1 \kappa_2)$

$$\kappa_j \equiv 2\pi \frac{k_j}{N_j}, \quad j = 1, 2 \Rightarrow |\mathbf{k}| \leq 2\pi$$

- Linearization about IFE gives rise to the alternative characteristic equation with cubic factor

$$C_c(\lambda; \mathbf{k}) = \lambda^3 + b_2 \lambda^2 + b_1 \lambda + b_0$$

where

$$b_0 = (\Delta_3(k + \Delta_2) - kL_I\varphi)(\Lambda + D_V|\mathbf{k}|^2) - kpL_V$$

$$b_1 = \Delta_3(k + \Delta_2 + \Lambda + D_V|\mathbf{k}|^2) + (\Lambda + D_V|\mathbf{k}|^2)(k + \Delta_2) - kL_I\varphi$$

$$b_2 = \Delta_3 + k + \Delta_2 + \Lambda + D_V|\mathbf{k}|^2$$

where $|\mathbf{k}| \leq 2\pi$

Routh-Hurwitz yields essentially same outcome as before:

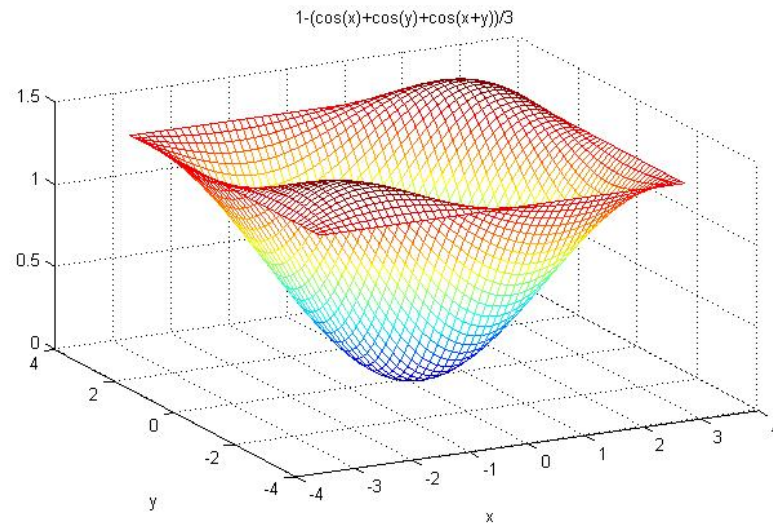
For sufficiently large killing rate (Δ_3) of infected cells by CTLs, the system is stable against spread of infection, independent of the diffusion rate D_V .

However: Critical Δ_3 will be different between 'discrete' and 'continuous' diffusion models

Approximations to $-\nabla^2$

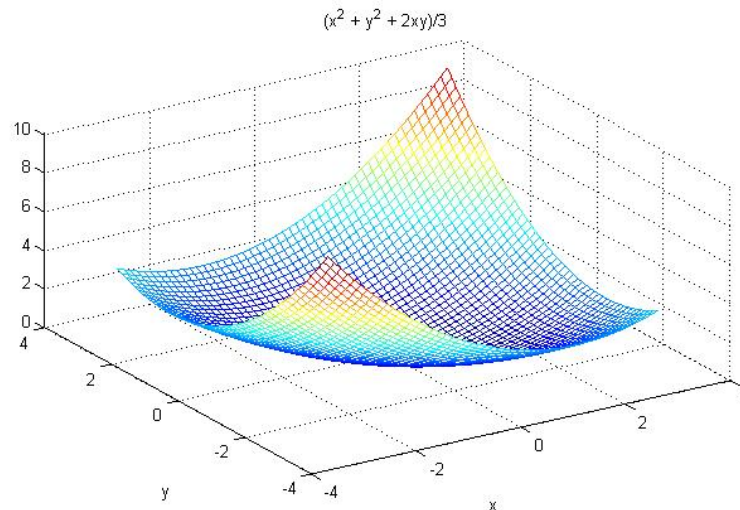
Numerical
simulations

$$1 - \frac{1}{6} \varphi(\mathbf{k})$$



Continuous

$$|\mathbf{k}|^2$$

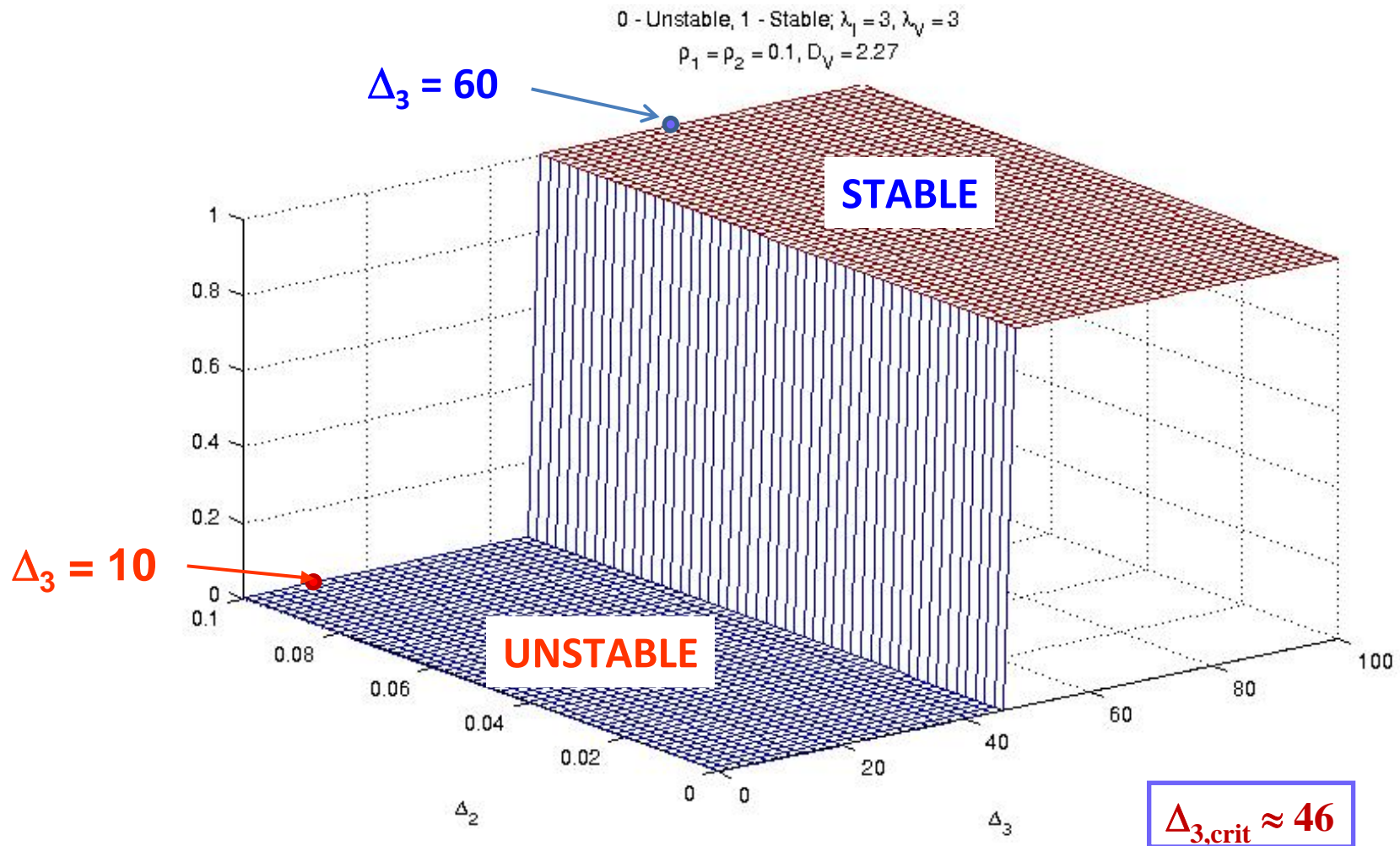


Parameter values

Parameter		Value
λ_I	Infection rate due to contact with I_2 cells	3 d ⁻¹
λ_V	Infection rate due to free virus	3 d ⁻¹
γ	Replacement rate, vacant epithelial cell sites (R)	3 d ⁻¹
k	Rate of $I_1 \rightarrow I_2$ (infected \rightarrow infectious) transition	4 d ⁻¹
δ_1	Death rate of I_2 by viral infection alone	0.1 d ⁻¹
δ_2	Death rate of I_1 by NK*	0.1 d ⁻¹
δ_3	Death rate of I_2 due to CTLs	(variable)
$\beta_{APC,1}$	Rate of viral phagocytosis (by APC)	*
β_{NEU}	Rate of viral phagocytosis (by neutrophils)	*
$\beta_{d,1}$	Rate of uptake of dead cells by APC	*
$\beta_{d,2}$	Rate of uptake of dead cells by neutrophils	*
p	Rate of production of free virus by infected cell	50
c	Viral clearance rate	2 d ⁻¹
D_V	Diffusion constant (scaled to cell spacing)	2.27 d ⁻¹

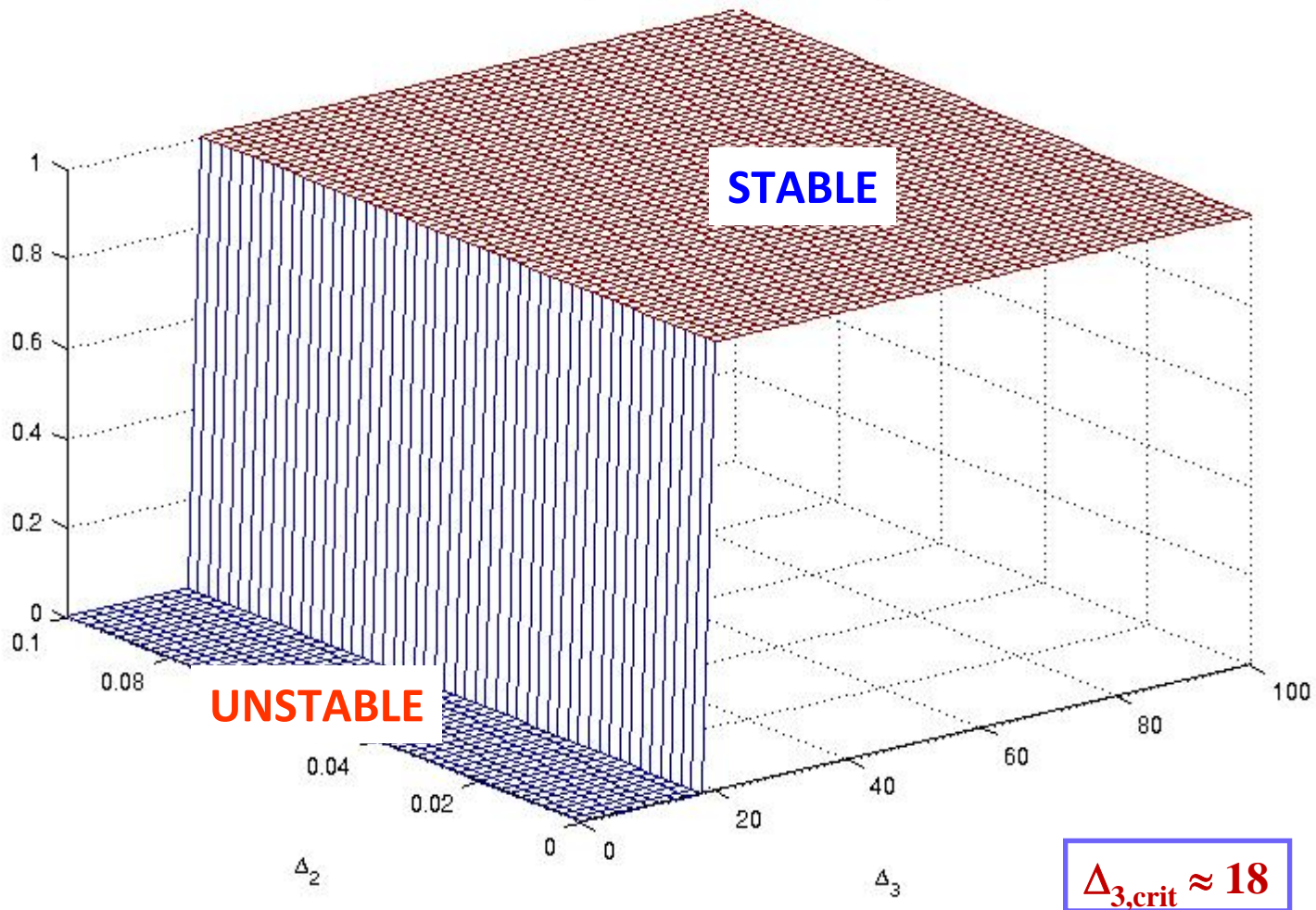
* Level-1 model incorporates these, and [APC], [NEU], into constants ρ_1 , ρ_2 16

Stability (of IFE) conferred by CTLs: **Discrete** diffusion approximation



Stability (of IFE) conferred by CTLs: **Continuous** diffusion approximation

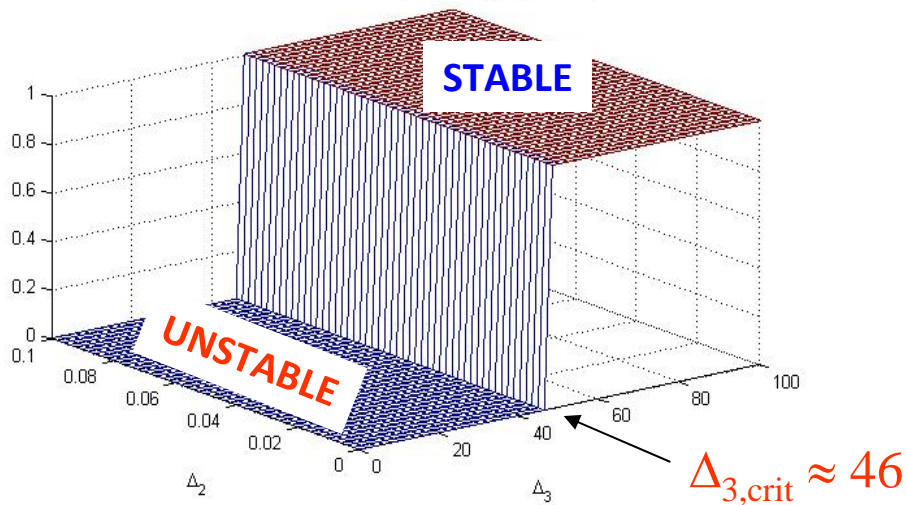
Continuous diffusion: $\lambda_I = 3$, $\lambda_V = 3$, $\rho_1 = \rho_2 = 0.1$, $D_V = 2.27$



Continuous diffusion approximation: Dependence on $|\mathbf{k}|$

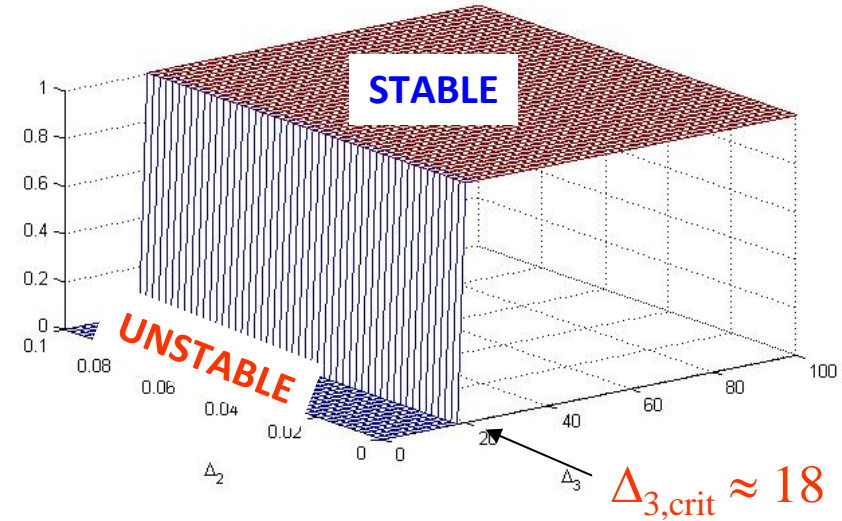
$$|\mathbf{k}| = 0$$

Continuous Diffusion: $|\mathbf{k}| = 0$, $\lambda_1 = \lambda_V = 3$, $\rho_1 = \rho_2 = 0.1$, $D_V = 2.27$



$$|\mathbf{k}| = 2\pi$$

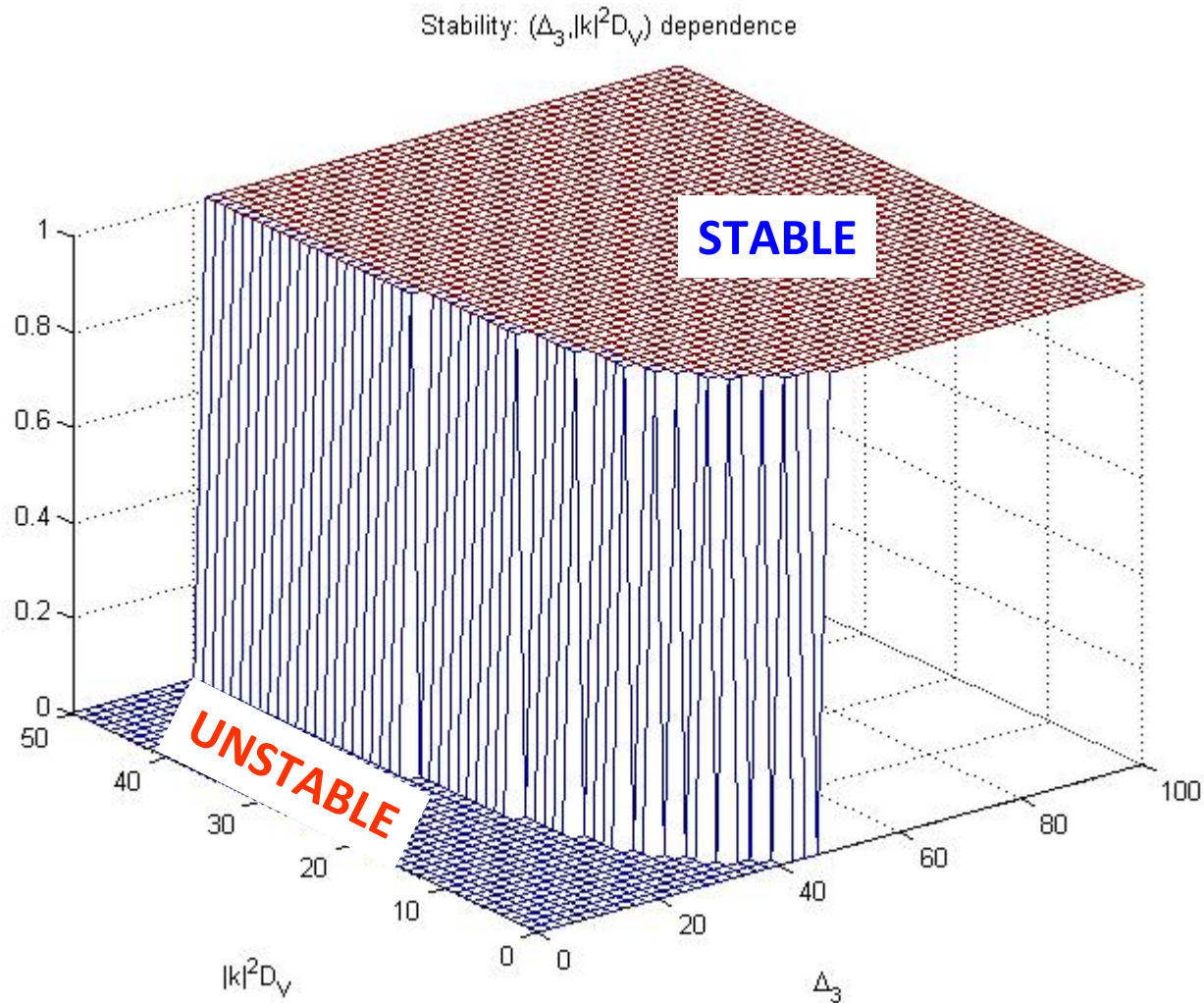
Continuous Diffusion: $|\mathbf{k}| = 2\pi$, $\lambda_1 = \lambda_V = 3$, $\rho_1 = \rho_2 = 0.1$, $D_V = 2.27$



Thus,

- As [CTL] increases, fine-scale ($|\mathbf{k}|$ large) infection regions disappear before coarse-scale ($|\mathbf{k}|$ small)
- As D_V increases, $\Delta_{3,\text{crit}}$ (and hence $[\text{CTL}]_{\text{crit}}$ required to prevent infection) increases

Stability: Dependence on [CTL], diffusion

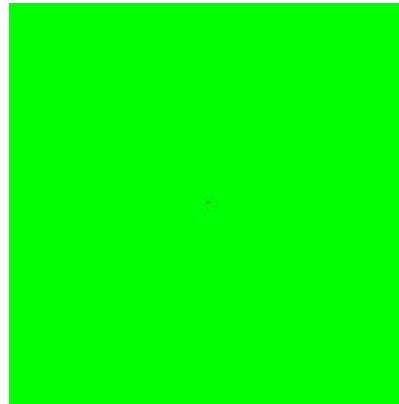


SPATIAL PROGRESSION OF INFECTION

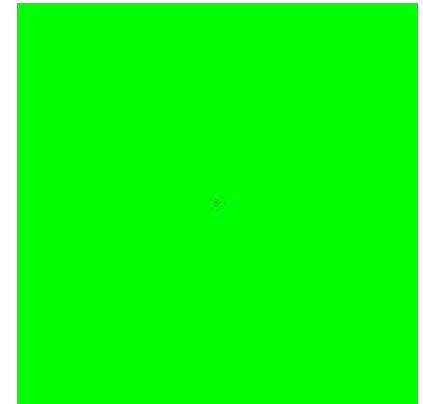
(Seed infection: $V = 1$ at centre)

Stable configuration: $\Delta_3 = 60$

$$\Delta_{3,\text{crit}} \approx 46$$

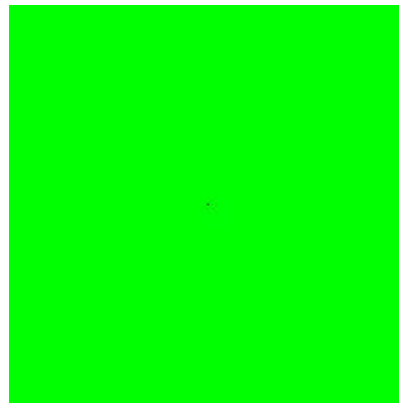


$t = 0$



$t = 10$ days

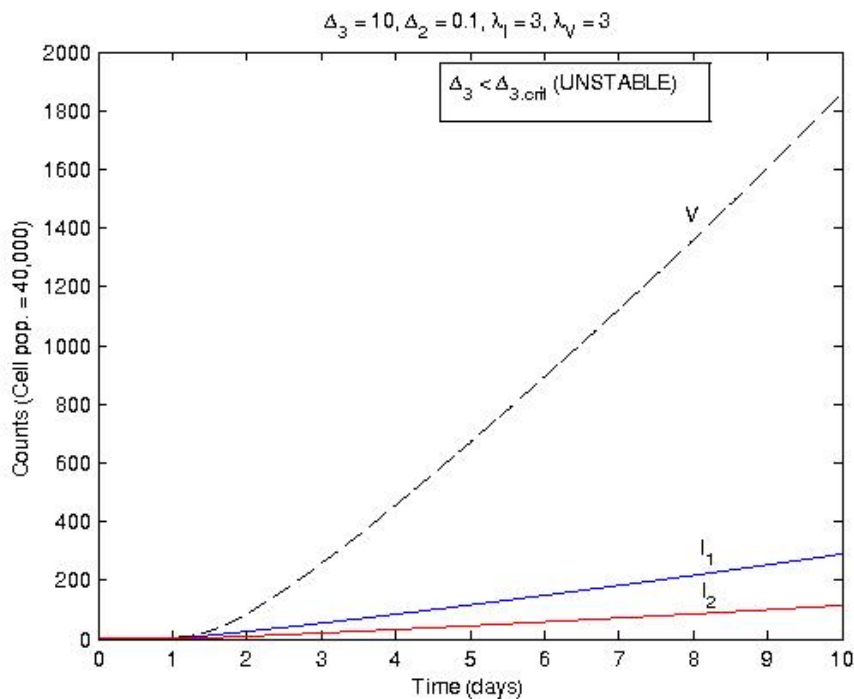
Unstable configuration: $\Delta_3 = 10$



Population counts for virus & infected cells

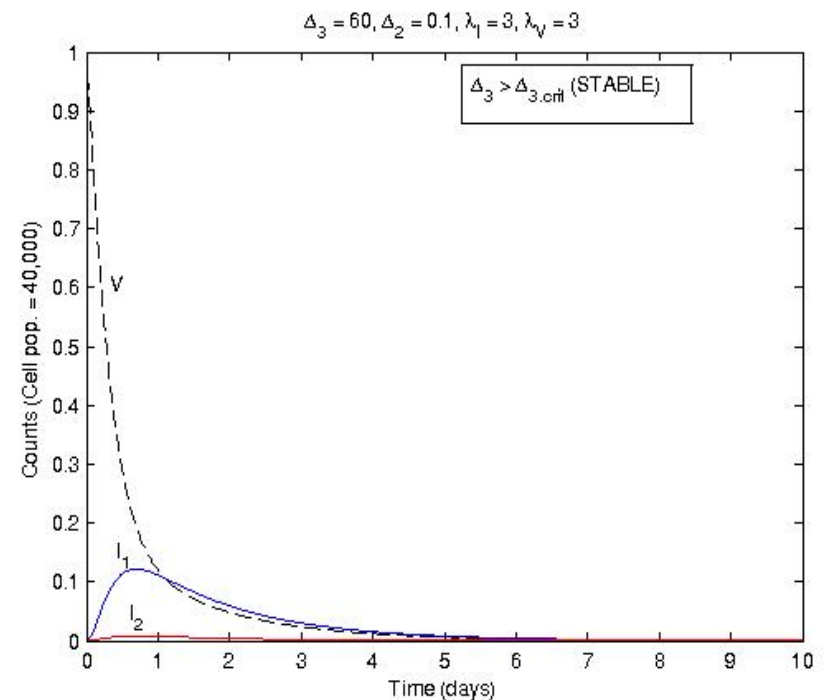
$$\Delta_3 \equiv \delta_1 + \delta_3[\text{CTL}]; \quad \Delta_2 \equiv \delta_2[\text{NK}^*]$$

Unstable ($\Delta_3 = 10 < \Delta_{3,\text{crit}}$)



$\Delta_{3,\text{crit}} \approx 46$

Stable ($\Delta_3 = 60 > \Delta_{3,\text{crit}}$)



Linear stability analysis correctly predicts:

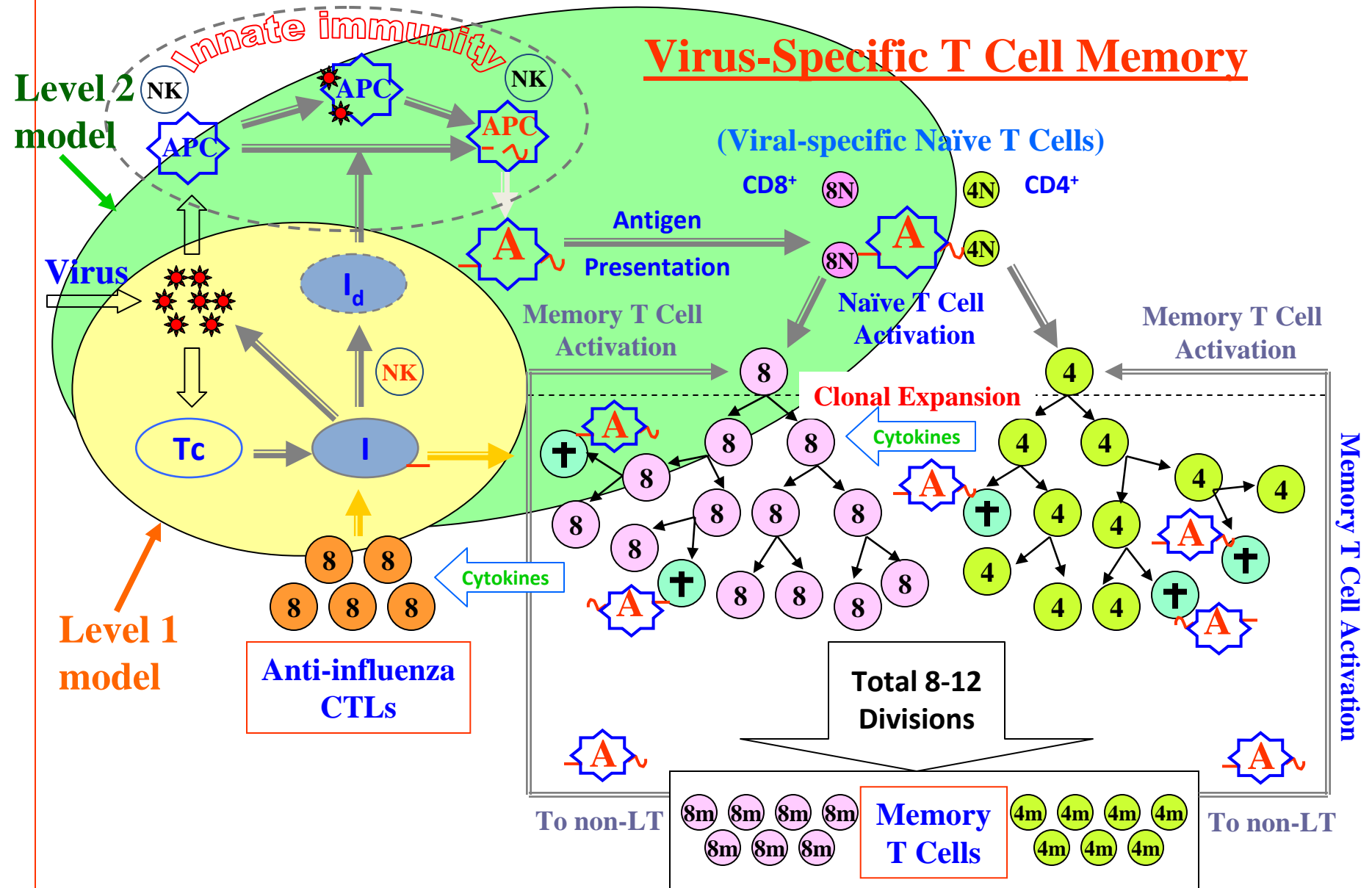
1. Transition between outbreak and suppression, when [CTL] is sufficiently raised;
2. Insensitivity to $[NK^*]$;
3. Critical [CTL] depends on value of diffusion constant D_v , with fine-scale features disappearing before coarse-scale

Under construction:

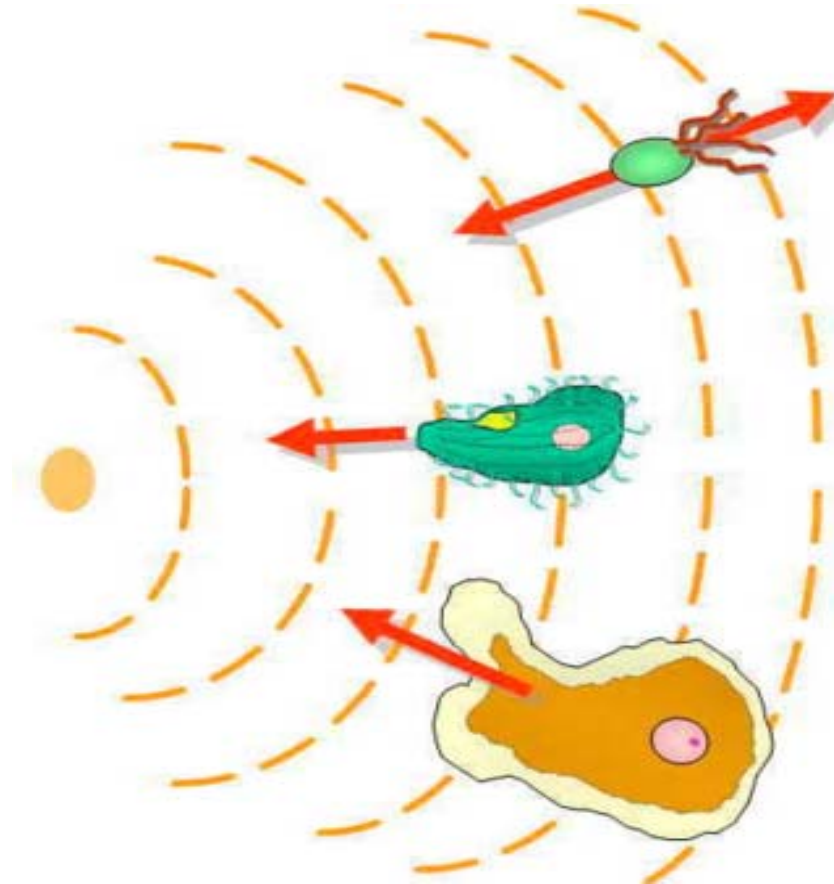
‘Level 2’ model of innate immune system

- Level-1 includes immune system only in a parametric sense, via death rates of infected epithelial cells
- **NEXT STEP:** Include strategic components of innate immune system, and couple its dynamics to Level 1 system. Current Level-2 model comprises
 - $INF\alpha/\beta, APC \rightarrow APC^*$, **chemotactic** and inflammatory cytokines, $INF\alpha, \beta, INF\gamma$, $NK \rightarrow NK^*$, neutrophils, Dendritic Cells
 - Diffusion and **chemotaxis**

Virus-Specific T Cell Memory



Example of chemotaxis



Directional cell migration in case of leukocytes (bottom two cells) but not bacteria (cell on top)

From: Francis Lin, Physics Dept., U of Manitoba