



# SITC 2017

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NATIONAL HARBOR  
MARYLAND

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Society for Immunotherapy of Cancer

November 8-12 • NATIONAL HARBOR, MD

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2017

# Systems Biology Modeling of the Immune System

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Society for Immunotherapy of Cancer

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# Presenter Disclosure Information

*Doron Levy*

The following relationships exist related to this presentation:

*No Relationships to Disclose*

# Mathematics and Immunotherapy

- CML – Cancer Vaccines
- CML – Signature of the (Individual) Immune Response
- Solid Tumors
- Complexity



# Immunotherapy and CML

OPEN  ACCESS Freely available online

PLOS COMPUTATIONAL BIOLOGY

## Dynamics and Potential Impact of the Immune Response to Chronic Myelogenous Leukemia

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### Abstract

Recent mathematical models have been developed to study the dynamics of chronic myelogenous leukemia (CML) under imatinib treatment. None of these models incorporates the anti-leukemia immune response. Recent experimental data show that imatinib treatment may promote the development of anti-leukemia immune responses as patients enter remission. Using these experimental data we develop a mathematical model to gain insights into the dynamics and potential impact of the resulting anti-leukemia immune response on CML. We model the immune response using a system of delay differential equations, where the delay term accounts for the duration of cell division. The mathematical model suggests that anti-leukemia T cell responses may play a critical role in maintaining CML patients in remission under imatinib therapy. Furthermore, it proposes a novel concept of an “optimal load zone” for leukemic cells in which the anti-leukemia immune response is most effective. Imatinib therapy may drive leukemic cell populations to enter and fall below this optimal load zone too rapidly to sustain the anti-leukemia T cell response. As a potential therapeutic strategy, the model shows that vaccination approaches in combination with imatinib therapy may optimally sustain the anti-leukemia T cell

# Complex Dynamics: CML, TKI, Immune system

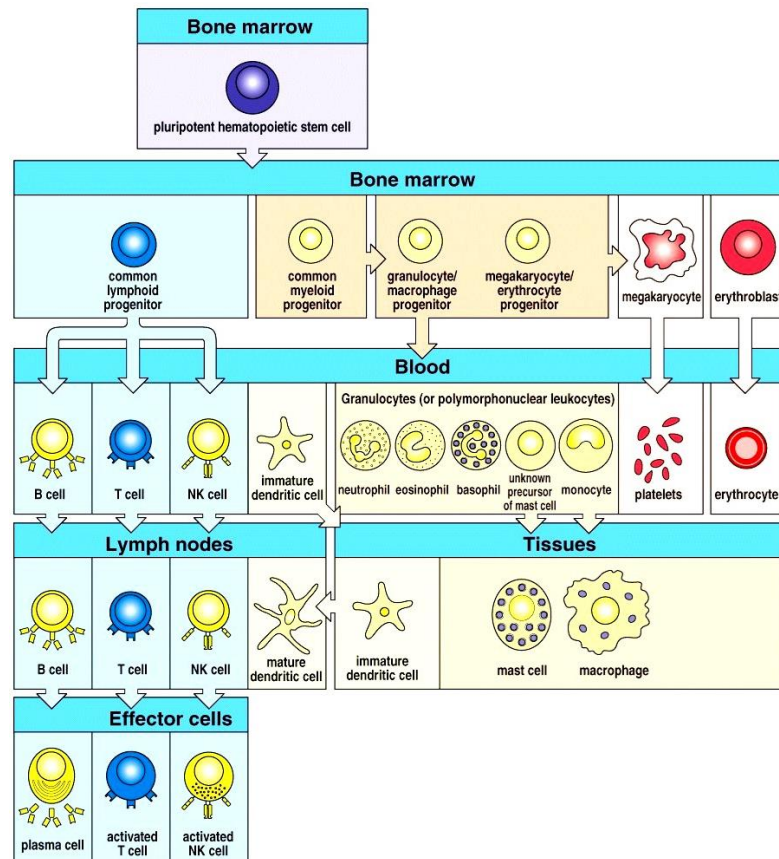
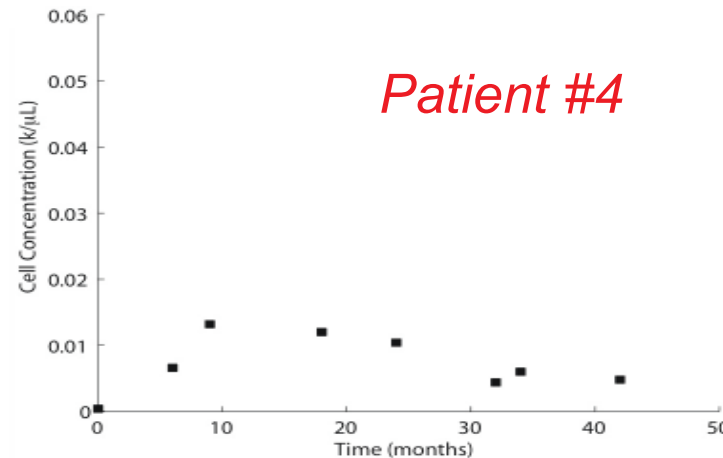
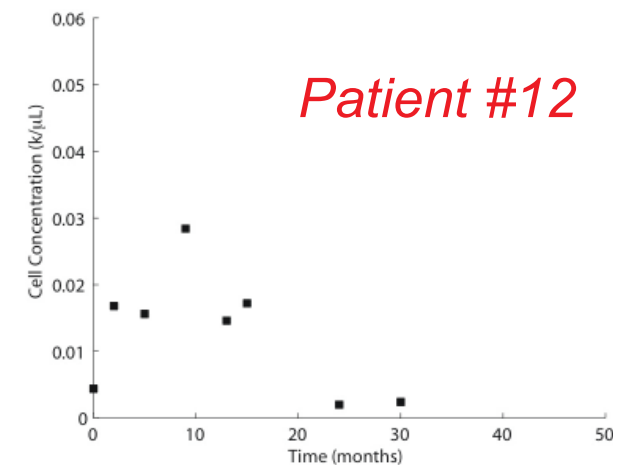


Figure 1-3 Immunobiology, 6/e. (© Garland Science 2005)



50 Months



50 Months

- Chen *et al.* (Blood, 2008)

# Complex Dynamics: CML, TKI, Immune system

- Converting the flowchart into a mathematical model

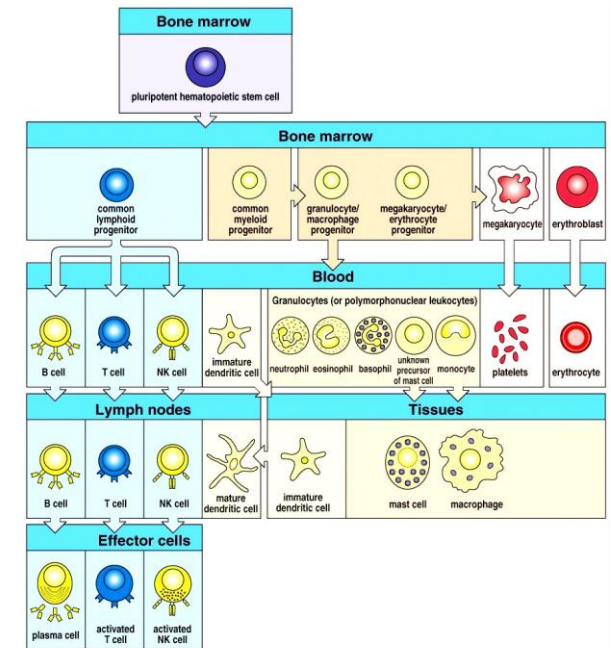
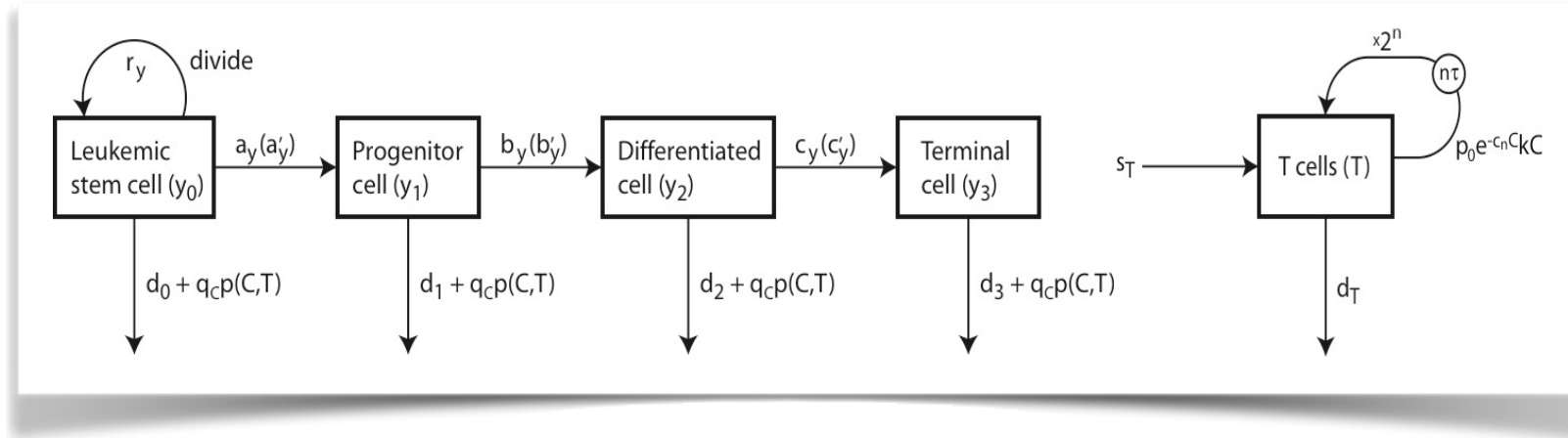
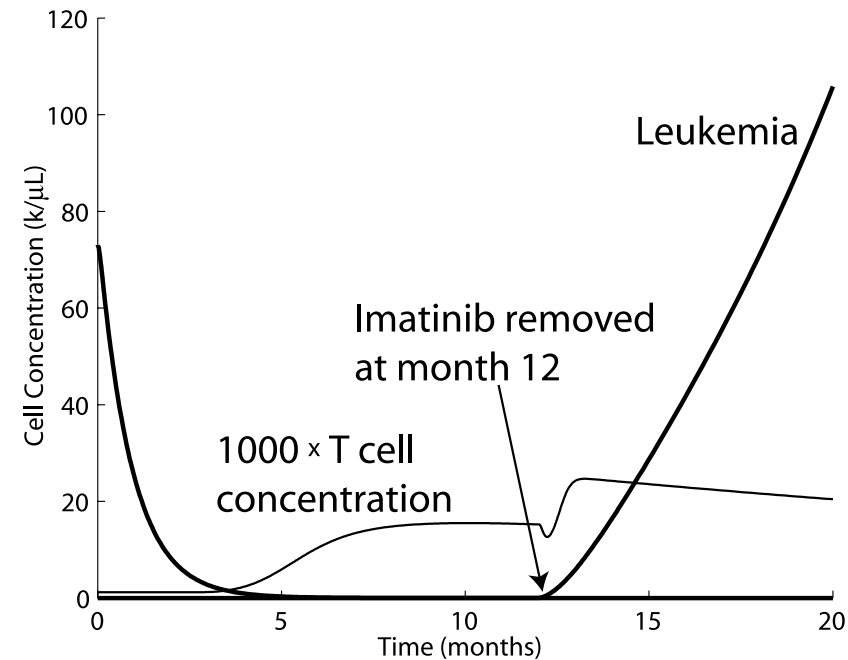
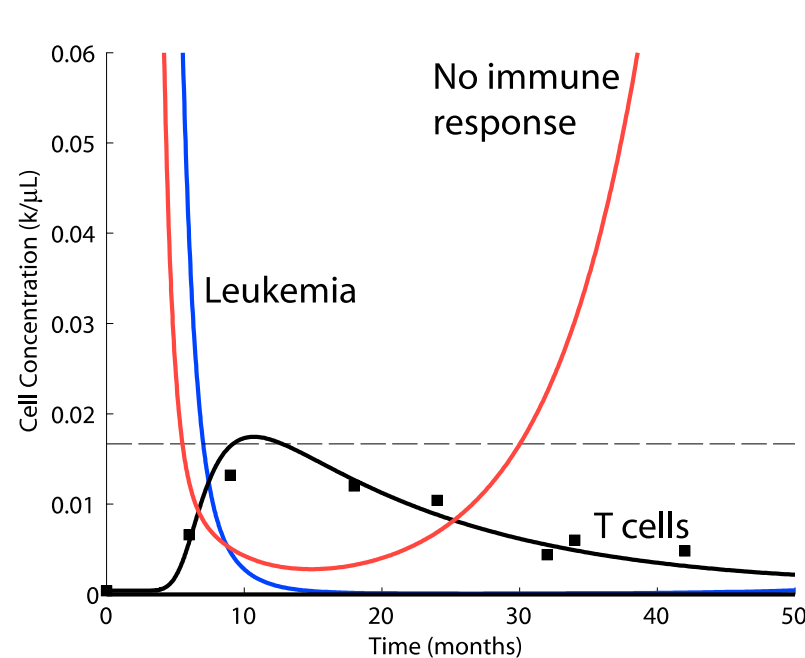


Figure 1-3 Immunobiology, 6/e. (© Garland Science 2005)

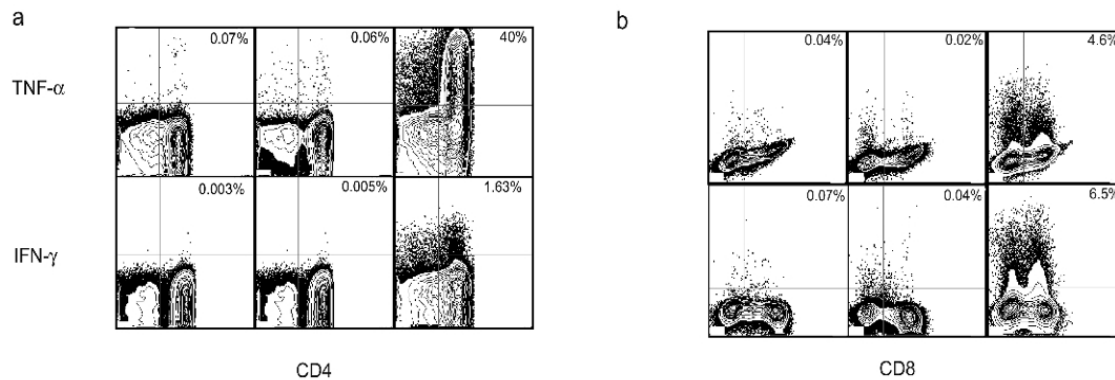
# Simulations of the Mathematical Model

- Red = Michor *et al.* (*Nature* '05). Blue = Kim *et al.* (*PLoS Comp Bio* '08)

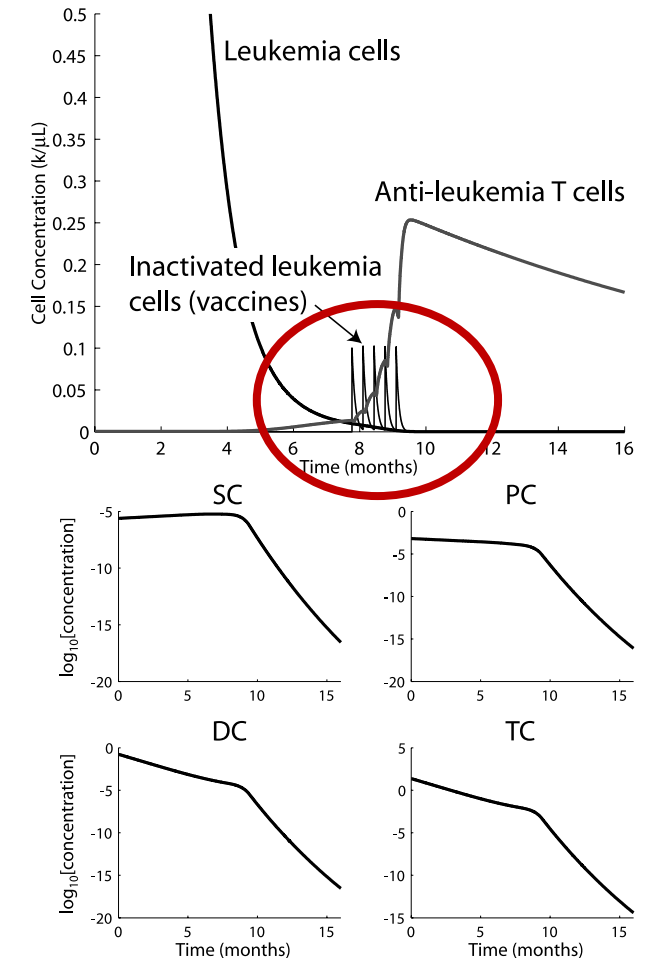




# Stimulating the Immune Response

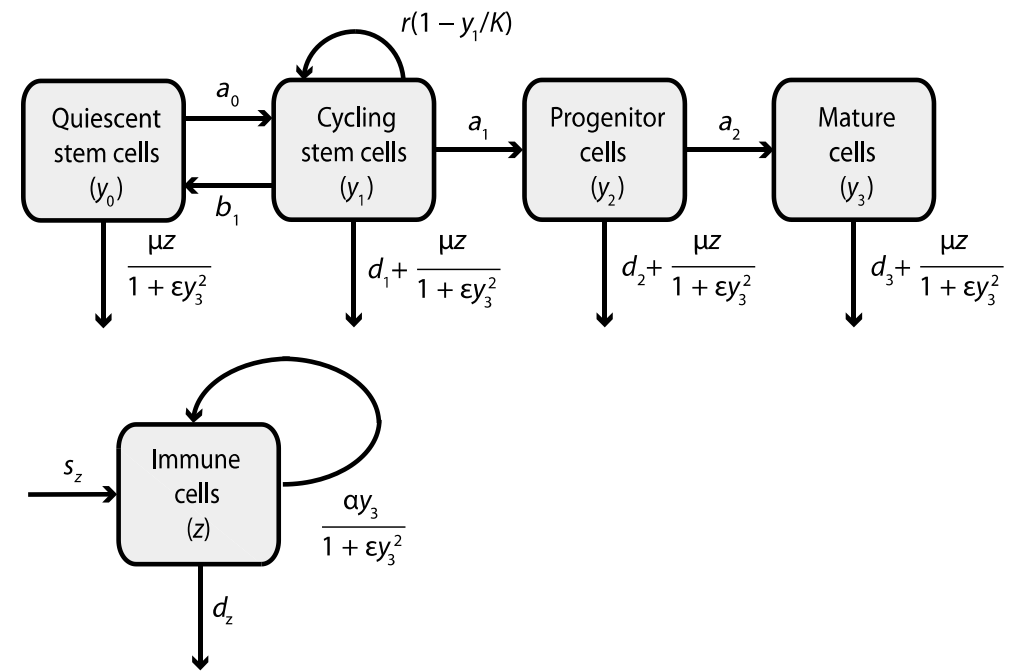
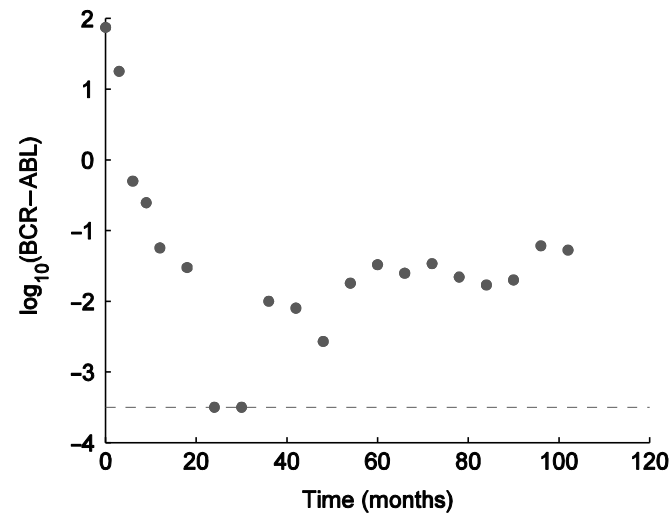
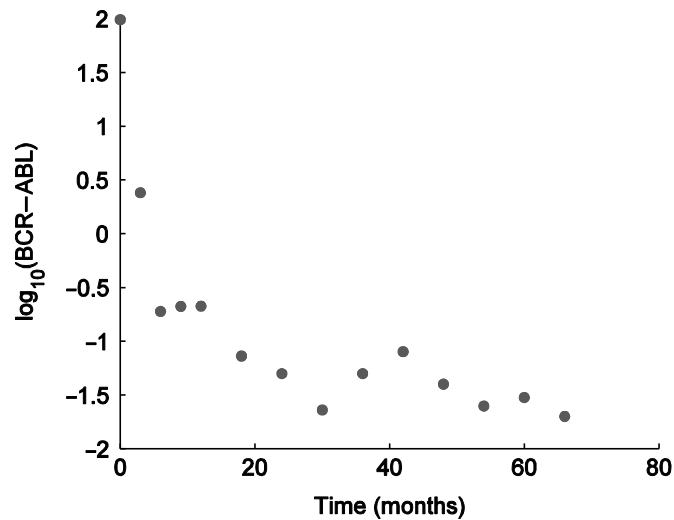


- Mathematical Model
- Dosage + Timing (!)
- Adjusted to each patient



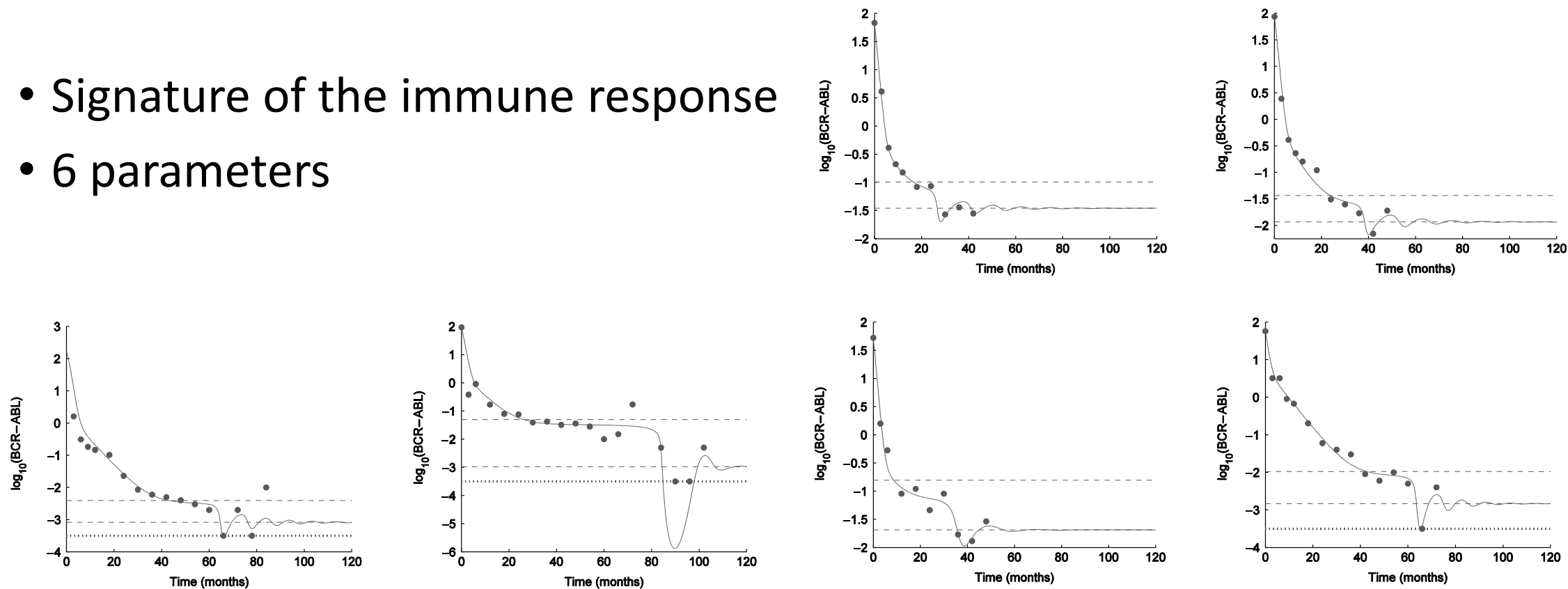
# Clapp et al., Cancer Research 2015

- CML Patients (Nicolini, Lyon)



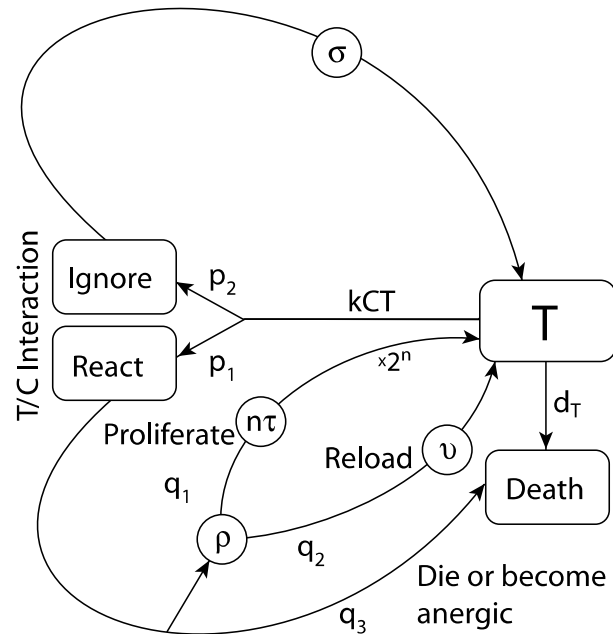
# Modeling immune system dynamics

- Signature of the immune response
- 6 parameters

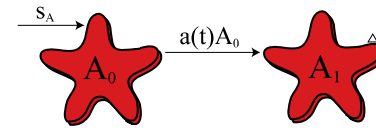


# Complexity of models

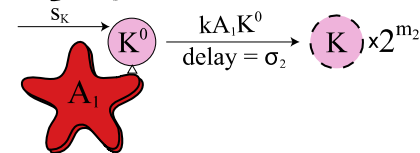
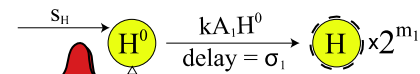
- Niculescu *et al.* (DCDS, 2008)



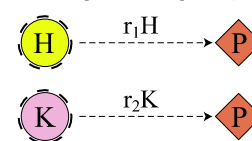
1) Migration of APCs to lymph node



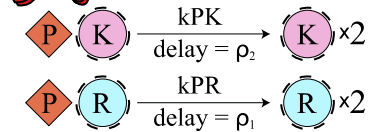
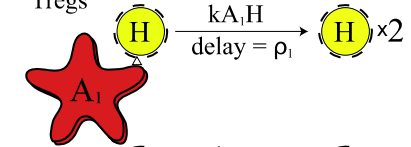
2) Initial T cell activation



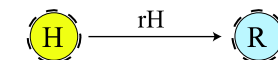
3) CD4+ and CD8+ T cells secrete positive growth signal (IL-2)



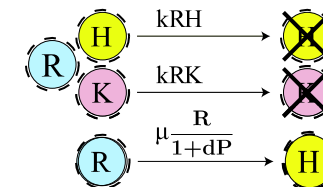
4) APC-driven proliferation of CD4+ cells. IL-2-driven proliferation of CD8+ cells & Tregs



5) CD4+ T cells differentiate into Tregs



6) Tregs suppress effector T cells, and transition into helper T cells.

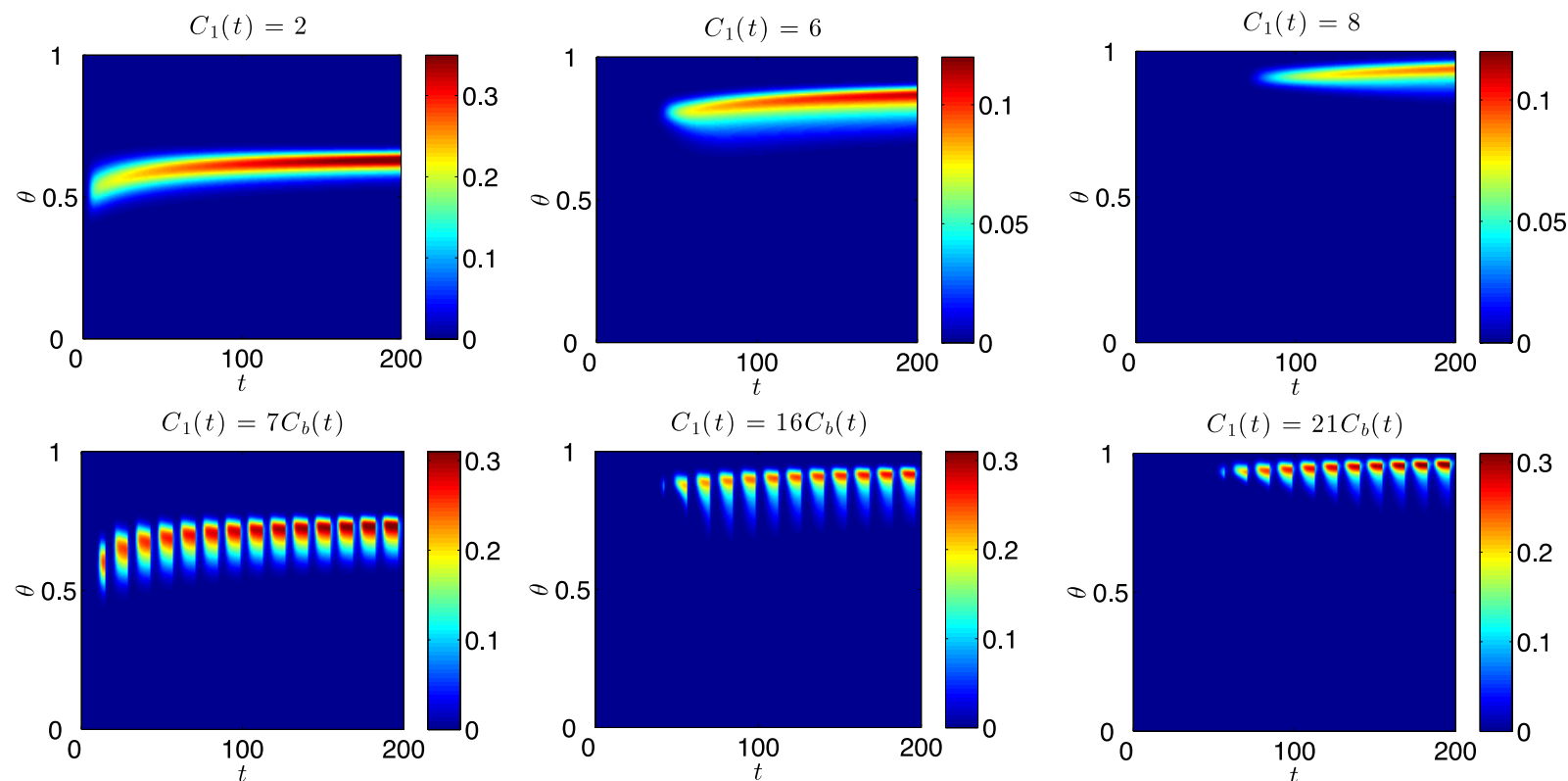


- Wilson & Levy (BMB, 2013)



# Heterogeneity of solid tumors

- Cho & Levy  
(BMB & JTB, 2018)



## Lessons and Take Home Messages

- Math complements other quantitative approaches
- Interdisciplinary: experiments/clinical/quantitative sciences
- Math can be useful in modeling **dynamics**
- Quantifying the **Timing + Dosage**
- **Personalized medicine**