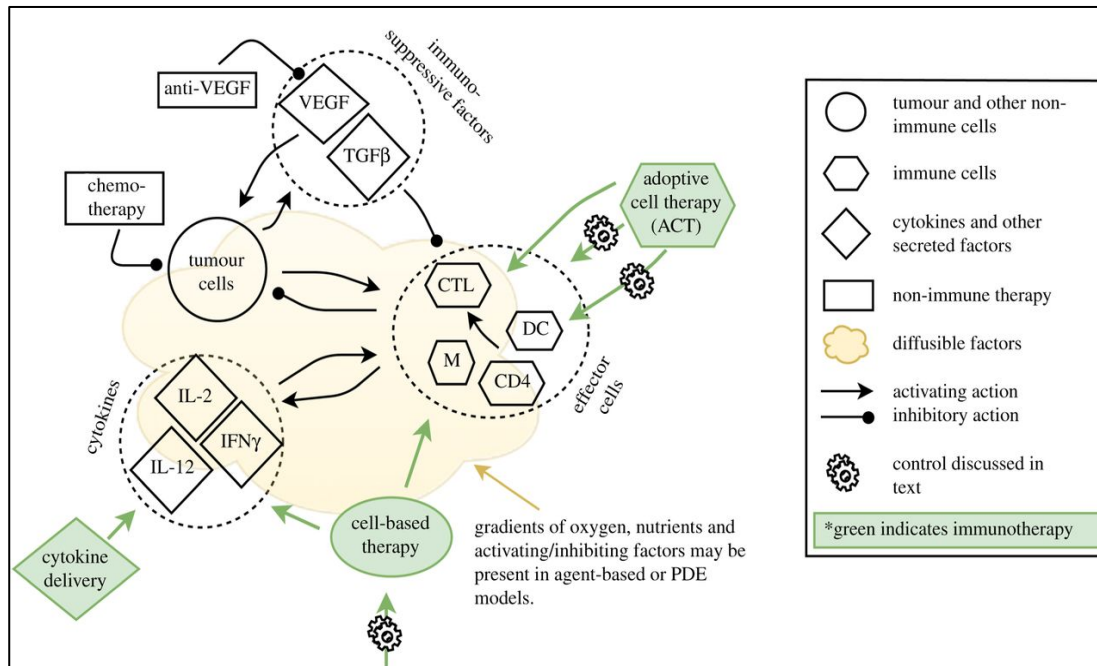


# Mathematical models for cancer immunotherapy: a review and new directions



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January 25<sup>th</sup>, 2019

# Acknowledgments

Center for Quantitative Medicine, UConn Health  
Reinhard Laubenbacher, Director



Department of Immunology, UConn Health  
Anthony Vella, Professor and Chair



Adam Adler, Professor



**UConn**  
**HEALTH**



NCI of the NIH postdoctoral  
fellowship award F32CA214030

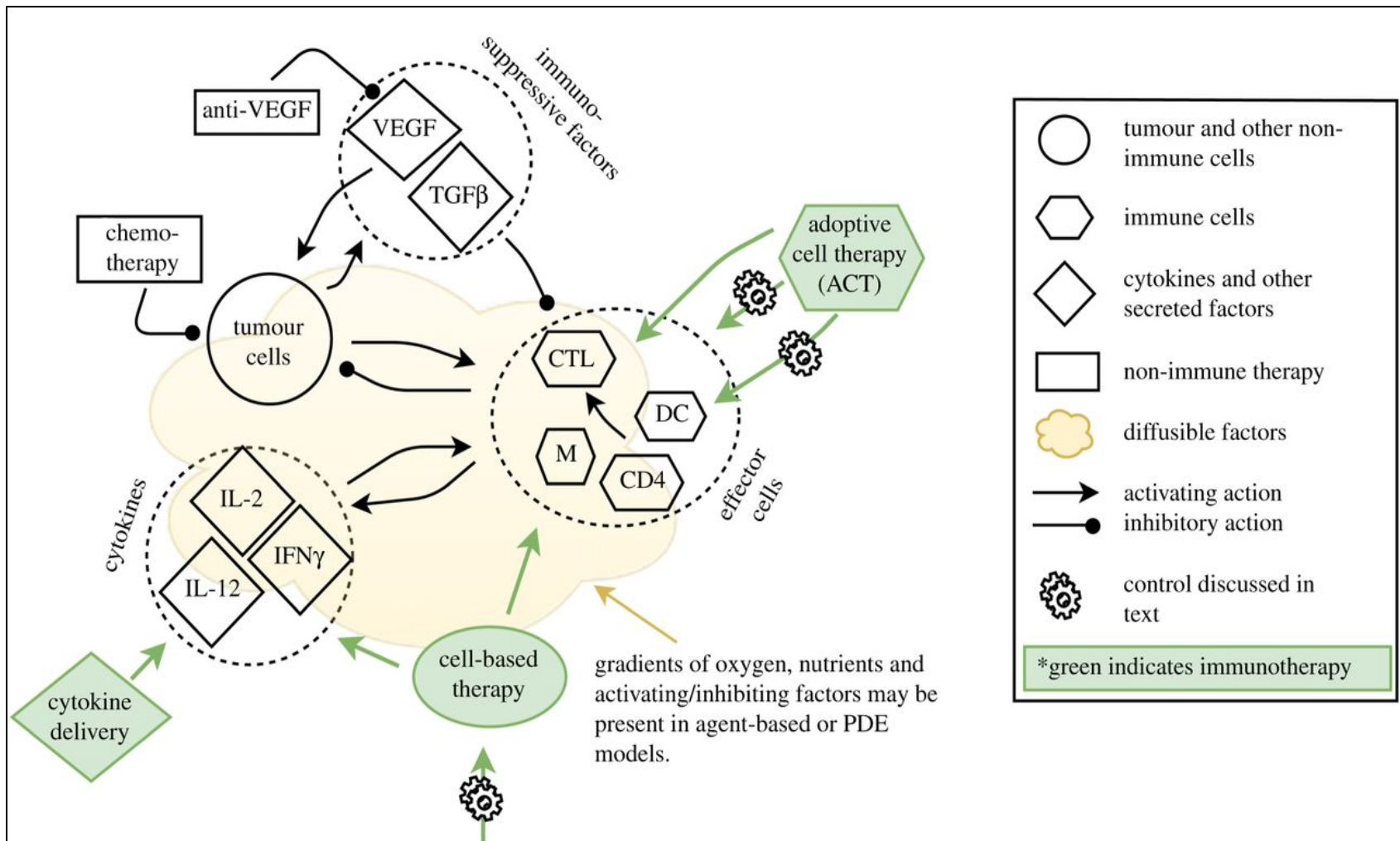
# Overview

- I. Mathematical models for immunotherapy: current progress and challenges<sup>1</sup>
  - i. Tumor classification for treatment and prediction of response
  - ii. Optimal scheduling and dosage of treatment
  - iii. Design and identification of combination treatment regimes
  - iv. Recommendations for further progress
  
- II. A mathematical model of combined CD8 T cell costimulation by 4-1BB (CD134) and OX40 (CD137) receptors<sup>2</sup>

<sup>1</sup>Konstorum A, Vella AT, Adler AJ, Laubenbacher RC (2017) Addressing current challenges in cancer immunotherapy with mathematical and computational modelling. *J. R. Soc. Interface* 14: 20170150.

<sup>2</sup>Currently manuscript in preparation, results not (yet!) published. Stay tuned!

# Summary of modeling efforts in immunotherapy



# Challenge: tumor classification for treatment and prediction of response

- Goal: to predict how a patient with a specific set of tumor characteristics will respond to a given treatment.
- Mathematical models can be used to predict effect of therapy that has not yet been tried in the clinic.

## Classic example: Panetta-Kirschner (PK) model<sup>1</sup>

- Models dynamics of effector ( $E$ ) and tumor ( $T$ ) cells, and the cytokine IL-2 ( $I_L$ ).
- Parameter of note:
  - antigenecity of tumor ( $c$ )
- Therapies represented by  $s_1, s_2$ .
  - $s_1$ ::= Adoptive Cellular Immunotherapy (ACI), injection of cultured immune cells with anti-tumor reactivity or Tumor Infiltrating Lymphocyte (TIL) therapy: tumor-derived lymphocytes cultured and reinjected into patient.
  - $s_2$ ::= external input of IL-2 into the system.

$$\frac{dE}{dt} = cT - \mu_2 E + \frac{p_1 E I_L}{g_1 + I_L} + s_1,$$

$$\frac{dT}{dt} = r_2(T)T - \frac{aET}{g_2 + T},$$

$$\frac{dI_L}{dt} = \frac{p_2 ET}{g_3 + T} - \mu_3 I_L + s_2,$$

with initial conditions

$$E(0) = E_0, \quad T(0) = T_0, \quad I_L(0) = I_{L_0}$$

<sup>1</sup>Kirschner, D and Panetta, JC (1998) Modeling immunotherapy of the tumor – immune interaction. *J. Math Biol* 37:235-252.

# Challenge: tumor classification for treatment and prediction of response

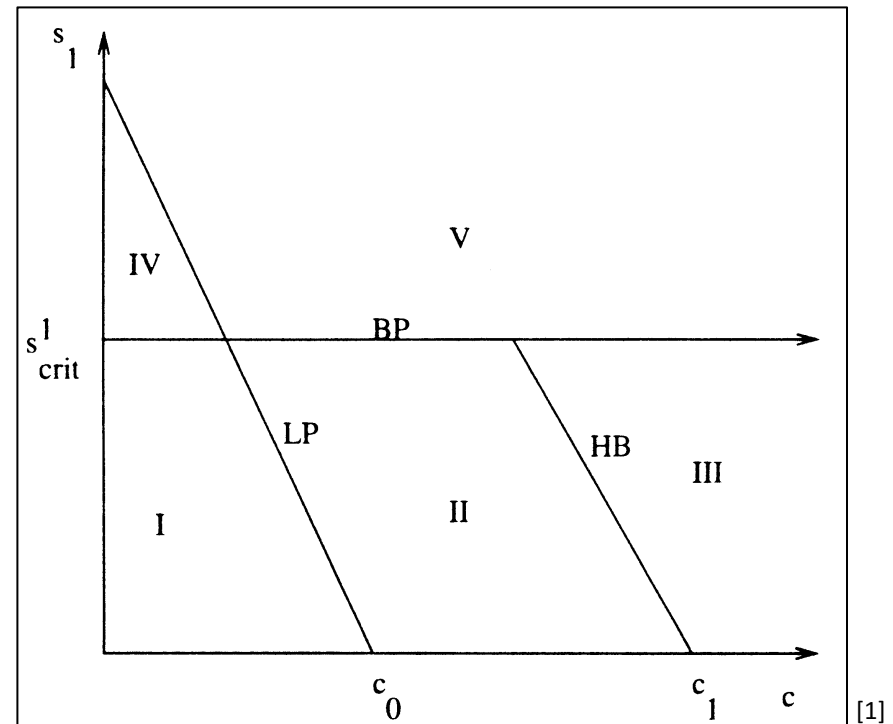
- Goal: to predict how a patient with a specific set of tumor characteristics will respond to a given treatment.
- Mathematical models can be used to predict effect of therapy that has not yet been tried in the clinic.
- Using linear stability analysis, identify

$$s_{\text{crit}}^1 = \frac{r_2 g_2 \mu_2}{a}$$

which impacts the tumor steady state.

- Region V has a stable steady state of tumor eradication, and Region IV may either tend to tumor eradication or survival depending on the initial conditions.
- Regions I-III do not produce tumor eradication.

*Therefore, can predict response to (and potentially modify) treatment with knowledge of system parameters.*



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# Challenge: tumor classification for treatment and prediction of response

- More complex systems require numerical analysis (vs. linear stability), and have focused on the concept of thresholds for predicting patient response. Some examples:
  - [Kronik et al. \(2012\)<sup>1</sup>](#) modeled *ex vivo* expanded tumor-specific T cell transfer for melanoma using a system of ODEs and used clinical data for retroactive validation.
    - Varied initial tumor size and growth rate to imitate a virtual population. Four different therapy regimens were simulated to correspond to four different clinical trials. Identified a tumor-size threshold for therapy effectiveness which matched patient data.
  - Wells et al (2015)<sup>2</sup> developed a hybrid discrete-continuous (HDC) agent-based model (ABM). These models treat cells as agents that can interact with and respond to other cells.
    - Observed that the ratio of M2 macrophages to other cell types was predictive of tumor survival. Spatial model necessary for predictive capability.
  - Eikenberry et al (2009)<sup>3</sup> developed a PDE of melanoma with immune infiltrate.
    - Showed that surgical removal of tumors with high levels of immune infiltrate could promote growth of satellite metastases, as was observed clinically.
    - Hence, provided a model-based hypothesis for tumor classification with respect to responsiveness to surgery.

<sup>1</sup>Kronik et al. (2012) Improving T-cell immunotherapy for melanoma through a mathematically motivated strategy: efficacy in numbers? *J. Immunother.* 35, 116-124.

<sup>2</sup>Wells et al. (2015) Spatial and functional heterogeneities shape collective behavior of tumor-immune networks. *PLoS Comput. Biol.* 11, e1004181.

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# Challenge: optimal scheduling and dosage of treatment

- If you know the treatment – how to determine the optimal schedule and dosage (not based on trial and error)?
- Techniques to identify optimal treatment schedules *in silico* include:

## Optimal control theory<sup>1</sup>

- Used for models based on continuum methods.
- States the problems of finding an optimal treatment plan in the framework of a controlled dynamical system.
- Example: identify optimal ACI therapy in PK model to minimize final tumor concentration<sup>2</sup>

## Genetic Algorithms<sup>3</sup>

- Belong to class of evolutionary algorithms.
- System can be agent-based, discrete, continuous, etc.
- Theory based on principles of genetic evolutionary theory.
- Example: identify optimal vaccine schedule for the Triplex vaccine (for HER-2/neu-positive BC) using an agent-based SimTriplex Model<sup>4</sup>

<sup>1</sup>Evans LC (2017). An introduction to mathematical optimal control theory, Version 0.2. See <https://math.berkeley.edu/evans/control.course.pdf>

<sup>2</sup>Burden et al (2004). Optimal control applied to immunotherapy. *Discr. Continuous Dyn. Syst. Series B* 4, 135-136.

<sup>3</sup>Whitley D. (1994). A genetic algorithm tutorial. *Stat. Comput.* 4, 65-85

<sup>4</sup>Lollini et al. (2006). Discovery of cancer vaccination protocols with a genetic algorithm driving an agent based simulator. *BMC Bioinform.* 7, 352.

# Challenge: design and identification of combination treatment regimes

- Mathematical modeling can help in rational design of combination immunotherapy (either with just immunotherapeutic agents or with immune- and non-immunotherapeutic agents) to maximize treatment response.

## Example 1: de Pillis et al. (2009)<sup>1</sup>: chemo-immunotherapy model.

- Model comprised of six ODEs for combination chemo- and immunotherapy that includes tumor and immune cells, and concentrations of chemo- and immuno-therapy drugs.
- Found that success of combination versus monotherapy different based on initial patient characteristics (derived from human clinical trials of metastatic melanoma).

Simulation	$T = 1 \times 10^6$ cells		$T = 1 \times 10^7$ cells		$T = 1 \times 10^8$ cells		$T = 1 \times 10^9$ cells	
	9	10	9	10	9	10	9	10
No treatment	<i>x</i>	<i>x</i>	<i>o</i>	<i>o</i>	<i>o</i>	<i>o</i>	<i>o</i>	<i>o</i>
Chemotherapy	<i>x</i>	<i>x</i>	<i>x</i>	<i>x</i>	<i>x</i>	<i>x</i>	<i>o</i>	<i>o</i>
Immunotherapy	<i>x</i>	<i>x</i>	<i>x</i>	<i>o</i>	<i>o</i>	<i>o</i>	<i>o</i>	<i>o</i>
Chemo-immuno	<i>x</i>	<i>x</i>	<i>x</i>	<i>x</i>	<i>x</i>	<i>o</i>	<i>o</i>	<i>o</i>

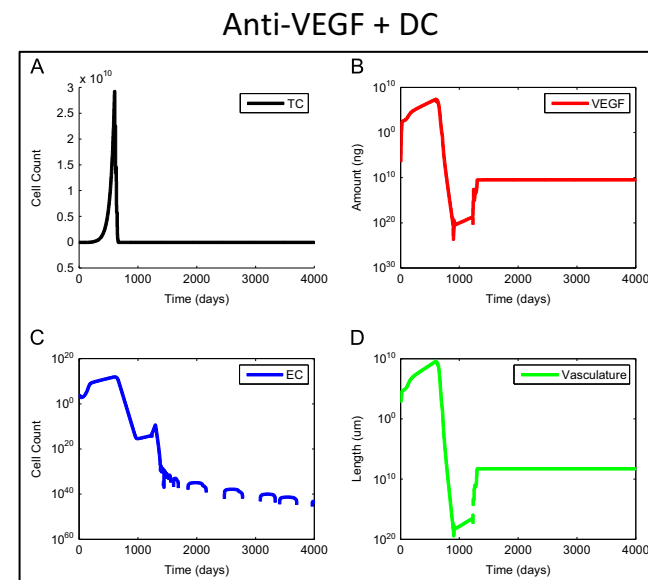
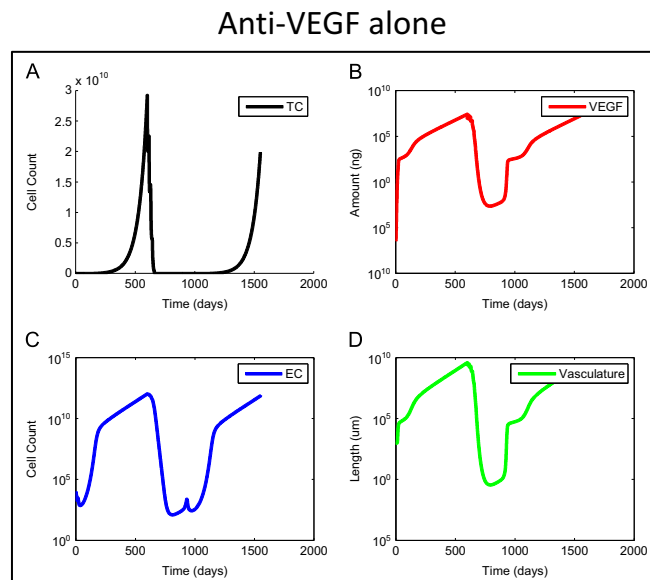
<sup>1</sup>de Pillis et al. (2006) Mixed immunotherapy and chemotherapy of tumors: modeling, applications and biological interpretation. *J. Theor. Biol.* 238: 841-862.

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## Example 2: Soto-Ortiz et al. (2016)<sup>1</sup>: anti-angiogenic and immunotherapy model

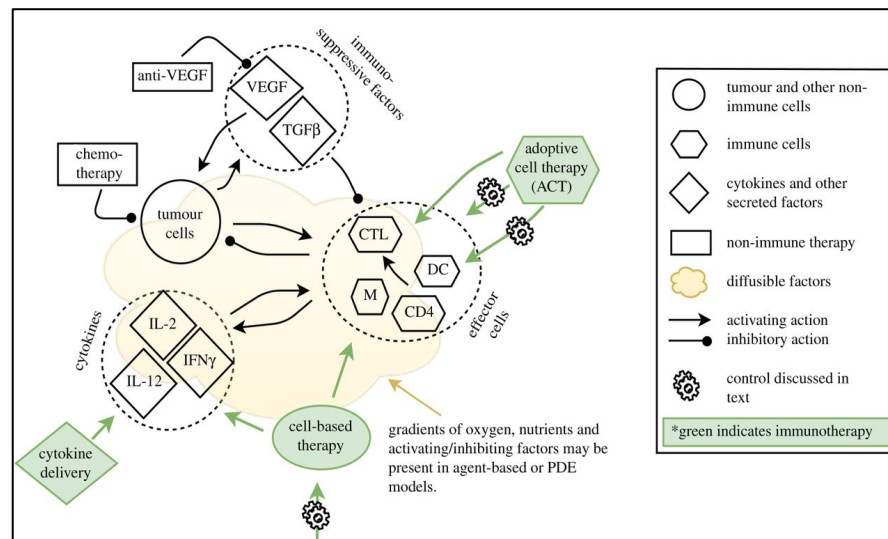
- Model comprised of 18 ODEs that include tumor, immune and vascular endothelial cells, and several cytokines and growth factors modeling anti-VEGF therapy (VEGF has pro-angiogenic and immunosuppressive activity) and administration of DC cells.



<sup>1</sup>Soto-Ortiz and Finley et al. (2016) A cancer treatment based on synergy between anti-angiogenic and immune cell therapies. *J. Theor. Biol.* 394: 197-211.

# Recommendations

- I. Intracellular and multi-scale modeling
  - i. Can give insights into therapeutic action at intracellular level, and relative contribution of cell-cell and intracellular activities.
  - ii. Can be developed from existing models of signaling cascades in cancers.
- II. Addressing toxicity
  - i. Incorporation of immunotherapy-related toxicity can help to optimize therapy predictions for maximum efficacy/minimum toxicity.
- III. Experimental and clinical validation of immunotherapy models.
  - i. Main bottleneck for wider validation and use of mathematical and computational models for purpose of developing novel therapies.
  - ii. Needs to be community-level initiative (at scale of organization or funding agencies).



# Recommendations

## Thank you!

