Ordinary Differential Equation (ODE) Models of Cancer-Immune Interactions

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Overview

Cancer Immunology Background

2 ODE Background

3 ODE Models of Cancer Immune Interaction

The Immune System

- The immune system defends the body against infections and abnormal cells.
- Composed of:
 - Innate immunity: fast, non-specific (e.g., macrophages, NK cells)
 - Adaptive immunity: slower, specific, with memory (e.g., T cells, B cells)
- T cells:
 - CD8⁺ T cells: directly kill infected or cancerous cells.
 - CD4⁺ T cells: help other immune cells through cytokine signaling.
- Immune surveillance: the immune system constantly checks for abnormal or foreign cells.

Cancer

- Cancer is the uncontrolled growth of abnormal cells in the body.
- Hallmarks of tumor cells:
 - Self-sufficiency in growth signals
 - Insensitivity to anti-growth signals
 - Resistance to cell death (Apoptosis)
 - Angiogenesis
 - Invasion & Metastasis

The Immune System and Cancer

• Immune surveillance: T cells and NK cells can detect and destroy abnormal cells, including cancerous ones.

• Tumor-immune dynamics:

- The immune system may control small tumors for years.
- Some cancer cells develop ways to avoid or suppress immune attacks.

• Immune evasion:

- Tumors may downregulate immune-activating signals.
- They may express immune checkpoint molecules or recruit suppressive cells.
- These complex interactions between cancer and the immune system can be explored using mathematical models.

Ordinary Differential Equations in Biology

- What are Ordinary Differential Equations? Equations used to describe how a system changes over time
- Why Use ODEs?
 - Capture dynamics of populations (e.g., bacteria, immune cells).
 - Model interactions between biological entities (e.g., predator-prey).
- ODEs can reveal steady states, growth/decay patterns, and oscillations, helping predict system behavior under varying and complex conditions.
- Use in Tumor-Immune Models: Tumor cells and immune cell populations can be represented using variables, allowing study of their interactions.

One Equation Tumor Growth Models

Simple equation for tumor growth:

$$\frac{dx}{dt} = x f(x), \qquad f(x) = p(x) - d(x)$$

- x(t): tumor cell population at time t (x(0) > 0).
- p(x): per-capita **proliferation** rate; d(x): per-capita **death** rate.
- States:

$$\begin{cases} p(x) > d(x) & \Rightarrow \text{growth} \\ p(x) = d(x) & \Rightarrow \text{dormancy} \\ p(x) < d(x) & \Rightarrow \text{decay} \end{cases}$$

Two-Equation Tumor-Immune Interaction Models

Immune cell interactions with tumors can be described with two equations.

Predator-prey model: immune cells are predators, tumor cells are prey.

$$\frac{dx}{dt} = xf(x) - d_x(x,y), \qquad \frac{dy}{dt} = p_y(x,y) - d_y(x,y) - a_y(y) + \phi(t),$$

- x: size/density of tumor cells y: size/density of effector (immune) cells
- a: apoptosis $\phi(t)$: treatment

Steady States

• A steady state (or equilibrium) is a value \bar{x} such that the system remains there:

$$f(\bar{x}) = \bar{x}.$$

- Stable steady state: When the system is disturbed it returns to \bar{x} . (eg. a ball in a valley)
- Unstable steady state: When the system is disturbed it does not return to \bar{x} . (eg. a ball at the top of a hill)

Stability

- Start with solution: $x_n = \bar{x} + x'_n$, where x'_n is a small perturbation.
- Then

$$x'_{n+1} = x_{n+1} - \bar{x} = f(x_n) - \bar{x} = f(\bar{x} + x'_n) - \bar{x}.$$

• Linearize:

$$x'_{n+1} \approx a x'_n, \qquad a = f'(\bar{x}).$$

• Stability: \bar{x} is stable if |a| < 1, the system will return to \bar{x} .

Bifurcation

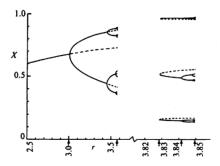


Figure 1: Bifurcation Diagram

Key ideas

- Horizontal axis: parameter value (e.g., r).
- Vertical axis: represents magnitudes of steady states of the equation.
- Branches: represents dependence of state level on parameter.
- Bifurcation points: where a change in the parameter's value changes the qualitative behaviour of the dynamical system (e.g. changes in number/stability of steady states, steady states may appear or disappear, oscillating steady states)

Simulating Oscillatory Dynamics

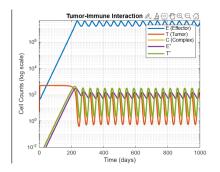


Figure 2: Oscillatory decay of tumor

- Model: Kuznetsov (1994) ODEs for tumor cells (T) and effector cells (E), with interaction/lysis terms and immune stimulation.
- Solver/Code: MATLAB ode45; parameters from: Interactions between the Immune System and Cancer: A Brief Review of Non-Spatial Mathematical Models
- Plot: Log-scale cell counts over time (days)

Impact of Parameters

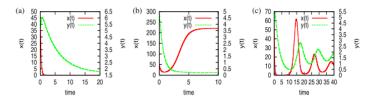


Figure 3: Effect of Parameters on System Behaviour

- Panel A: s=0.318 (high), d=0.1908 (low), $b=2\times 10^{-3}$ (moderate) exponential tumor decay
- Panel B: s = 0.318 (high), d = 2.0 (very high), $b = 4 \times 10^{-3}$ (high) tumor initially shrinks, but immune cell population decays and tumor grows again
- Panel C: s = 0.1181 (low) d = 0.3743 (moderate), $b = 2 \times 10^{-3}$ (moderate) tumor size decays in an oscillatory manner

Parameters:

- s immune cell recruitment rate (day⁻¹) d – immune cell death rate (day⁻¹)
- b the strength to which tumor cells stimulate an immune response (cell⁻¹ day⁻¹)

Applications of ODEs in Cancer Immunology

- Guide treatment choices: Simulate effects of immunotherapies to identify strategies with the highest chance of success.
- Predict patient outcomes: Forecast whether a tumor will regress, remain dormant, or relapse under different conditions or treatment protocols.
- Optimize therapy schedules: Determine optimal doses and timing that maximize tumor destruction while minimizing side effects.
- Explore Tumor Immune dynamics: Identify conditions leading to steady states, oscillations, or bifurcations to better understand clinical phenomena such as remission and recurrence.
- Test scenarios outside of clinical settings: Explore the impact of changes in tumor growth rate, immune activation, or other variables, without needing to experiment directly on patients.

References I

Edelstein-Keshet, L. (2005).

 $\begin{tabular}{ll} Mathematical Models in Biology, volume 46 of Classics in Applied Mathematics. \end{tabular}$

Society for Industrial and Applied Mathematics.

Eftimie, R., Bramson, J. L., and Earn, D. J. D. (2011). Interactions between the immune system and cancer: A brief review of non-spatial mathematical models. *Bulletin of Mathematical Biology*, 73(1).

Hanahan, D. and Weinberg, R. A. (2011). Hallmarks of cancer: The next generation. *Cell*, 144(5).

References II



Kuznetsov, V. A., Makalkin, I. A., Taylor, M. A., and Perelson, A. S. (1994).

Nonlinear dynamics of immunogenic tumors: Parameter estimation and global bifurcation analysis.

Bulletin of Mathematical Biology, 56(2).



Sadeghalvad, M., Mohammadi-Motlagh, H. R., and Rezaei, N. (2023).

Introduction to cancer immunology.

In Rezaei, N., editor, *Handbook of Cancer and Immunology*. Springer, Cham.



Waldman, A. D., Fritz, J. M., and Lenardo, M. J. (2020).

A guide to cancer immunotherapy: from t cell basic science to clinical practice.

Nature Reviews Immunology, 20.

Thank you!

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