



Tumor-Normal-Immune Interaction Model

Aditi Deshpande
Mentored by Justin Hager
Fall 2025

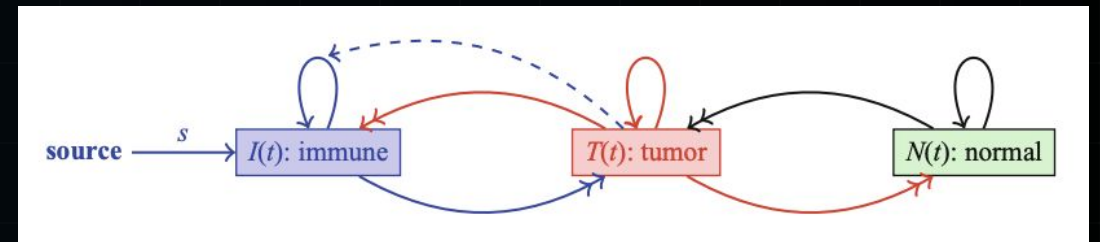
Motivation

Understanding how normal cells, tumor cells, and immune cells interact is critical for predicting cancer progression and optimizing treatment.

Key Questions:

- Do these populations reach a stable balance, or do they oscillate?
- What conditions allow tumors to escape immune control?

Goal: Develop mathematical models to characterize tumor-immune dynamics, predict bifurcations and limit cycles, and evaluate how different treatment strategies affect system behavior.



| 1. The Dimensional Model

We model the interactions between three distinct populations over time t .

- > **N (Normal Cells)**: Logistic growth, inhibited by tumors.
- > **T (Tumor Cells)**: Fast growth, killed by immune cells.
- > **I (Immune Cells)**: Stimulated by tumor presence.

$$dN/dt = r_2N(1 - N/K_2) - c_4TN$$

$$dT/dt = r_1T(1 - T/K_1) - c_2IT - c_3TN$$

$$dI/dt = s + aT - c_1IT - d_1I$$

2. Scaling & Numerical Method

Dimensionless System

We scale variables to reduce parameters and simplify analysis.


$$N \rightarrow n, T \rightarrow t, I \rightarrow i$$

This reveals the intrinsic dynamics independent of physical units.

Forward Euler Scheme

The core integration method used for solving the ODEs.

$$y_{k+1} = y_k + \Delta\tau \cdot f(y_k)$$

 We use a small time step $\Delta\tau = 0.01$ to ensure numerical stability and accuracy for the oscillatory dynamics.

| 3. Stability Analysis

We analyze the system's behavior near the coexistence equilibrium.

Methodology

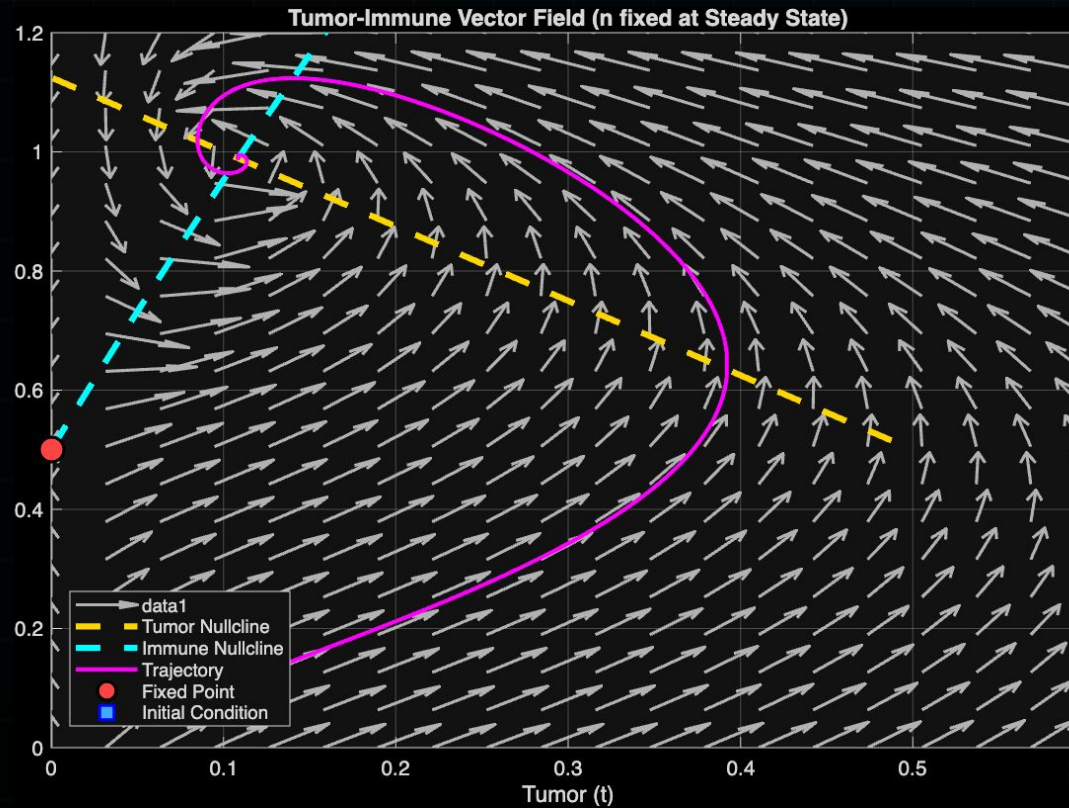
1. **Find Fixed Point (FP):** Solve $f(n,t,i) = 0$ using `fsolve`.
2. **Jacobian Matrix (J):** Linearize the system at the
3. **Eigenvalues (λ):** Calculate `eig(J)`.

Result Interpretation

$$\text{Re}(\lambda) > 0$$

A positive real part indicates an **Unstable Spiral**. The system does not settle at the fixed point but enters a sustained oscillation known as a **Limit Cycle**.

4. Phase Plane & Nullclines



Visualizing Dynamics

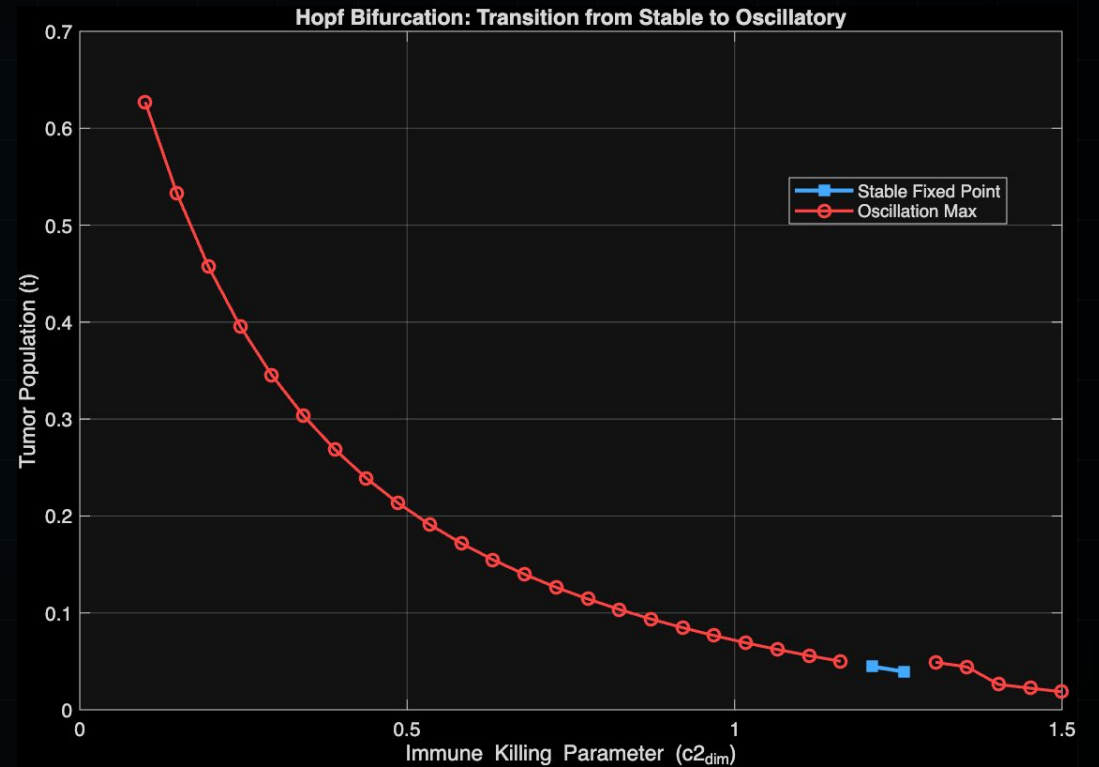
- > **Nullclines:** Curves where $\frac{dt}{dt} = 0$ (Yellow) and $\frac{di}{dt} = 0$ (Blue).
- > **Vector Field:** Arrows show the direction of flow at any point.
- > **Trajectory:** The magenta path shows the system spiraling away from the unstable fixed point towards the limit cycle.

5. Hopf Bifurcation

We varied the Immune Killing Parameter $c2_dim$ to observe behavioral changes.

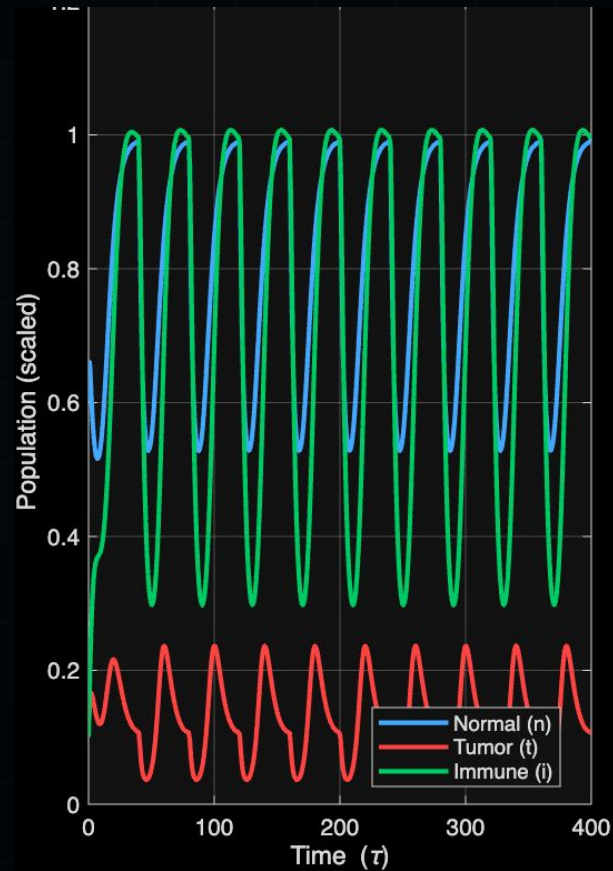
Parameter: $c2_dim$
Range: 0.1 to 1.5

As $c2_dim$ decreases, the system undergoes a **Hopf Bifurcation**, transitioning from a stable steady state to a stable limit cycle (oscillation).



6. Baseline Oscillations

The resulting time series for the dimensionless baseline model.



7. Adding Chemotherapy

We extend the ODE system to 4 equations by adding a drug variable u .

$$du/d\tau = v(\tau) - d_u u$$

Treatment Schedules $v(\tau)$:

- > Constant Infusion
- > Pulsed Therapy (Periodic)
- > Single Bolus

Kill Term Mechanics

The drug affects populations based on a saturation function:

$$\text{Kill} = a_x(1 - e^{-u})$$

Selectivity: $a_2 > a_1 > a_3$. We assume the drug kills Tumor cells most effectively, followed by Normal cells, with minimal impact on Immune cells.

8. Reaction-Diffusion PDE

Moving from a "well-mixed" flask to spatial tissue dynamics.

$$\partial T / \partial \tau = D \nabla^2 T + f(T, I, N)$$

New Variables

- > $T(x,y)$: Tumor density in 2D space.
- > $N(x,y)$: Nutrient concentration (replaces Normal cells ODE).
- > D : Diffusion coefficient.

Physical Meaning

Tumors now cluster and consume local nutrients. The nutrient N diffuses from blood vessels (boundaries) into the tissue.

9. Solving the PDE

Finite Difference Method (FDM)

We discretize space into a grid (N_x, N_y) and approximate the Laplacian operator ∇^2 .

5-Point Stencil:

Calculated using array shifting ('circshift') to compute the difference between a cell and its 4 neighbors.

Stability Constraints

Explicit PDE solvers require strict time steps.

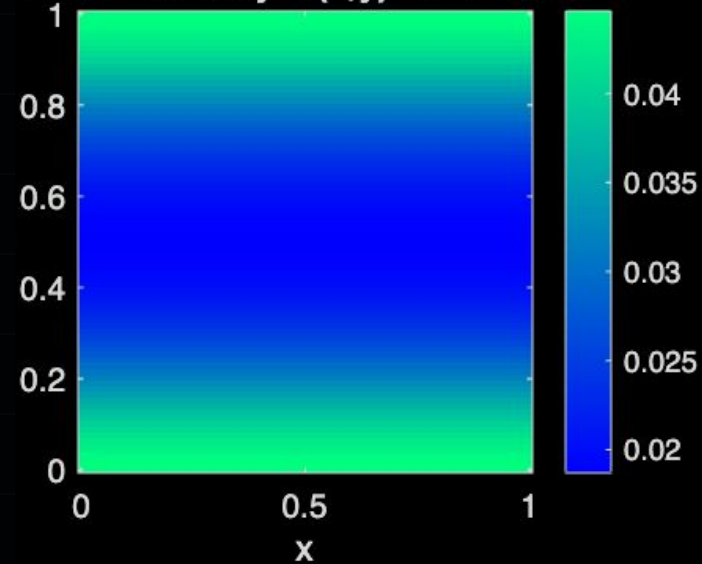
CFL Condition:
 $dt < dx^2 / (4D)$

If dt is too large, the simulation creates artificial artifacts and explodes.

10. Spatial Heterogeneity

Unlike the ODE model, the PDE shows localized effects.

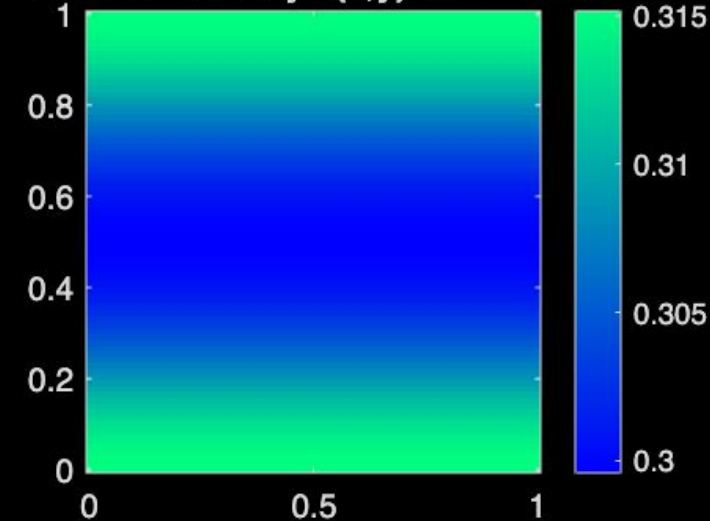
Tumor density $T(x,y)$ at $\tau=100.0$



Tumor $T(x,y)$

Localized growth centers.

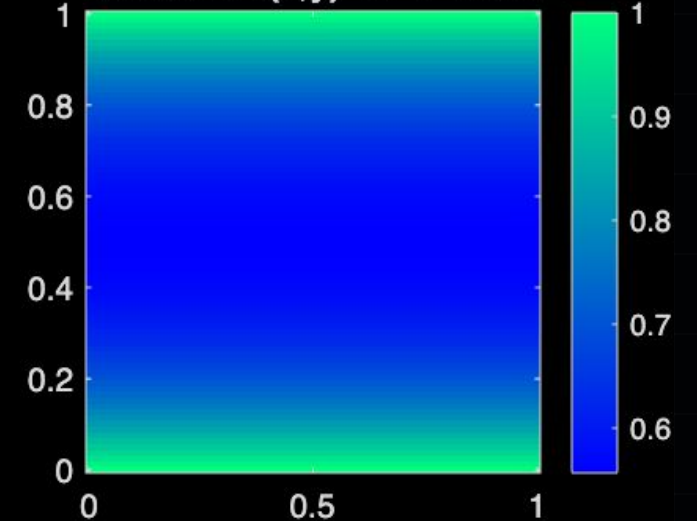
Immune density $I(x,y)$ at $\tau=100.0$



Immune $I(x,y)$

Infiltration at tumor edges.

Nutrient $N(x,y)$ at $\tau=100.0$



Nutrient $N(x,y)$

Depletion in tumor cores.

11. Citations & Acknowledgements

References

Mathematical Oncology 2013

d'Onofrio, A., & Gandolfi, A. (Eds.). Birkhäuser, 2014.

ISBN: 978-1-4939-0458-7

Source:

<file:///Users/aditideshpande/Downloads/978-1-4939-0458-7.pdf>

Mathematical Models in Biology

Müller, J. (2004). Lecture Notes.

Source:

<http://www.bio-nica.info/biblioteca/muller2004mathematicalmodelsinbiology.pdf>



Special Thanks

To the **UNC Directed Reading Program** for providing this research opportunity.

And to my mentor,

Justin Hager



Questions?

Thank you for listening.