

Numerical Solutions of an Immunology Model Using Reaction-Diffusion Equations with Stochastic Source Terms

Tim Lucas
Center for Computational Immunology
Duke University

January 7, 2008

Background

- Immune cells secrete **cytokines** in order to communicate with other immune cells. Cytokines are modeled continuously as **soluble factors**.
- A subset of cytokines called **chemokines** direct the motion of immune cells along their concentration gradient by a process known as **chemotaxis**. Cell motion is inherently **stochastic**.
- Based on these signals, the cells will group together to respond to an infection.

Reaction-Diffusion System for the Soluble Factors

For $i = 1, 2, \dots, N$, let U_i be the concentration of soluble factor i .

$$\begin{aligned} \frac{\partial U_i}{\partial t}(\mathbf{x}, t) &= \text{diffusion} + \text{reaction} + \text{source terms} \\ &= D_i \Delta U_i(\mathbf{x}, t) - \left\{ \sum_{j=1}^N r_{ij} U_j(\mathbf{x}, t) + \lambda_i \right\} U_i(\mathbf{x}, t) \\ &\quad + \sum_{\mu=1}^M J_{i\mu}(t) g_{s_\mu}(\mathbf{x} - \mathbf{x}_\mu(t)), \quad \mathbf{x} \in \Omega \end{aligned}$$

- D_i is the diffusion coefficient for U_i .
- r_{ij} is the rate at which U_i is removed by interaction with U_j .
- $\mathbf{x}_\mu(t)$ is the position of cell μ .
- g_{s_μ} is a smoothly cut-off Gaussian centered at \mathbf{x}_μ .
- $J_{i\mu}(t)$ is the secretion rate of U_i by cell μ .

Stochastic Differential Equations for the Cell Motion

Cell motion is modeled by a Langevin process $X_t = (\mathbf{x}_t, \mathbf{v}_t)$

- Position: $d\mathbf{x}_t = \mathbf{v}_t dt$
- Velocity: $d\mathbf{v}_t = [h(\mathbf{U}(\mathbf{x}_t)) - \gamma\mathbf{v}_t] dt + \sigma(\mathbf{U}(\mathbf{x}_t))\sqrt{\gamma}d\mathbf{W}_t$
- The motion is biased toward the direction of $h(\mathbf{U}(\mathbf{x}_t))$ where

$$h(\mathbf{U}) = \sum_{i=1}^N \frac{\chi_i \nabla U_i}{h_0 + |\nabla U_i|}.$$

Here χ_i is the chemotactic constant that controls how much the drift is influenced by the gradient of soluble factor i .

- The magnitude of the Wiener process depends on a soluble factor concentration by

$$\sigma(U) = \sum_{i=1}^N \sigma_0 \left(1 + \frac{U_i}{U_i^0}\right)^q e^{-\lambda U_i}$$

Operator Splitting

We can split the model into the following three problems for which there are well-known numerical methods:

- 1 The diffusion of the soluble factors,

$$\frac{\partial U_i}{\partial t}(\mathbf{x}, t) = D_i \Delta U_i + \sum_{\mu=1}^n J_{i\mu}(t) g_{s_\mu}(\mathbf{x} - \mathbf{x}_\mu(t)), \quad U_i(0) = u_i.$$

Note: This is now a decoupled set of *linear* PDEs.

- 2 The reactions of the soluble factors,

$$\frac{\partial U_i}{\partial t} = - \left\{ \sum_{j=1}^m r_{ij} U_j + \lambda_i \right\} U_i, \quad U_i(0) = u_i.$$

Note: This is now a system of ODEs.

- 3 The motion of the cells,

$$d\mathbf{x}_t = \mathbf{v}_t dt, \quad \mathbf{x}(0) = \xi,$$

$$d\mathbf{v}_t = [h(\mathbf{u}(\xi)) - \gamma \mathbf{v}_t] dt + \sigma(\mathbf{u}(\xi)) \sqrt{\gamma} d\mathbf{W}_t, \quad \mathbf{v}(0) = \nu.$$

Operator Splitting

For each time step:

- 1 Solve the diffusion using the initial concentrations and cell positions.
- 2 Solve the reaction the using the updated concentrations.
- 3 Move the cells according to the Langevin process using the updated concentrations and initial cell positions.

Important Estimates:

$$\sup_{0 \leq t \leq \Delta t} |\mathbb{E} [\|U(t) - \bar{U}(t)\|_k + |X_t - \bar{X}_t|] | \leq C(\sup |u|, \|u\|_{(k+2)\vee 3})(\Delta t)^2.$$

$$\sup_{0 \leq t \leq \Delta t} \mathbb{E} \left[\|U(t) - \bar{U}(t)\|_k^2 + |X_t - \bar{X}_t|^2 \right]^{\frac{1}{2}} \leq C(\sup |u|, \|u\|_{(k+2)\vee 3})(\Delta t)^{\frac{3}{2}}.$$

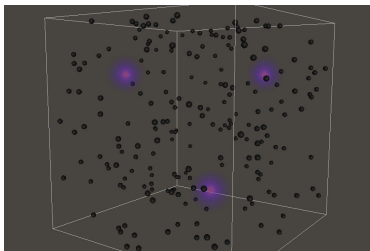
The resulting error using the two estimates above is $\mathcal{O}(\Delta t)$.

Numerical Methods

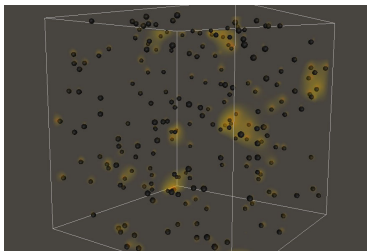
- Diffusion: Backward Euler with Multigrid
- Reaction: Simple ODE method
- Cell Motion: Hold the soluble factor concentration constant in space. Solve the Langevin process exactly.

All three methods are $\mathcal{O}(\Delta t)$.

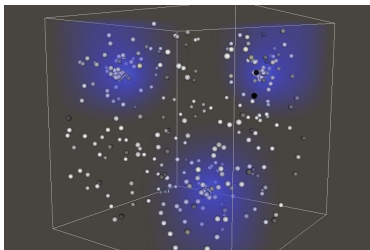
Numerical Simulation - Response to MCP1



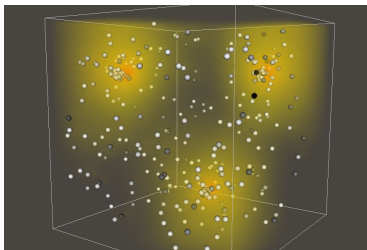
(a) Concentration of MCP1 at $t = 10\text{min}$



(b) Concentration of TNF at $t = 10\text{min}$



(c) Concentration of MCP1 at $t = 120\text{min}$



(d) Concentration of TNF at $t = 120\text{min}$

Acknowledgements

From Duke University:

- William K. Allard, Department of Mathematics
- Thomas B. Kepler, Department of Biostatistics and Bioinformatics
- Cliburn Chan, Department of Biostatistics and Bioinformatics
- Eric Monson, Department of Computer Science