

Image-based modelling using geometric surface partial differential equations (GS-PDEs)

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European Union



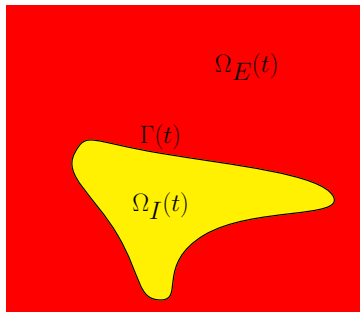
The Leverhulme Trust



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Data-driven coupled bulk-surface and ECM models for cell motility and pattern formation



Aim: To develop a mathematical and computational framework for modelling cell motility and pattern formation

- 1 Data-driven Modelling and Analysis:**
 - **Mechanics:** Viscoelastic, poroelastic, hyperelastic, morphoelastic in the cell interior and ECM, **Geometric Surface PDEs** on the cell surface: *Protrusion, Retraction, Adhesion, Membrane forces such as surface tension and bending rigidity, ...*
 - **Biochemistry:** Polarisation, Receptor-Ligand Dynamics, Spatio-temporal dynamics of RhoGTPases, ...
- 2 Numerical Analysis and HPC Scientific Computing:** Development of accurate, robust and efficient numerical methods for simulating the model equations
- 3 Validation and Model Predictions:** Calibrate and test model predictions with experimental data
- 4 Parameter Identification and Model Selection:** Bayesian and optimal control approaches (useful for guiding data acquisition)

- 1 Biological Motivation: Zebrafish as model organism**
- 2 Whole cell tracking using geometric surface PDEs
- 3 Optimal control of phase fields formulation using geometric PDEs
- 4 Adaptive multigrid method
- 5 Numerical tests and applications
- 6 Funding - Acknowledgement

Whole cell tracking through an optimal control of geometric evolution laws

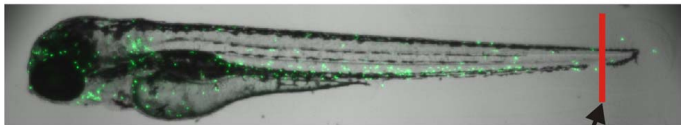
- Yang, F., Venkataraman, C., Gu, S., Styles, V. and Madzvamuse, A., 2022. *Force estimation during cell migration using mathematical modelling*. **Journal of Imaging**, **8**(7), p.199. doi.org/10.3390/jimaging8070199
- Yang, F.W. Venkataraman, C. Styles, V. and Madzvamuse, A. (2017). *A robust and efficient adaptive multi-grid solver for the optimal control of phase field formulations of geometric evolution laws*. **Communications in Computational Physics**. **21**(1):65-92. doi: 10.4208/cicp.240715.080716a
- Yang, F.W. Venkataraman, C. Styles, V. Kutenberger, V. Horn, E. von Guttenberg, Z. and Madzvamuse, A. (2016). *A computational framework for particle and whole cell tracking applied to a real biological dataset*. **Journal of Biomechanics**. ISSN 0021-9290
- Blazakis, K.N. Madzvamuse, A. Reyes-Aldasoro, C.C. Styles, V. and Venkataraman, C. (2015) *Whole cell tracking through the optimal control of geometric evolution laws* **Journal of Computational Physics**, **297**. pp. 495-514. ISSN 0021-9991
- Yang, F.W. Venkataraman, C, Styles, V and Madzvamuse, A. (2015) *A parallel and adaptive multi-grid solver for the solutions of the optimal control of geometric evolution laws in two and three dimensions*. In: **4th International Conference on Computational and Mathematical Biomedical Engineering - CMBE2015**. P. Nithiarasu and E. Budyn (Eds.) 29 June-1 July 2015, Paris, France.
- Blazakis, K.N. Reyes-Aldasoro, C C, Styles, V, Venkataraman, C and Madzvamuse, A. (2015) *An optimal control approach to cell tracking*. In: Louis, Alfred K, Arridge, Simon and Rundell, Bill (eds.) **Proceedings of the Inverse Problems from Theory to Applications Conference (IPTA2014)**. IOP Publishing Ltd , pp. 74-77. ISBN 9780750311069

Biological Motivation: Zebrafish as model organism



- Zebrafish are transparent - observe morphological changes during development.
- Use the GFP to label individual cells, organs or even organelles.
- Embryo development in normal/abnormal situations (mutated or drugs)
- Zebrafish are more closely related to humans than invertebrate models such as the worm *C. elegans* and the fly *D. Melanogaster*,

Zebrafish larva three days post-fertilisation



Tail transection

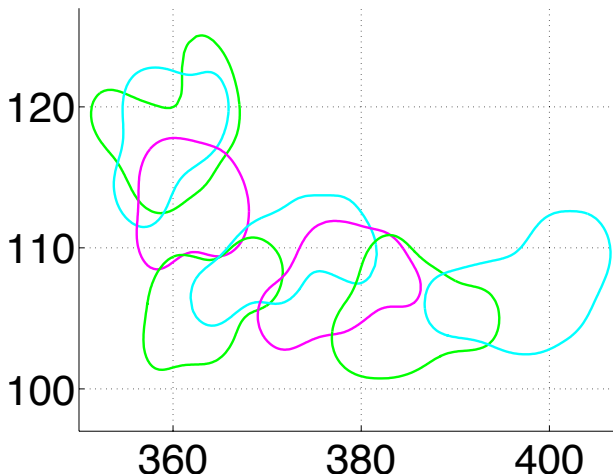
Neutrophil cell migration during wound healing process

Cell migration during wound healing process

3-D reconstruction of the cell migration

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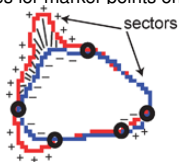
Discrete sequence of cross-sections of 3D image datasets



- Can we model "optimally" the evolution of the cell from one image to the next?
- Reconstruct dynamic evolution of the cell from static images?

Cell Tracking

- Reconstruct a dynamic (2D+t) or (3D+t) model from static imaging data (2D or 3D).
 - **Particle tracking (e.g. Agent Based Tracking)**: recover trajectories, “connect the dots”, enables the computation of many motility related statistics such as velocities, persistence lengths, MSD, etc.
 - **Whole cell tracking**: recover morphologies, allows investigation of the dynamics of geometric features (surface area, volume, curvature, aspect ratio, ...)
- For example
 - **Particle tracking**: reconstruct (often only centroid) trajectories by linear interpolation
 - **Whole Cell tracking**: Level set and electrostatic based methods that generate trajectories for marker points on the membrane



[Tyson, Epstein, Anderson, and Bretschneider, 2010]

- **Our approach**: fitting a mathematical model to static imaging data.

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Whole Cell Tracking

Basic Idea: Reconstruct (dynamic) whole cell morphologies from static imaging data.

- Majority of existing approaches (level set, electrostatics, . . .) incorporate only geometric considerations equidistribution of vertices,
- Our approach: **Fit models for the cell evolution to imaging data.** Focus is on geometric evolution law based models for the motion of the cell membrane as considered in [Barreira, Elliott, and Madzvamuse, 2011; Elliott, Stinner, and Venkataraman, 2012; Marth and Voigt, 2013; Neilson, Veltman, van Haastert, Webb, Mackenzie, and Insall, 2011; Shao, Levine, and Rappel, 2012].
- For many cell tracking scenarios, only the membrane location is available with no other fluorescence data provided, hence we focus on this most basic setting.

Geometric Evolution Law Model

$$\mathbf{V}(\vec{x}, t)\mathbf{n}(\vec{x}, t) = (\eta(\vec{x}, t) + \lambda(t) - \sigma H(\vec{x}, t))\mathbf{n}(\vec{x}, t) \quad (\vec{x}, t) \in (\Gamma(t), [0, T])$$

Velocity = **Forcing** + Vol. Cons. + Regularisation

The problem is to find $\eta(\vec{x}, t)$ such that the solution to the model and the image data are “close”.

Problem is in effect the optimal control of the sharp interface formulation of a geometric evolution law for which no adequate theory is available yet (how to define a smooth notion of “closeness”, regularity for $\eta(\vec{x}, t)$, etc).

Diffuse Interface Formulation

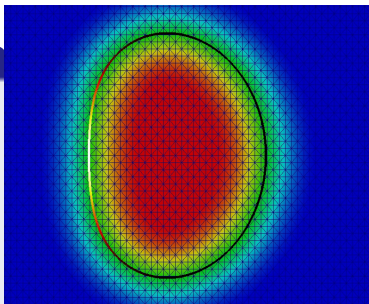
From Sharp to Diffuse Interface

Given a surface

$$\Gamma = \{\vec{x} \in \Omega \mid d_\Gamma(x) = 0\},$$

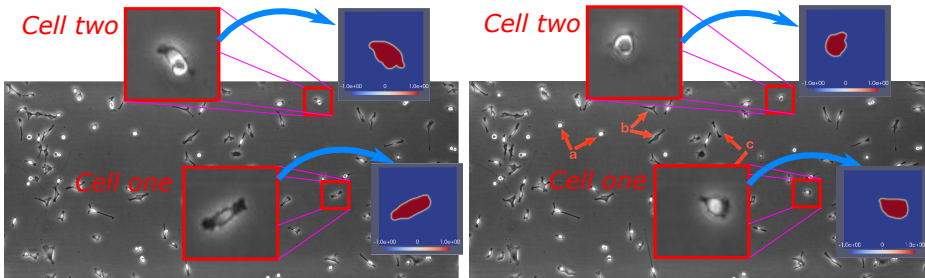
where d_Γ is the signed distance function to Γ , compute $\varphi : \Omega \rightarrow \mathbb{R}$ such that

$$\varphi_\epsilon(\vec{x}) = \begin{cases} 1 & \text{if } d_\Gamma(\vec{x}) > \epsilon \\ \sin\left(\frac{\pi d_\Gamma(\vec{x})}{2\epsilon}\right) & \text{if } |d_\Gamma(\vec{x})| < \epsilon \\ -1 & \text{if } d_\Gamma(\vec{x}) < -\epsilon \end{cases}$$



Alternatively one may simply work with (high contrast) raw data without further segmentation of the cell outline (with a few steps of de-noising).

Individual cells



Diffuse Interface Formulation

Adopting a phase field (diffuse interface) formulation \rightsquigarrow Allen-Cahn with forcing.

Allen-Cahn with forcing

$$\varepsilon \partial_t \varphi(\mathbf{x}, t) = \varepsilon \Delta \varphi(\mathbf{x}, t) - \frac{1}{\varepsilon} G'(\varphi(\mathbf{x}, t)) + c_w \eta(\mathbf{x}, t) + \lambda(t) \quad (\mathbf{x}, t) \in \Omega \times [0, T] \quad (1)$$

where $G(\varphi) = \frac{1}{4}(\varphi^2 - 1)^2$ is a double well potential with minima at ± 1 .

Volume constraint interpreted as constraint on the mass, i.e., $\int_{\Omega} \varphi$.

Example: fitting to a single image φ_{des} (using an observation at a previous time as initial data).

PDE constrained optimisation problem

$$\min J(\varphi, \eta) = \frac{1}{2} \int_{\Omega} (\varphi(\mathbf{x}, T) - \varphi_{des}(\mathbf{x}))^2 d\mathbf{x} + \frac{\gamma}{2} \int_0^T \int_{\Omega} \eta^2 d\mathbf{x} dt$$

subject to (1).

Following [Haußer, Rasche, and Voigt, 2010; Haußer, Janssen, and Voigt, 2012] we formally derive the first order optimality conditions and propose an adjoint based solution method for the minimisation problem.

Optimality conditions [Tröltzsch, 2010]

Introducing the Lagrange multiplier (adjoint state) p , we define the Lagrangian functional

$$\begin{aligned} \mathcal{L}(\varphi, \eta, p) = J(\varphi, \eta) - \int_0^T \int_{\Omega} & \left(\varepsilon \partial_t \varphi(\mathbf{x}, t) - \varepsilon \Delta \varphi(\mathbf{x}, t) \right. \\ & \left. + \frac{1}{\varepsilon} G'(\varphi(\mathbf{x}, t)) + c_w \eta(\mathbf{x}, t) + \lambda(t) \right) p(\mathbf{x}, t) d\Omega dt. \end{aligned}$$

Requiring stationarity of the Lagrangian with respect to the adjoint state yields the state (forward) equation and requiring stationarity of the Lagrangian, at the optimal control η^* and associated optimal state φ^* , with respect to the state and the control, yields the (formal) first order optimality conditions

$$\begin{aligned} \delta_{\varphi} \mathcal{L}(\varphi^*, \eta^*, p) \varphi &= 0, \forall \varphi : \varphi(\mathbf{x}, 0) = 0, \\ \delta_{\eta} \mathcal{L}(\varphi^*, \eta^*, p) \eta &= 0, \forall \eta. \end{aligned}$$

Optimality conditions [Tröltzsch, 2010]

Seeking optimality conditions yield the adjoint equation, which is the following linear parabolic PDE (posed backwards in time) for the adjoint state p ,

$$\begin{cases} \partial_t p(\mathbf{x}, t) &= -\Delta p(\mathbf{x}, t) + \frac{1}{\epsilon^2} G''(\varphi(\mathbf{x}, t)) p(\mathbf{x}, t) & \text{in } \Omega \times (T, 0], \\ \nabla p \cdot \nu &= 0 & \text{on } \partial\Omega \times (T, 0], \\ p(\mathbf{x}, T) &= \varphi(\mathbf{x}, T) - \varphi_{obs}(\mathbf{x}) & \text{in } \Omega, \end{cases}$$

and together with the Riesz representation theorem yields the optimality condition for the control [Tröltzsch, 2010]

$$\delta_\eta \mathcal{L}(\varphi^*, \eta^*, p) = \theta \eta^* + \frac{1}{\epsilon} p = 0.$$

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HPC Scientific Computing

- Cell-centred 2nd order finite difference method (FDM) on rectangular grids
- PARAMESH library for mesh generation and parallelism
- Dynamic load-balancing for adaptive mesh refinement (AMR)
- Fully implicit 2nd order backward differentiation formula (BDF2)
- Geometric nonlinear multigrid solution method using full approximation scheme (FAS) and multi-level adaptive technique (MLAT)
- Point-wise Newton linearisation with Gauss-Seidel iteration
- Full weighting restriction and multi-linear interpolation

Numerical challenges

- Each η iteration covers all time steps (both forward and backward)
- Memory requirement (let's consider double precision and 100 time steps)
 - 2-D: 512^2 requires 0.4 gigabytes
 - 3-D: 512^3 requires 215 gigabytes

Adaptive iterative update for the optimal control

- We denote a superscript ℓ for the η iteration, and at $\ell = 0$, we take

$$\eta^{\ell=0} = 0 \text{ on } \Omega \times [0, T)$$

as our initial guess for the control.

- A gradient-based iterative update of the control, following the steepest descent approach, and the update is given by

$$\eta^{\ell+1} = \eta^\ell - \alpha \left(\theta \eta^\ell + \frac{1}{\epsilon} p^\ell \right), \text{ on } \Omega \times [0, T),$$

where $\ell + 1$ denotes the next η iteration and ℓ indicates the current η iteration.

- The whole procedure is repeated until the objective function J satisfies some pre-defined tolerances.

Adaptive α

Algorithm

- 1 **While** the difference between consecutive J s is still large or J has not reached below a pre-defined tolerance **do**
- 2 Solve the forward Allen-Cahn equation in $\Omega \times (0, T]$
- 3 Compute the objective functional J^ℓ
- 4 **if** $J^\ell > J^{\ell-1}$ **and** $\ell > 0$ **then**
 $\alpha = \max(\alpha \times \mathcal{P}_l, \alpha_{min})$
 restart = **TRUE**
- else if** $J^\ell < J^{\ell-1}$ **and** $\ell > 0$ **then**
 $\alpha = \alpha \times \mathcal{P}_u$
 restart = **FALSE**
- end if**
- 5 **if** restart == **FALSE** **then**
 Solve the backward adjoint equation in $\Omega \times [T, 0)$
 Backup the current η
 Compute the next η using α
 Continue to the next η iteration
- else**
 Compute a new η using the latest backup with α
 Restart the current η iteration
- end if**
- 6 **End**

Space and time discretisations of the forward and adjoint equations

- Spatial discretisation – Finite differences (2D and 3D)
- Time-stepping:
 - 1 BDF1 (also known as backward Euler method) is employed for the very first time step.
 - 2 BDF2 is employed for the remaining time steps

$$\epsilon \frac{\varphi_{i,j,k}^{n+1,\ell+1} - \frac{4}{3}\varphi_{i,j,k}^{n,\ell+1} + \frac{1}{3}\varphi_{i,j,k}^{n-1,\ell+1}}{\tau} = \frac{2\epsilon}{3} D(\varphi_{i,j,k}^{n+1,\ell+1}) - \frac{2\left(-\varphi_{i,j,k}^{n+1,\ell+1} + \left(\varphi_{i,j,k}^{n+1,\ell+1}\right)^3\right)}{3\epsilon} + \frac{2\eta_{i,j,k}^{n+1,\ell+1}}{3} + \frac{2\lambda^{n+1}}{3},$$

- $\ell + 1$ denotes the current η iteration, $n + 1$, n and $n - 1$ indicate solutions from current, previous and the one before the previous time steps, respectively.
- We denote the 3-D Laplacian operator D as

$$D(\varphi_{i,j,k}) = \frac{\varphi_{i+1,j,k} + \varphi_{i-1,j,k} + \varphi_{i,j+1,k} + \varphi_{i,j-1,k} + \varphi_{i,j,k+1} + \varphi_{i,j,k-1} - 6\varphi_{i,j,k}}{h^2}.$$

Space and time discretisations of the forward and adjoint equations

- Within each time step, while solving for the solution of the above systems, we are also required to satisfy a given mass constraint.
- Iteratively determine the time-dependent, spatially-uniform volume constraint λ for the imposed mass constraint
- The Allen-Cahn system has to be solved multiple times, until a stopping criterion for λ is met.
- We denote this λ iteration using a superscript Λ , and its update follows the multi-step approach which is given as

$$\lambda^{n+1, \Lambda+1} = \lambda^{n+1, \Lambda} + \frac{(\lambda^{n+1, \Lambda} - \lambda^{n+1, \Lambda-1}) \left[M_{\varphi}^{n+1} - \int_{\Omega} \varphi^{n+1, \Lambda} \right]}{\left(\int_{\Omega} \varphi^{n+1, \Lambda} - \int_{\Omega} \varphi^{n+1, \Lambda-1} \right)}, \quad \text{for } \Lambda > 1,$$

where M_{φ} is defined earlier, $\Lambda + 1$, Λ and $\Lambda - 1$ indicate values of λ from current, previous and the one before the previous λ iterations, respectively.

- Initial guesses

$$\lambda^{\Lambda=0} = -\frac{2\epsilon}{\tau} + 1, \quad \lambda^{\Lambda=1} = \frac{2\epsilon}{\tau} - 1.$$

- The stopping criterion used here is based upon the difference between consecutive values of λ .
- Providing a tolerance tol_{λ} , we consider the algorithm to have converged when $|\lambda^{n+1, \Lambda+1} - \lambda^{n+1, \Lambda}| < tol_{\lambda}$.

Approximation of the Optimal Control Problem

We propose a simple steepest-descent based iterative algorithm for the solution of the OC problem.

Algorithm

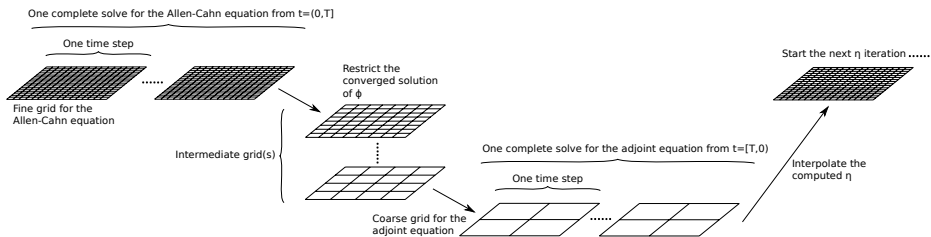
While tolerance not met:

- 1 Compute solution φ^n corresponding to control η^n . Mass constraint computed following [Blowey and Elliott, 1993].
- 2 Compute objective functional $J(\varphi^n, \eta^n)$ check if tolerance met.
- 3 Compute adjoint state p^n (for efficient computation of the gradient of J) **requires the φ^n** .
- 4 Update control according to gradient (i.e., compute η^{n+1}).
- 5 $n + 1 \rightarrow n$.

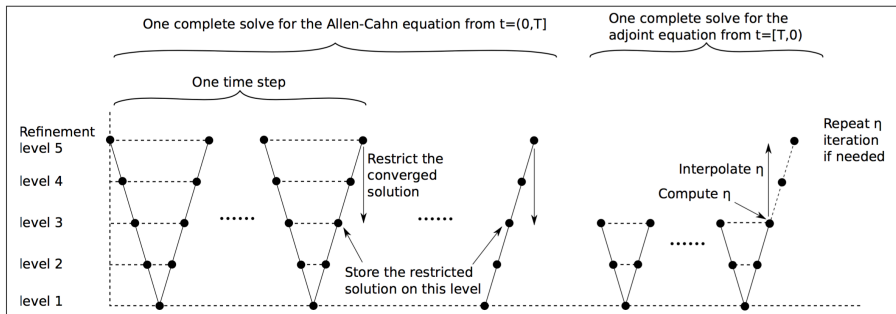
Efficiency

- 1 Adaptive parallel multigrid based solution method for the forward and adjoint problems.
- 2 “Two-grid” strategy: **adaptive** grid for forward problem, **coarse uniform** grid for (linear) adjoint problem. \rightsquigarrow **Massive memory savings**.
- 3 Adaptive step-size selection for control update.

Multigrid: Two grid solution strategy



Two-grid scheme within multigrid V-cycles

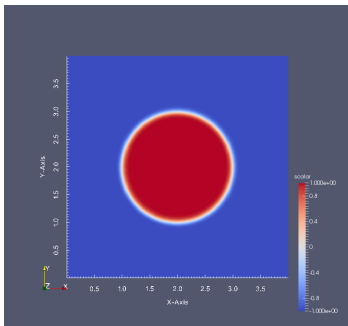


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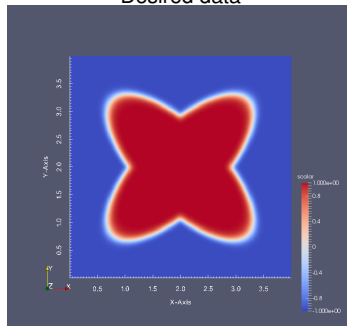
Circle to ellipse

A circle becomes two ellipses.

Initial data



Desired data



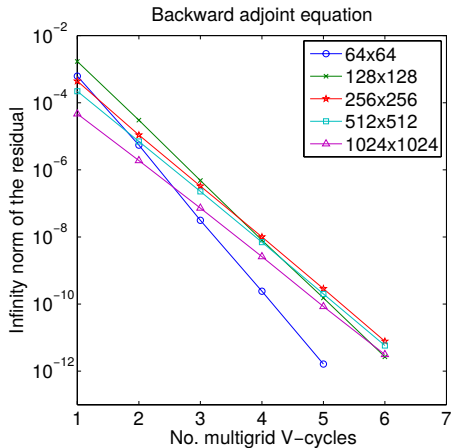
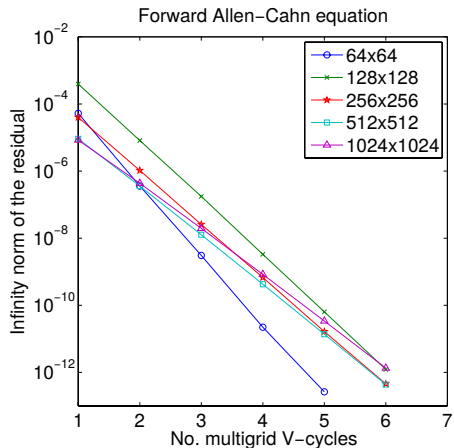
Geometric evolution of a circle to an ellipse

Convergence

$L_2(\Omega)$ error for φ			
	$m = t_1$	$m = t_2$	$m = t_3$
$d_{64^2}^m$	3.4264×10^{-2}	4.9226×10^{-2}	6.8561×10^{-2}
$d_{128^2}^m$	1.9721×10^{-2}	4.8058×10^{-2}	6.0397×10^{-2}
$d_{256^2}^m$	8.1793×10^{-3}	2.2300×10^{-2}	3.4557×10^{-2}
$d_{512^2}^m$	2.7850×10^{-3}	8.0407×10^{-3}	1.3192×10^{-2}
$L_2(\Omega)$ error for adjoint p			
	$m = t_1$	$m = t_2$	$m = t_3$
$d_{64^2}^m$	1.6773×10^{-2}	1.8048×10^{-2}	4.9344×10^{-2}
$d_{128^2}^m$	9.9721×10^{-3}	1.0158×10^{-2}	3.1554×10^{-2}
$d_{256^2}^m$	7.9290×10^{-3}	8.5311×10^{-3}	2.2551×10^{-2}
$d_{512^2}^m$	6.5082×10^{-3}	7.5551×10^{-3}	1.4901×10^{-2}
$L_2(\Omega)$ error for η			
	$m = t_1$	$m = t_2$	$m = t_3$
$d_{64^2}^m$	1.6976×10^{-1}	2.0752×10^{-1}	7.5240×10^{-1}
$d_{128^2}^m$	1.1923×10^{-1}	1.5793×10^{-1}	6.2554×10^{-1}
$d_{256^2}^m$	8.5093×10^{-2}	1.0601×10^{-1}	5.3023×10^{-1}
$d_{512^2}^m$	3.2344×10^{-2}	3.9359×10^{-2}	2.6302×10^{-1}

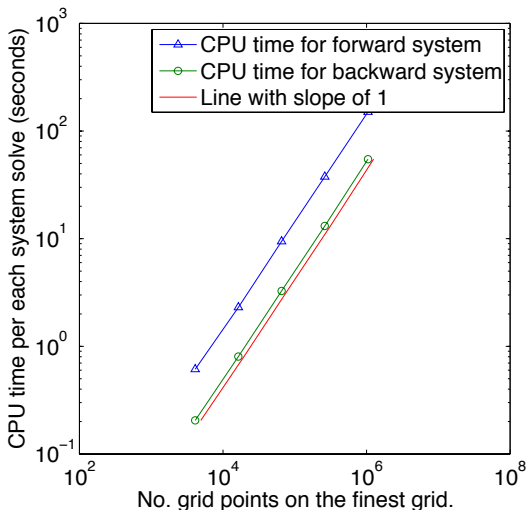
The convergence tests for the solutions of φ , adjoint p and η .

Multigrid convergence rates



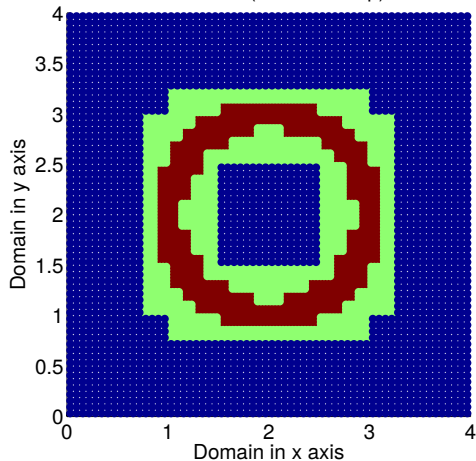
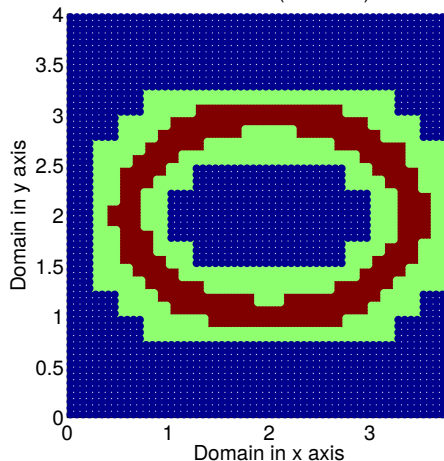
The multigrid convergence rates for the forward Allen-Cahn and backward adjoint equations.

Linear complexity of the multigrid method



A log-log plot to illustrate the linear complexity of our multigrid solver. For comparisons, a line of slope 1 is included.

Dynamic Adaptive Mesh Refinement

AMR at t_1 (first time step)AMR at $t=T$ (end time)

Two colour plots show the dynamic AMR in our solver. The blue region shows the 64^4 grid; light green region indicates the 128^2 grid; and finally red region illustrates the finest 256^2 grid. The colour version of this figure is [online](#).

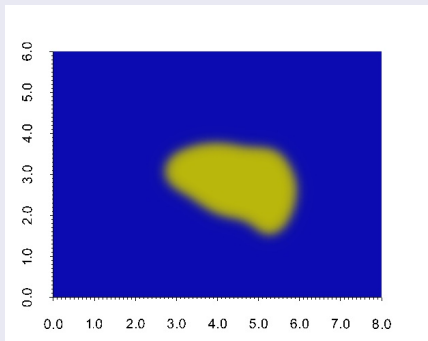
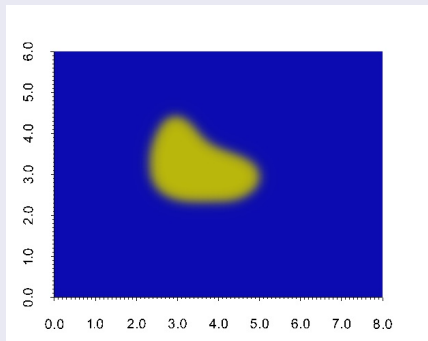
Convergence: Two-grid strategy

$L_2(\Omega)$ error for φ			
	$m = t_1$	$m = t_2$	$m = t_3$
$d_{128^2}^m$	1.2327×10^{-2}	2.6664×10^{-2}	3.8450×10^{-2}
$d_{256^2-64^2}^m$	8.6270×10^{-3}	1.6895×10^{-2}	3.2925×10^{-2}
$L_2(\Omega)$ error for adjoint p			
	$m = t_1$	$m = t_2$	$m = t_3$
$d_{128^2}^m$	9.2021×10^{-3}	9.7122×10^{-3}	3.0233×10^{-2}
$d_{256^2-64^2}^m$	1.0004×10^{-2}	1.4886×10^{-2}	2.7392×10^{-2}
$L_2(\Omega)$ error for η			
	$m = t_1$	$m = t_2$	$m = t_3$
$d_{128^2}^m$	9.7196×10^{-2}	1.2153×10^{-1}	5.3632×10^{-1}
$d_{256^2-64^2}^m$	7.7932×10^{-2}	1.4930×10^{-2}	5.9167×10^{-1}

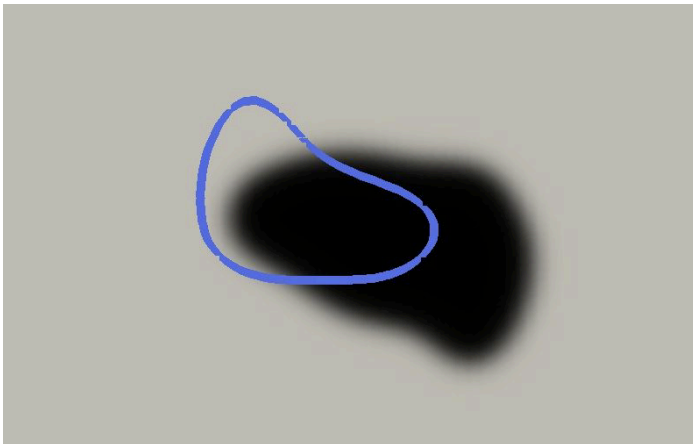
Comparisons of errors between an adaptive two-grid simulation ($256^2 - 64^2$) with adaptive mesh refinement and a standard 128^2 uniform grid simulation.

Single cell imaging data

Initial and target Neutrophil shapes



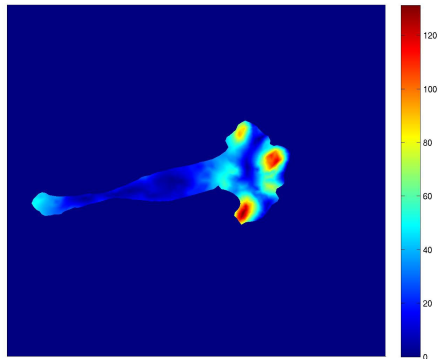
Single cell imaging data



Zero level set of the computed surface shaded by the value of η and phase field representation of the target surface.

Cell one video

Cancer cell migration



Data from [Peschetola, Laurent, Duperray, Michel, Ambrosi, Preziosi, and Verdier, 2013].

Keratocyte migration

Tracking topological changes during cell migration

For a smooth, oriented, compact (i.e., no boundary) surface Γ , the Euler number of a Riemann manifold (a topological invariant) is given by [Allendoerfer, 1940]

$$\chi = \frac{1}{2\pi} \int_{\Gamma} K d\sigma$$

where K is the Gauss curvature of the surface.

2D case [Du, Liu, and Wang, 2005]

For $-1 < b < 0 < a < 1$

$$\chi_{\varepsilon} = \frac{1}{2\pi(a-b)} \int_{\Omega(a,b)} \left(-\Delta\varphi + \frac{\nabla|\nabla\varphi|^2 \cdot \nabla\varphi}{2|\nabla\varphi|^2} \right) d\vec{x} \approx \# \text{ of cells}$$

Future Work

Penalise changes in topology [Dondl, Mugnai, and Röger, 2011, 2014].

Application to quantifying cell proliferation rates: The MDCK cell line

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- 5 Numerical tests and applications
- 6 Funding - Acknowledgement**

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