

# Attraction and Repulsion in Biological Tissues: Challenges for Models, Analysis, and Numerics

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## Overview

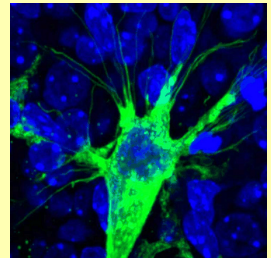
- ▶ Cell attraction and repulsion: some biological background.
- ▶ The nonlocal PDE model and its application.
- ▶ Analytical results.
- ▶ Derivation from a position-jump model.
- ▶ The need for efficient numerics.
  - ▶ Approximation in a periodic setting.
  - ▶ Generalizations and extensions
- ▶ Summary and outlook.

# Cell contact and response

- ▶ Cells can communicate via direct contacts, e.g. membrane-membrane molecular binding.
- ▶ Contacts also occur at long distances, up to 50 cell diameters, via cell protrusions. →
- ▶ Reality is full of detail: cells are complex, morphing objects with a lot of structure and chemistry.
- ▶ Direct contact can lead to many responses, such as movement.

Cells *explore* their surrounding in search of contact sites.

Filopodia (green) of endothelial cell ↓



[Gerhardt et al., J. Cell Biol. (161), '03]

# Cell contact and response

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Focus here: **Direct contacts** between cells

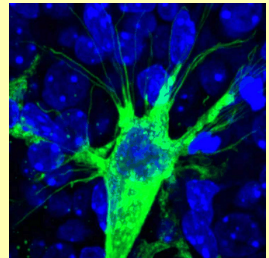
as, e.g., in cell-cell adhesion or contact inhibition.

Not considered: **Indirect contacts** between cells

as, e.g., mediated by diffusible chemical signalling (chemotaxis).

Cells *explore* their surrounding in search of contact sites.

Filopodia (green) of endothelial cell ↓

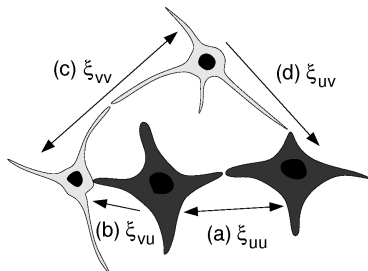


[Gerhardt et al., J. Cell Biol. (161), '03]

Instructing others to move is fundamental for many animal/cellular populations.

## Commands

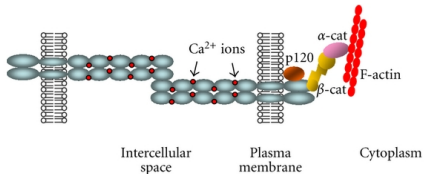
- ▶ are transmitted by contact over variable distances (short or long range),
- ▶ effect cells of the same (homotypic) or different (heterotypic) type,
- ▶ give rise to an attractive or repelling response.



These mechanisms can have a significant impact on the organisation of a tissue.

# Example 1: Cell adhesion

## Molecular aspects and applications



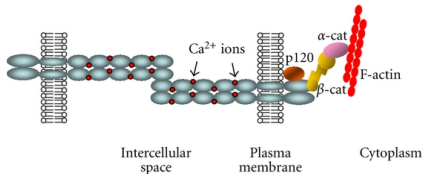
[Gama&Schmitt, Veterinary Medicine International, '12]

Adhesion [latin *adhaesio*] of cells in the body determined by expression and regulation of **cell adhesion molecules**

- ▶ Cadherines (cell-cell adhesion)
- ▶ Integrines (cell-matrix adhesion)
- ▶ ...a few others.

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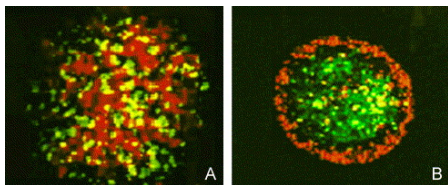
Adhesion important for **tissue integrity** and **cell migration!**

Selected applications:

- ▶ Embryonic development: cells adhere selectively to each other and sort out to form tissue and organs.
- ▶ Cancer invasion: modified adhesive properties of cancer cells are implicated as an important factor.

# Example 1: Cell adhesion

## Cell sorting

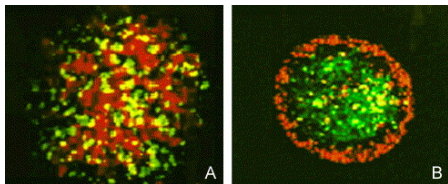


[Foty & Steinberg, Dev. Biol. (278), '05]

2 cell types, differing in number of cadherin molecules on their cell surface only.

Cell type with larger number sorts to the core of the cell pellet.

## Example 1: Cell adhesion Cell sorting



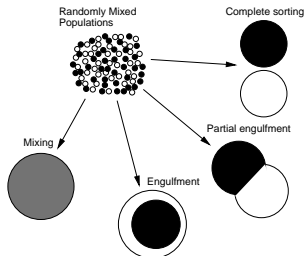
[Foty & Steinberg, Dev. Biol. (278), '05]

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### Differential Adhesion Hypothesis (Steinberg)

A mixture of two cell types sorts always to the same final configuration, independent of its initial distribution. This final configuration depends solely on the adhesive properties (self- and cross-adhesion parameters) of the cell types.



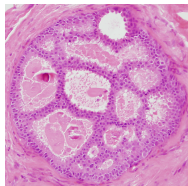
[G. & Painter, '10]



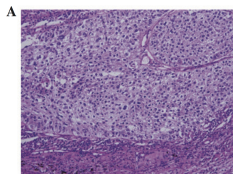
## Example 1: Cell adhesion

### Cancer invasion

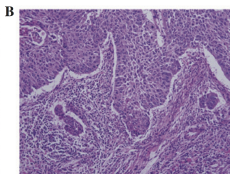
- ▶ Invasion is the process of extension of the cancer into surrounding tissue.
- ▶ It is part of the metastatic cascade, giving cancer its deadly characteristic.
- ▶ Modulation of adhesive properties of cancer cells is one of the hallmarks of **cancer** [Hanahan & Weinberg, '00 & '11]



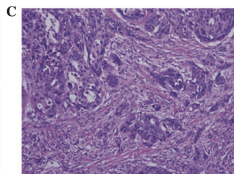
ductal carcinoma in  
situ, intermediate grade,  
[[www.breastpathology.info](http://www.breastpathology.info)]



INF a



INF b



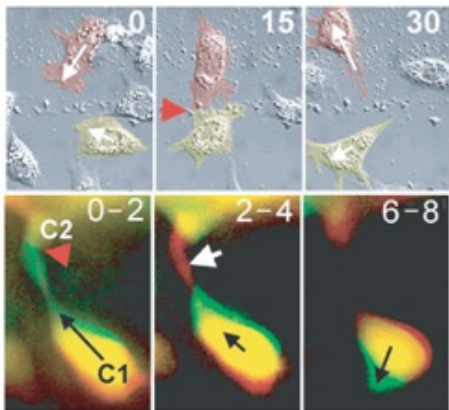
INF c

lung squamous cell carcinoma,  
types (a) to (c) of tumour infiltrative pattern,  
[Masuda et al., '12]

## Example 2: Contact repulsion

### Contact inhibition in neural crest cell migration

Certain cells “repel” each other on contact, a process known as “contact inhibition of locomotion” [Abercrombie & Heaysman, *Exp Cell Res*, 5, '50], [Abercrombie, *Nature*, 281, '79].



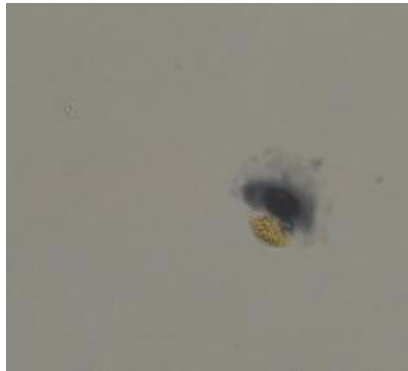
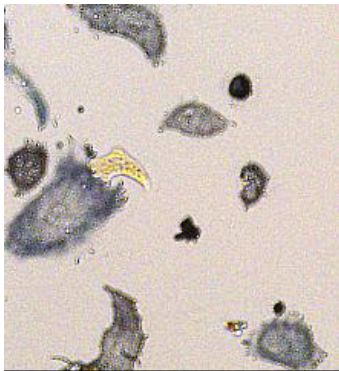
Contact inhibition of locomotion controls neural crest cell directional migration

[Carmona-Fontaine et al., *Nature*, 456, '08]

## Example 3: Mixed interaction Zebrafish pigmentation pattern

Certain heterogeneous populations: attracting and repelling interactions observed.

“run-and-chase” behaviour (xanthophores chase melanophores)



[Yamanaka & Kondo, PNAS, '14]

# Modelling of attraction and repulsion

## Focus on cell adhesion



Two general classes of models

- ▶ individual cell based (discrete) models
  - ↪ dynamics of individual cells,
- ▶ continuous models
  - ↪ dynamics of population level behaviour.
    - ▶ Cells represented through their density at the tissue level.
    - ▶ Cellular scale events captured in model parametrisation.

# Modelling of attraction and repulsion

## Focus on cell adhesion

### Examples of continuous modelling approaches

- ▶ Cell-matrix adhesion:
  - ▶ haptotactic migration, modelled by advective type flux term (cf. chemotaxis).
- ▶ Cell-cell adhesion: Is problematic!
  - ▶ *Some* aspects of adhesion captured by density-dependent cell motility coefficients.
  - ▶ Direct incorporation of surface tension (e.g. Byrne, Chaplain, Lowengrub, Cristini,...)
    - ↔ Cahn-Hilliard type PDE models arising from expansion of nonlocal terms.
  - ▶ **Nonlocal PDE model:** [Armstrong, Painter & Sherratt, J Theor Biol, 243, '06].  
Had and has a substantial influence on the subject with  $\approx 140$  citations to date!

# The Armstrong-Painter-Sherratt model

## Underlying idea

In a fluid (extracellular space) with dynamic viscosity  $\eta$  ...

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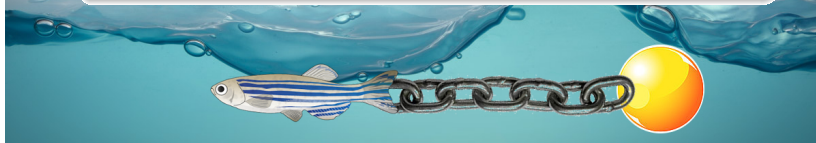
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Given by Stokes's law:  $F = 6\pi\eta\hat{R}v$  or  $v = \frac{1}{6\pi\eta\hat{R}}F$



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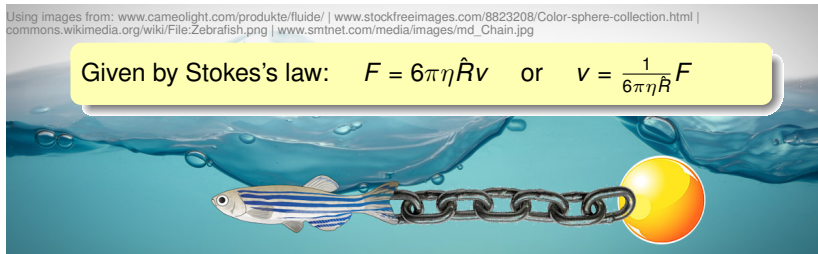
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Assume cells with density  $u(t, x)$  move with velocity  $v(t, x)$  due to adhesion.

$$\rightsquigarrow \text{flux of cells} \quad u(t, x)v(t, x) = u(t, x) \frac{1}{6\pi\eta\hat{R}} F(t, x),$$

where  $F(t, x)$  is now the net force (due to adhesion) acting on a cell at  $(t, x)$ .

# The Armstrong-Painter-Sherratt model

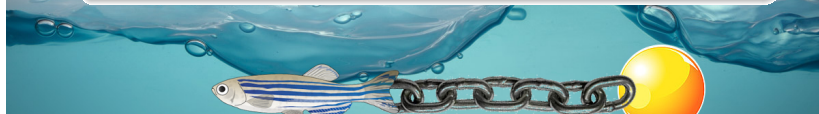
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Slight difficulty:

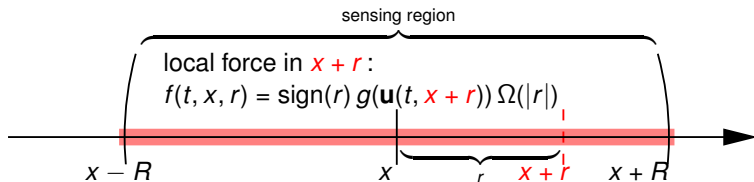
Assume **Stokes's law is for a single cell, not for interacting cell populations.**  
... but let's go ahead!

where  $F(t, x)$  is now the net force (due to adhesion) acting on a cell at  $(t, x)$ .

# The Armstrong-Painter-Sherratt model

## Net force due to adhesion

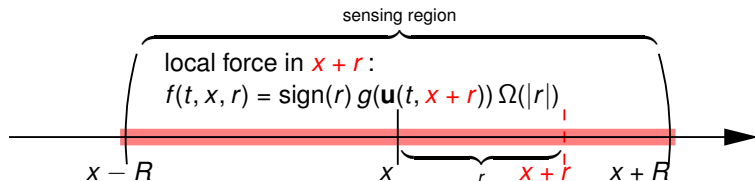
The net force  $F$  in  $(t, x)$  is the sum of “local” forces:  $F(t, x) := \int_{-R}^R f(t, x, r) dr$



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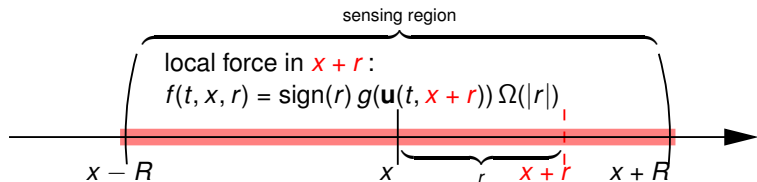


$\text{sign}(r) = \frac{r}{|r|}$  force direction between the cell at  $x$  and those at  $x+r$ .

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$\text{sign}(r) = \frac{r}{|r|}$       force direction between the cell at  $x$  and those at  $x+r$ .

$g(\mathbf{u})$       force magnitude between the cell at  $x$  and those at  $x+r$ :

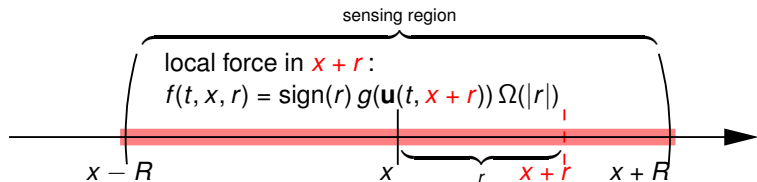
One cell type:  $g(u) = Cu$  or  $g(u) = Cu(1-u)^+$ ;

Two cell types:  $g_1(u_1, u_2) = (C_{11}u_1 + C_{12}u_2)(1-u_1-u_2)^+$ .

# The Armstrong-Painter-Sherratt model

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$\Omega(|r|) \geq 0$       radial dependency: e.g. constant or decaying on  $[0, R]$ .



# The Armstrong-Painter-Sherratt model

## The nonlocal PDE model

Results in nonlocal **adhesion velocity**  $\mathcal{A}_i\{\mathbf{u}(t, \cdot)\}(x)$  for cell type  $i$

$$\mathcal{A}_i\{\mathbf{u}(t, \cdot)\}(x) := \frac{1}{\Phi R} \int_{\mathcal{S}} \frac{r}{|r|} g_i(\mathbf{u}(t, x+r)) \Omega(|r|) dr,$$

where  $\mathcal{S}$  now denotes the sensing region relative to but independent of  $x$ .

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Conservation of mass framework leads to nonlocal PDE model

$$\partial_t u_i = -\nabla \cdot (-D_i \nabla u_i + u_i \mathcal{A}_i\{\mathbf{u}(t, \cdot)\})$$

where

- ▶  $-D_i \nabla u_i$  is a Fickian diffusive flux and
- ▶  $+u_i \mathcal{A}_i\{\mathbf{u}(t, \cdot)\}$  is the flux due to adhesion.

The system is complemented with initial and (periodic) boundary conditions.

# Application: Cell sorting

[G. & Painter, '10]

- ▶ Two cell populations with densities  $u_A(t, x)$  and  $u_B(t, x)$  for  $x \in (0, 10)^2$ .
- ▶ Model equations

$$\partial_t u_A = -\nabla \cdot (-D\nabla u_A + u_A \mathcal{A}_A\{\mathbf{u}(t, \cdot)\})$$

$$\partial_t u_B = -\nabla \cdot (-D\nabla u_B + u_B \mathcal{A}_B\{\mathbf{u}(t, \cdot)\})$$

where  $\mathcal{S}$  is the unit circle,  $D = R = \Phi = 1$ ,  $\Omega \equiv 1$ , and

$$g_A(\mathbf{u}) = (C_{AA}u_A + C_{AB}u_B)(1 - u_A - u_B)^+$$

$$g_B(\mathbf{u}) = (C_{BA}u_A + C_{BB}u_B)(1 - u_A - u_B)^+$$

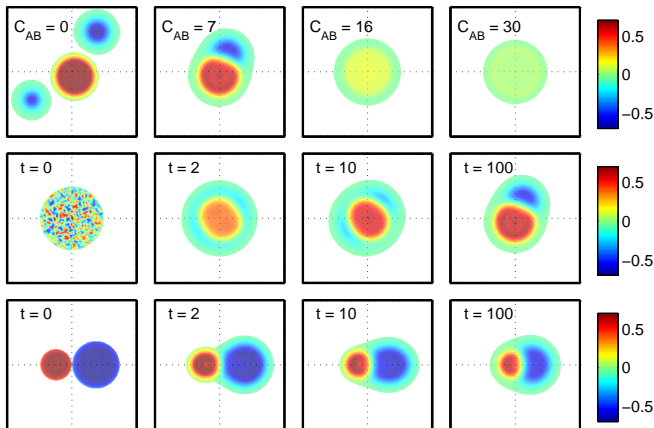
with self-adhesion coefficients  $C_{AA}$  and  $C_{BB}$   
and cross-adhesion coefficients  $C_{AB} = C_{BA}$ .

- ▶ Equations complemented with initial and periodic boundary conditions.

# Application: Cell sorting

[G. & Painter, '10]

→→→ increasing cross-adhesion →→→



Plots show the  
difference  $u_A - u_B$

Parameter:

$$C_{AA} = 30$$

$$C_{BB} = 15$$

Initial conditions:

← mixed

← separated

[G. & Painter, '10]

# Application: cancer invasion

[Domschke et al., J Theor Biol, 361, '14], also [G. & Chaplain, J Theor Biol, 250, '08]



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Model with three time- and space-dependent variables:

- ▶ the cancer cell density,  $c : \mathcal{I}_T \times \mathcal{D} \rightarrow \mathbb{R}$ ,
- ▶ the extracellular matrix (ECM) density,  $v : \mathcal{I}_T \times \mathcal{D} \rightarrow \mathbb{R}$ , and
- ▶ the matrix-degrading enzyme (MDE) concentration,  $m : \mathcal{I}_T \times \mathcal{D} \rightarrow \mathbb{R}$ .

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- ▶ the matrix-degrading enzyme (MDE) concentration,  $m : \mathcal{I}_T \times \mathcal{D} \rightarrow \mathbb{R}$ .

$$\begin{aligned}\frac{\partial c}{\partial t} &= -\nabla \cdot [-D_1 \nabla c + c \mathcal{A}\{t, \mathbf{u}(t, \cdot)\}] + \mu_{1,1} c (1 - u - v), \\ \frac{\partial v}{\partial t} &= -\gamma m v + \mu_2 (1 - u - v)^+, \\ \frac{\partial m}{\partial t} &= -\nabla \cdot [-D_3 \nabla m] + \alpha_1 c - \lambda m,\end{aligned}$$

where  $\mathbf{u} := (c, v)$  and function  $g$  of the nonlocal term  $\mathcal{A}$  specified by:

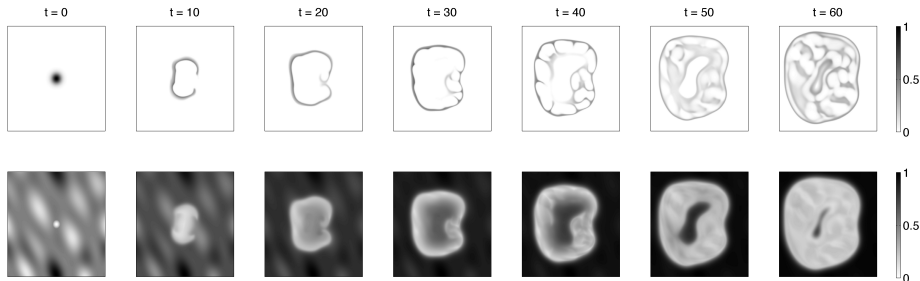
$$g(t, \mathbf{u}) = [S_{cc}(t)c + S_{cv}(t)v] \cdot (1 - u - v)^+.$$

Complemented with initial and zero-flux boundary conditions.

# Application: cancer invasion

[Domschke et al., J Theor Biol, 361, '14], also [G. & Chaplain, J Theor Biol, 250, '08]

- ▶ Heterogeneous initial ECM density; ECM remodelling at rate  $\mu_2 = 0.05$ .
- ▶ Cell-matrix adhesion coefficient increases from 0.25 to 0.5 at  $t = 10$
- ▶ Cell-cell adhesion coefficient decreases from 0.5 to 0.25 at  $t = 40$ .
- ▶ Plots of cell density (top row) and ECM density (bottom row) at various times  $t$ .



# Application: zebrafish pigmentation patterning

[Painter et al., Bull Math Biol, 77, '15]

- ▶ Stripe/interstripe pattern with thin separating strip without pigment cells:
  - ▶ black melanophores ( $u$ ) and some light reflecting iridophores;
  - ▶ yellowish xanthophores ( $v$ ) and light reflecting iridophores.
- ▶ Turing-type, morphogen-based models in the '90, but...  
...no chemical morphogenes were found!
- ▶ **Here:** minimal set of pattern-generating interactions, Cf. [Yamanaka & Kondo, PNAS, '14].
- ▶ “Run and Chase” proposed to explain pigmentation in zebrafish and related species:

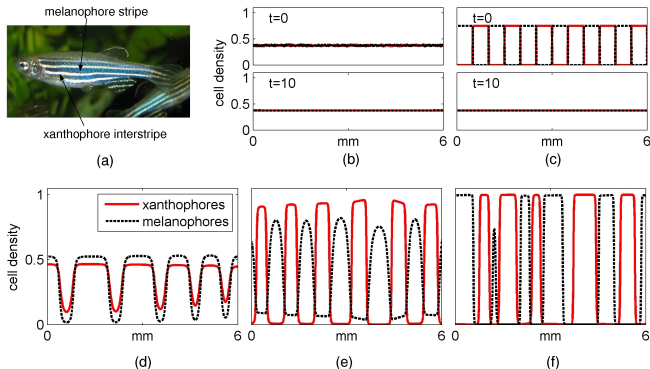


$u$  is repelled by  $v$ , i.e.  $C_{uv} < 0$ ,  $v$  is attracted by  $u$ , i.e.  $C_{vu} > 0$ .



# Application: zebrafish pigmentation patterning

[Painter et al., Bull Math Biol, 77, '15]



(b,c) no homotypic interaction  $\rightsquigarrow$  no pattern; (d)  $C_{uu} > 0 \rightsquigarrow$  mixed aggregates;  
(e)  $C_w > 0 \rightsquigarrow$  realistic pattern; (f)  $C_{uu}, C_{vv} > 0 \rightsquigarrow$  no robust pattern.

More likely: include the impact of iridophore cells to explain patterning.

# The Armstrong-Painter-Sherratt model

## Analytical results

- ▶ Existence and boundedness of solution
  - ▶ [Sherratt et al., '09] boundedness in 1D requires additional assumptions on  $g$  and  $\Omega$ .
  - ▶ [Chaplain et al., '11] local and global existence for nonlocal cancer-ECM adhesion model.
  - ▶ [Hillen et al., '17] local and global existence and boundedness for Armstrong-Painter-Sherratt and cancer invasion model.

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- ▶ One cell population: aggregation takes place for  $C > 0$  sufficiently large.
- ▶ For sensing radius  $R \rightarrow 0$ : nonlocal model reduces to [G., Chaplain, '08]
  - ▶ standard taxis model for linear  $g$ ,
  - ▶ volume-filling taxis [Hillen, Painter, '01] model for logistic (volume filling)  $g$ .

# Derivation of the nonlocal PDE model from a position-jump model



In [Gerisch & Painter, '10] we state:

A highly desirable objective is to develop continuous models for cellular adhesion as the appropriate limit from an underlying individual model for cell movement [...].

We propose to fill this gap via a position-jump model (spatial stochastic random walk).

[Buttenschön et al, J Math Biol, 76, '18]

Goals:

- ▶ a better understanding of underlying modelling assumptions and
- ▶ a framework to modify the continuous model as needed.

Master equation for a position jump process [Othmer, Dunbar, Alt, '88]:

$$\partial_t u(t, x) = \lambda \int_{\mathcal{D}} T(x, y) u(t, y) - T(y, x) u(t, x) d\mu(y)$$

where

- ▶  $(\mathcal{D}, \mu)$  measure space representing physical space (domain or grid),
- ▶  $\lambda$  jump rate,
- ▶  $T(x, y)$  probability density function for jump from  $y$  to  $x$ .

# A position-jump model

$$\partial_t u(t, x) = \lambda \int_{\mathcal{D}} T(x, y) u(t, y) - T(y, x) u(t, x) d\mu(y)$$

Define:

- ▶ Heading  $z := x - y$ , such that  $T_y(z) := T(y + z, y) = T(x, y)$ ,
- ▶ symmetric set  $D^y$ , the set of possible headings from  $y$ .

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## Lemma (Even and Odd Decomposition)

There is a decomposition

$$T_y(z) = S_y(z) + \mathbf{A}_y(z) \cdot \frac{z}{|z|},$$

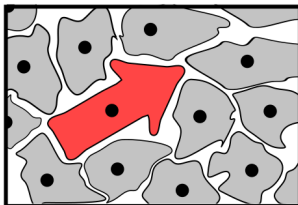
where  $S_y(z) = S_y(-z)$  and  $\mathbf{A}_y(z) = \mathbf{A}_y(-z)$  are symmetric.



# A position-jump model

$$T_y(z) = S_y(z) + \mathbf{A}_y(z) \cdot \frac{z}{|z|},$$

- ▶  $S_y(z)$  will become the **motility** and lead to the **diffusion term**.
- ▶  $\mathbf{A}_y(z)$  will define the cell **polarization** and lead to the **adhesion integral term**.



## A position-jump model

- ▶ Consider myopic random walk, that is  $S_y(z) = S_y$  and  $\mathbf{A}_y(z) = \mathbf{A}_y$ .
- ▶ Substitute  $T_y(z)$  into the master equation and rearrange.
- ▶ Consider small jumps of length  $h$  in any direction in  $\mathbb{S}^{n-1}$ .
- ▶ Use Taylor expansion in  $h$ .

Then

$$\partial_t u = \lambda h^{n-1} \left[ \frac{h^2 |\mathbb{S}^{n-1}|}{2n} \Delta (S_x u(t, x)) - \frac{h |\mathbb{S}^{n-1}|}{n} \nabla \cdot (\mathbf{A}_x u(t, x)) \right] + h.o.t.$$

# A position-jump model

## Advection-diffusion limit

Assume  $\mathbf{A}_x = O(h)$  and consider the parabolic scaling, i.e.  $1/\lambda \sim h^{n+1}$ .

Then the following limits exist:

$$\lim_{h \rightarrow 0, \lambda \rightarrow \infty} \frac{\lambda h^n |\mathbb{S}^{n-1}|}{n} \mathbf{A}_x = \alpha(x) \quad \text{and} \quad \lim_{h \rightarrow 0, \lambda \rightarrow \infty} \frac{\lambda h^{n+1} |\mathbb{S}^{n-1}|}{n} S_x = D(x)$$

...leading to the following limit equation

$$\partial_t u(t, x) + \nabla \cdot (\alpha(x) u(t, x)) = \Delta (D(x) u(t, x))$$

Note: spatial diffusion parameter appears inside Laplacian (expected for transition rates based on local information, cf. [Stevens & Othmer, '97]).

# A position-jump model

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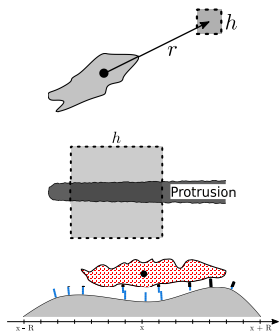
Note: spatial diffusion parameter appears inside Laplacian (expected for transition rates based on local information, cf. [Stevens & Othmer, '97]).

We assume

- ▶  $S_x$  is constant, i.e.  $D$  is constant.
- ▶  $\mathbf{A}_x$  is given by the net adhesive force acting on the cell that is located at  $x$ .

# A position-jump model

## Microscopic biological assumptions



$V_h$ : small test volume inside sensing region of cell at  $x$ .

- ▶ The distance from  $V_h$  to the cell body is  $r$ .
- ▶ Direction of generated force is  $r/|r|$ .
- ▶ The free space in  $V_h$  is  $f(x+r)$ .
- ▶ The part of the cell protrusion that is inside  $V_h$  is independent of  $x$  and called  $\Omega(r)$ .
- ▶ The density of adhesion bonds formed with background population in  $V_h$  is called  $N_b(x+r)$ .
- ▶ The adhesive strength per bond is  $\gamma$ .

↪ adhesive force generated in  $V_h$

$$\mathbf{F}_h(x+r) = \frac{r}{|r|} \gamma \underbrace{h^n N_b(x+r)}_{\text{\#of adhesion bonds}} \underbrace{f(x+r)}_{\text{free space}} \underbrace{h\Omega(r)}_{\text{amt. of cell in } V_h}.$$

# A position-jump model

## Net adhesive force

Summing  $\mathbf{F}_h(x+r)$  over all test volumina in the sensing region  $\mathcal{S}$

$$\rightsquigarrow \text{ net adhesive force } \mathbf{F}_{\text{net}}(x) = O(h)$$

and we let  $\mathbf{A}_x := \mathbf{F}_{\text{net}}(x)$ .

# A position-jump model

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Now, letting  $h \rightarrow 0$  and  $\lambda \rightarrow \infty$  yields

$$\alpha(x) = \int_{\mathcal{S}} \gamma \underbrace{N_b(x+r) f(x+r)}_{=g(\mathbf{u}(t,x+r))} \frac{r}{|r|} \Omega(r) dr.$$

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Next task: define specific  $N_b$ ,  $f$ , and  $\Omega$  for particular models.



# A position-jump model

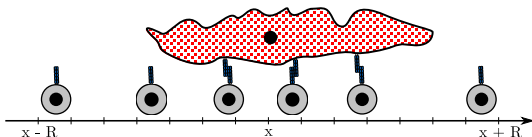
## The linear Armstrong-Painter-Sherratt model

1. One dimensional domain with sensing region  $\mathcal{S} = [-R, R]$ .
2. Assume law of mass action kinetics for the adhesion bonds, i.e.

$$N_b(x) \sim u(x)$$

3. Assume there is always free space, i.e.  $f \equiv 1$ .
4. Let  $\Omega(r)$  be the uniform distribution on  $\mathcal{S}$ , i.e.  $\Omega(r) = \frac{1}{2R}$ .

$$\partial_t u = Du_{xx} - \left( u \int_{-R}^R \gamma u(t, x+r) \frac{r}{|r|} \frac{1}{2R} dr \right)_x$$



# A position-jump model

## Adhesion with volume filling

1. Assume law of mass action kinetics for the adhesion bonds, i.e.

$$N_b(x) \sim u(x)$$

2. Assume space is limited  $f(u(t, x)) = (1 - u(t, x))^+$ .
3. Let  $\Omega(r)$  be the uniform distribution on  $\mathcal{S}$ .

$$\partial_t u = D\Delta u - \nabla \cdot \left( u \int_{\mathcal{S}} \gamma u(t, x+r) (1 - u(t, x+r))^+ \frac{r}{|r|} \Omega(r) dr \right)$$

Objective from spatial constraint:

Areas of high cell density in  $x + \mathcal{S}$  contribute less to the adhesive force in  $x$ .

# A position-jump model

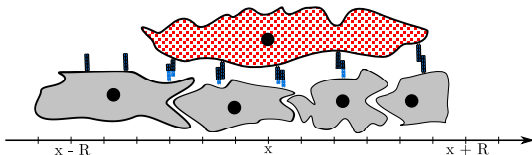
## Non-local background population

1. Assume cells in the background have spatial distribution of adhesion bonds  $\eta(r)$  around their cell center:

$$N_b(x) \sim \int_S u(t, x+r) \eta(r) dr.$$

2. Assume there is always free space, i.e.  $f \equiv 1$ .

$$\partial_t u = D \Delta u - \nabla \cdot \left( u \int_S \gamma \int_S u(t, x+y+r) \eta(y) dy \frac{r}{|r|} \Omega(r) dr \right)$$



# Numerical technique

## The need for efficient numerics

$$\partial_t u_i(t, x) = -\nabla \cdot (u_i(t, x) \mathcal{A}_i\{\mathbf{u}(t, \cdot)\}(x)) + \dots$$

- ▶ Spatial discretisation (FV, FD, FE) [we use 2nd order finite volumes]:  
 *$\mathcal{A}_i\{\mathbf{u}(t, \cdot)\}(x)$  must be evaluated for given approximations of  $\mathbf{u}(t, \cdot)$  in many points  $x$ , related to the spatial grid, of spatial domain  $\mathcal{D}$ .*
- ▶ Time integration [we use linearly implicit Runge-Kutta method ROWMAP]:  
*The spatial discretisation must be evaluated repeatedly over time for changing  $\mathbf{u}(t, \cdot)$ .*

**The evaluation of the nonlocal term quickly becomes the computational bottleneck of any numerical scheme.**

# Numerical technique

## Nonlocal term approximation — periodic setting



$$\mathcal{A}\{\mathbf{u}(t, \cdot)\}(x) := \frac{1}{\Phi R} \int_S \frac{r}{|r|} g(\mathbf{u}(t, x+r)) \Omega(|r|) dr ,$$

- ▶ Uniform  $N_1 \times N_2 \times \dots \times N_d$  grid on spatial domain  $\mathcal{D}$ .
- ▶ Grid cells  $\mathcal{D}_i$  of size  $h_1 \times h_2 \times \dots \times h_d$  indexed with  $i \in \mathcal{N}$  (multi-index set).
- ▶ Assume  $\mathbf{u}$  is given as centre value or volume average  $\mathbf{u}_i$  for each  $\mathcal{D}_i$ .
- ▶ Define  $g_i := g(\mathbf{u}_i)$  for all  $i \in \mathcal{N}$  and extend periodically beyond  $\mathcal{N}$ .
- ▶ Approximate  $g(\mathbf{u}(t, x))$  by piecewise constant reconstruction from

$$g(\mathbf{u}(t, x)) \approx \tilde{g}(x) := \sum_{i \in \mathcal{N}} g_i \chi_i(x)$$

The replacement of  $g(\mathbf{u}(t, \cdot))$  by  $\tilde{g}$  is the only approximation in the scheme.  
From now on everything will be essentially exact.

# Numerical technique

## Nonlocal term approximation — periodic setting



Nonlocal term after approximation for  $x^* \in \mathcal{D}$

$$v(x^*) := \frac{1}{\Phi R} \int_{\mathcal{S}} \frac{r}{|r|} \tilde{g}(x^* + r) \Omega(|r|) dr = \sum_{i \in \mathcal{N}} g_i \underbrace{\frac{1}{\Phi R} \int_{\mathcal{S}} \frac{r}{|r|} \chi_i(x^* + r) \Omega(|r|) dr}_{:= w_i^* \text{ (integration weight)}} .$$

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All points  $x_m^* := x^* + (m_j h_j)_{j=1}^d$  with  $m \in \mathbb{Z}^d$  share the same integration weights!

$$v(x_m^*) = \sum_{i \in \mathcal{N}} g_{i+m} w_i^*.$$

Integration weights  $w_i^*$

- ▶ depend only on known quantities,
- ▶ can be precomputed with your favourite method to arbitrary accuracy, and
- ▶ can be applied to evaluate the nonlocal term for arbitrary  $\tilde{g}$  and all  $x_m^*$ .

# Numerical technique

## Nonlocal term approximation — periodic setting



$$v(x_m^*) = \sum_{i \in \mathcal{N}} g_{i+m} w_i^* .$$

Do we need all  $w_i^*$ ? — If  $\{x^* + \mathcal{S}\} \cap \mathcal{D}_i = \emptyset$  then  $w_i^* = 0$ .

Let  $\tilde{\mathcal{N}} \subset \mathbb{Z}^d$  be an interval such that for all  $i \in \mathcal{N} \setminus \{i^* + \tilde{\mathcal{N}}\}$  holds  $w_i^* = 0$ .



# Numerical technique

## Nonlocal term approximation — periodic setting

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- ▶ Apply integration rule by executing the sum. Can become expensive!
- ▶ Combined evaluation at same location within each grid cell  $\mathcal{D}_n, \forall n \in \mathcal{N}$ :

$$v_n := v(x_{n-i^*}^*) = \sum_{i \in \tilde{\mathcal{N}}} g_{i+n} w_i, \quad w_i := w_{i+i^*}^* .$$

This provides a linear map from  $G := (g_n)_{n \in \mathcal{N}}$  to  $V := (v_n)_{n \in \mathcal{N}}$ .

**Structure of corresponding matrix  $W$ :**

$d = 1$ : (banded) circulant matrix  $W$  of weights;

$d = 2$ : (banded) block-circulant matrix with (banded) circulant blocks; ...

# Numerical technique

## Nonlocal term approximation — periodic setting



Evaluation of the matrix-vector product  $V = WG$  (case  $d = 1$ ).

- ▶ Circulant matrix  $W$  defined by its first column, denoted  $w$ .
- ▶ Any circulant  $W$  is diagonalized by the Fourier transform matrix  $F$  and

$$WF = F \operatorname{diag}(Fw).$$

- ▶ Thus

$$V = WG = F \operatorname{diag}(Fw)F^*G = \operatorname{FFT}(\operatorname{FFT}(w) .* \operatorname{iFFT}(G)),$$

where FFT denotes the Fast Fourier Transform algorithm and iFFT its inverse;  $.*$  is element-wise multiplication.

- ▶ This cuts the operation count down from  $\mathcal{O}(N_1^2)$  to  $\mathcal{O}(N_1 \log(N_1))$  operations.

**General  $d$ :** replace FFT by its  $d$ -dimensional counterpart FFT $d$ .

# Numerical technique

## Nonlocal term approximation — extensions

- ▶ From periodic to non-periodic boundary conditions:
  - ▶ Need to specify values of  $g(\mathbf{u})$  outside of  $\mathcal{D}$  for definition of the integral.
  - ▶ Banded circulant matrices replaced by banded Toeplitz matrices.
  - ▶ Toeplitz-to-circulant embedding saves efficient FFT-based algorithm.
- ▶ From uniform to non-uniform grids:
  - ▶ The integration weights independent of evaluation position property breaks down and so the combined evaluation via FFT.
  - ▶ Combined evaluation via fixed uniform intermediate grid works (it's not perfect).

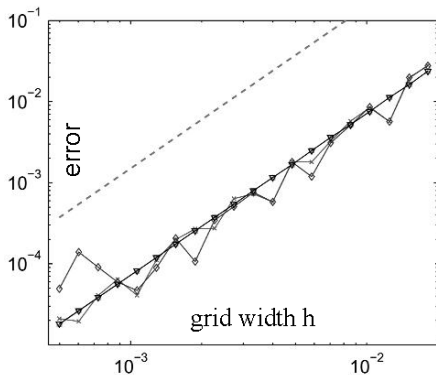
# Numerical technique

## Nonlocal term approximation — Accuracy

- ▶  $\mathcal{D} \subset \mathbb{R}$
- ▶  $\tilde{g}(x) = \sin(8\pi x)$
- ▶  $S = B(0, 0.1)$
- ▶  $\Omega \equiv 1$

↪ nonlocal term in analytical form

error for decreasing grid width →



Nonlocal term evaluation converges with order two for grid width to zero.

# Numerical technique

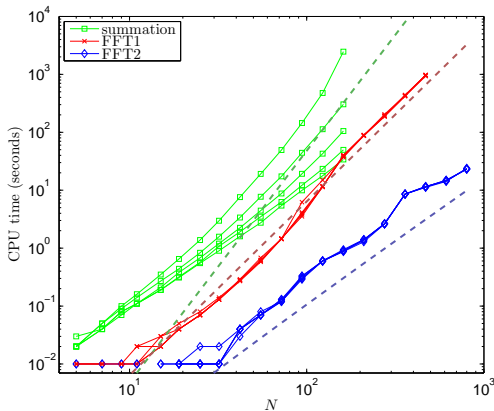
## Nonlocal term approximation — Efficiency

$d = 2$ -dimensional square domain  $\mathcal{D}$  with  $N \times N$  grid, periodic BCs.

Tests for increasing  
sensing region  $\mathcal{S}$ .

FFT2 vs. summation:  
reduction of operations ( $h = \frac{1}{N}$ )  
 $\sim N^4 \rightarrow \sim N^2 \log(N)$ .

Matrix-vector product:  
speed-up: 10 – 100



# Numerical technique

## Nonlocal term approximation — Efficiency

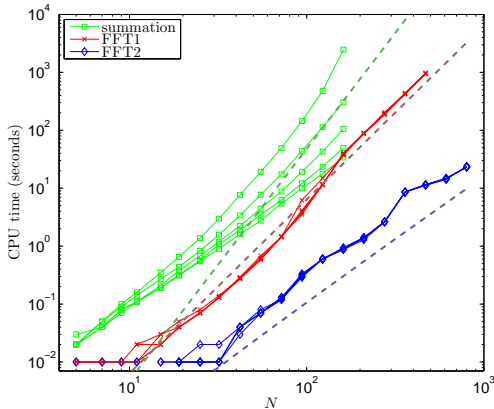
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Matrix-vector product:  
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Full cell sorting simulation:  
speed-up:  $\approx 20$  (2 h to 6 min).



- ▶ Cell attraction and repulsion are important basic mechanisms in biology.
- ▶ A flexible nonlocal continuous model is available and successfully applied in models in developmental and cancer biology.
- ▶ These models can be efficiently simulated by making use of the FFT; spatially highly resolved long time simulations are feasible.
- ▶ The requirements of periodic boundary conditions and uniform spatial grids can be relaxed while maintaining favourable algorithmic properties.
- ▶ The nonlocal continuous model can be derived from a stochastic random walk. This allows for better insight into the parametrisation of the continuous model.

- ▶ Cross-diffusion, instead of Fickian diffusion, in models with multiple cell types can significantly sharpen interfaces.
- ▶ Modelling neural crest cell invasion requires nonlocal term on growing domains. Efficient numerics?
- ▶ More tests and experience for non-uniform grids.



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Thank you very much for your attention!