

# Modeling crawling cell movement

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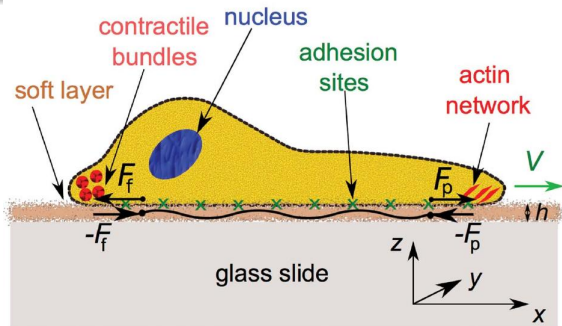
Group Seminar, April 2014

- several moving cells<sup>1</sup>
- Top left: mouse fibroblasts moving into an artificial wound (total video time: 3h)
- Bottom left: chick fibroblasts (total video time: 2h)
- Top right: mouse melanoma cell (total video time: 20min)
- Bottom right: trout epidermal keratocyte (total video time: 4min)

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<sup>1</sup>Video from: A Video Tour of Cell Motility, <http://cellix.imba.oeaw.ac.at/> 

# Sketch of cell cross section



2

- 2D cell shape modeled by phase field  $\rho(x, y, t)$
- $\rho = 1$ : cell,  $\rho = 0$ : no cell
- we neglect variations in height of cell
- nucleus rolls behind the lamellipodium front<sup>3</sup>

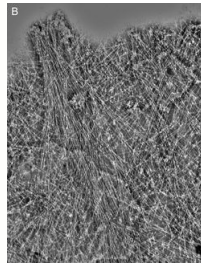
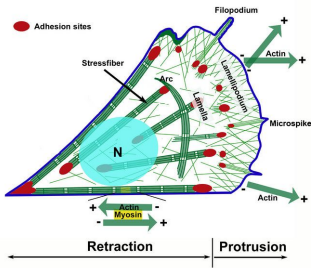
<sup>2</sup>Image from: F. Ziebert and I. S. Aranson, PLOS ONE, **8**, e64511.

<sup>3</sup>Video from: A Video Tour of Cell Motility, <http://cellix.imba.oeaw.ac.at/>

# Actin cytoskeleton

- cell crawling is driven by the continuous reorganization and turnover of the actin cytoskeleton
- two functions
  - protrusion by polymerization
  - contraction by interaction with myosin

- modeled by average actin orientation field  $\mathbf{p} = \begin{pmatrix} p_x(x, y, t) \\ p_y(x, y, t) \end{pmatrix}^4$

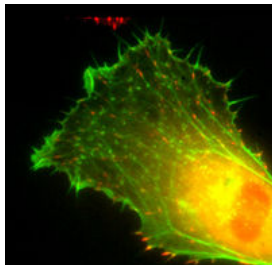


(a) Schematics of actin network (b) Closeup of actin filaments

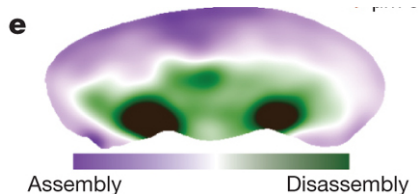
<sup>4</sup>Images from: A Video Tour of Cell Motility, <http://cellix.imba.oeaw.ac.at/>

# Adhesion sites

- adhesion sites connect the actin network to the substrate
- video: [adhesion sites \(red\)](#)<sup>5</sup>
- modeled by concentration of adhesion sites  $A(x, y, t)$
- adhesion sites do not move with the cell
- [rupture of adhesion sites in the retracting region of the cell](#)

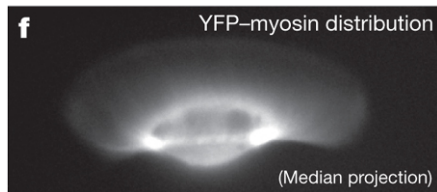


<sup>5</sup>JV Small, B Geiger, I. Kaverina, A. Bershadsky, Nat. Rev. Mol. Cell Biol. **3**, 957 (2002).



(c) Sites of actin assembly and disassembly

- myosin concentration is high where actin is disassembled
- could be modeled by an extra field  $m(x, y, t)$  but is eliminated in our model

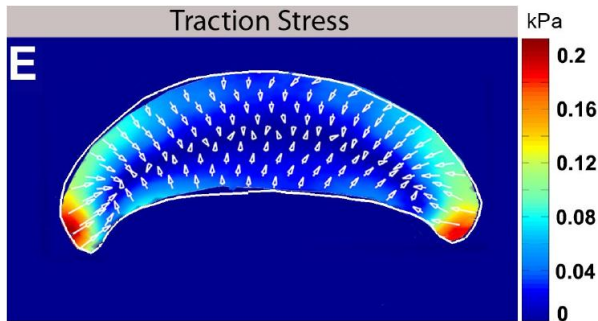


(d) Concentration of myosin

<sup>6</sup>CA Wilson et al. Nature **465**, 373 (2010).

# Traction and substrate displacements

- cell exerts traction forces  $\mathbf{T} = \begin{pmatrix} T_x(x, y, t) \\ T_y(x, y, t) \end{pmatrix}$  on substrate
- leads to substrate displacements<sup>7</sup>:  $\mathbf{u} = \begin{pmatrix} u_x(x, y, t) \\ u_y(x, y, t) \end{pmatrix}$



<sup>7</sup>MF Fournier, R Sauser, D Ambrosi, JJ Meister, AB Verkhovsky, J. Cell Biol. **188**, 287 (2010).

# Phase field $\rho(x, y, t)$

- phase field:  $\rho = 1$ : cell,  $\rho = 0$ : no cell,  $\nabla\rho \neq 0$ : cell boundary

$$\partial_t \rho = D_\rho \Delta \rho - (1 - \rho)(\delta - \rho) \rho - \alpha \mathbf{A} \mathbf{p} \cdot (\nabla \rho)$$

- $\rho(x) = 1 / (1 + \exp(x / \sqrt{D_\rho 2}))$  is a steplike stationary solution for  $\delta = 1/2$ : [Mathematica](#)
- volume conservation by feedback
  - $\langle \rho \rangle$  = volume integral over  $\rho$
  - $V_0$ : initial volume
  - $\sigma |\mathbf{p}|^2$  models actin network contraction

$$\delta = \frac{1}{2} + \mu (\langle \rho \rangle - V_0) - \sigma |\mathbf{p}|^2$$

- advection of  $\rho$  along the actin orientation vector  $\mathbf{p}$ ,  
 $\alpha$ : propulsion strength



# Actin orientation field $\mathbf{p}(x, y, t)$

$$\partial_t \mathbf{p} = D_\rho \Delta \mathbf{p} - \tau_1^{-1} \mathbf{p} - \tau_2^{-1} (1 - \rho^2) \mathbf{p} - \beta f(\nabla \rho) - \gamma [(\nabla \rho) \cdot \mathbf{p}] \mathbf{p}$$

- nearest neighbour interaction by diffusion  $D_\rho$
- degradation of actin by depolymerization inside ( $\tau_1$ ) and outside ( $\tau_2$ ) of the cell
- actin created by polymerization at the cell front,  
 $f(\kappa) = \frac{\kappa}{\sqrt{1 + \epsilon \kappa^2}}$  saturates for large  $\kappa$
- reflection symmetry broken due to myosin motors

# Myosin concentration $m(x, y, t)$

- actin disassembles where myosin concentration is higher than equilibrium value  $m_0$

$$\partial_t \mathbf{p} = D_\rho \Delta \mathbf{p} - \tau_1^{-1} \mathbf{p} - \tau_2^{-1} (1 - \rho^2) \mathbf{p} - \beta f(\nabla \rho) - (m - m_0) \mathbf{p}$$

- myosin
  - diffuses with coefficient  $D_m$
  - relaxes to  $m_0$  with rate  $\tau_m$
  - moves along the actin filaments with velocity  $V_m$
  - is suppressed near to front of the cell with rate  $\bar{\gamma} \nabla \rho \cdot \mathbf{p}$

$$\partial_t m = D_m \Delta m - \tau_m^{-1} (m - m_0) + V_m \mathbf{p} \cdot \nabla m + \bar{\gamma} \nabla \rho \cdot \mathbf{p}$$

- assume  $\tau_m \ll 1$

$$m - m_0 \approx \tau_m \bar{\gamma} \nabla \rho \cdot \mathbf{p}$$

# Concentration of adhesion sites $A(x, y, t)$

$$\partial_t A = D_A \Delta A + a_0 \rho p^2 + a_{nl} \rho A^2 - s A^3 - d(|\mathbf{u}|) A$$

- adhesion sites form only if actin is present but independent of actin direction: linear attachment  $\sim \rho p^2$
- already formed adhesion complex favors formation of more adhesive contacts nearby: nonlinear attachment  $\sim A^2$
- nonlinear detachment  $\sim A^3$  locally saturates concentration of adhesion sites
- breakup of adhesion sites if substrate displacement  $|\mathbf{u}|$  exceeds critical displacement  $U_c$ : linear step-like detachment rate

$$d(|\mathbf{u}|) = \frac{d}{2} \left( 1 + \tanh \left[ b (\mathbf{u}^2 - U_c^2) \right] \right)$$

# Substrate model: Kelvin-Voigt material

- stress tensor of 3D incompressible isotropic visco-elastic (Kelvin-Voigt) material

$\mathbf{u}$ : displacements,  $p$ : pressure,  $\tilde{G}$ : shear modulus,  $\tilde{\eta}$ : viscosity

$$\sigma_{ik} = \tilde{G}(u_{i,k} + u_{k,i}) + \tilde{\eta}(\dot{u}_{i,k} + \dot{u}_{k,i}) - p\delta_{ik}$$

- overdamped motion:  $\ddot{u}_i = 0$ ,  $\sigma_{ik,k} = 0$

$$\tilde{G}\nabla^2\mathbf{u} + \tilde{\eta}\nabla^2\dot{\mathbf{u}} = \nabla p, \quad \nabla \cdot \mathbf{u} = 0$$

- lower boundary conditions:  $\mathbf{u}(x, y, z = 0, t) = 0$
- upper boundary conditions: traction force  $\mathbf{T}$ ,  $H$ : height of substrate layer

$$\sigma_{xz}(x, y, z = H, t) = T_x(x, y, t),$$

$$\sigma_{yz}(x, y, z = H, t) = T_y(x, y, t),$$

$$\sigma_{zz}(x, y, z = H, t) = 0,$$

- periodic boundary conditions in  $x$ -,  $y$ - direction with period  $L$

# Substrate model: traction forces $\mathbf{T}(x, y, t)$

- integrate over  $z$ - direction
- assume height  $\ll$  lateral extension:  $H \ll L$ , expand in  $H/L$

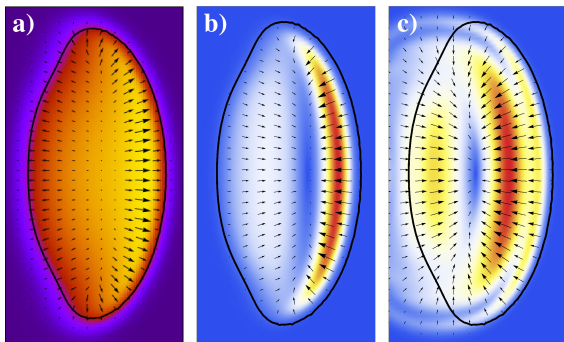
$$\partial_t \mathbf{u} = -\frac{1}{\eta} \left( G \mathbf{u} - \frac{1}{\xi} \left( \mathbf{T} + h \left[ 5 \Delta \mathbf{T} + 19 \nabla (\nabla \cdot \mathbf{T}) \right] \right) \right)$$

- traction due to actin polymerization:  $\mathbf{T}_{\text{pr}} = -\xi \rho A \mathbf{p}$
- traction due to friction:  $\mathbf{T}_{\text{fr}} = \rho A \zeta$
- cell does not exert a net force on substrate:  
determine  $\zeta$  by  $\langle \mathbf{T}_{\text{pr}} + \mathbf{T}_{\text{fr}} \rangle = 0$

$$\mathbf{T} = \xi A \rho \frac{\langle A \mathbf{p} \rho \rangle}{\langle A \rho \rangle} - \xi A \rho \mathbf{p}$$

- for heterogeneous substrate, shear modulus  $G$  (stiffness) depends on space

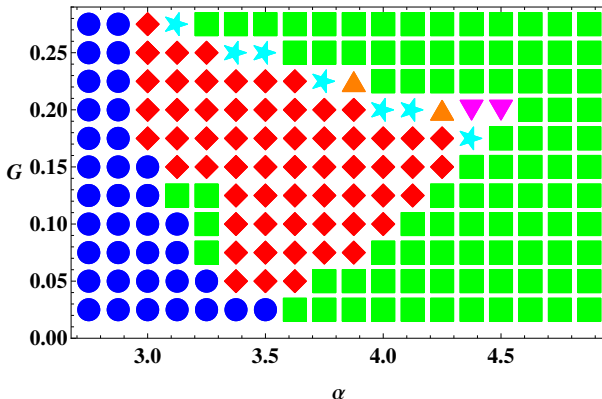
# Cell shape



**Figure:** Shape of cells in the steady moving regime. Black contour:  $\rho = 0.25$ . a) Actin orientation field  $\mathbf{p}$ . b) Traction force  $\mathbf{T}$ . Red (blue) corresponds to large (small) values of  $|\mathbf{T}|$ . c) Displacements field  $\mathbf{u}$ . Red (blue) corresponds to large (small) values of  $|\mathbf{u}|$ .

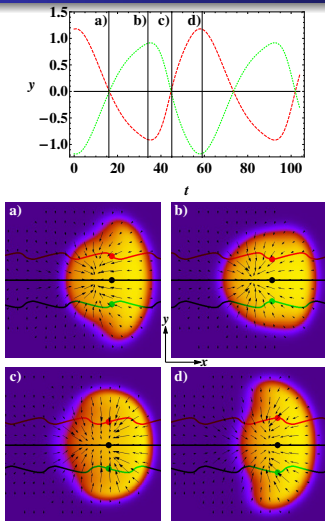
# Phase diagram

Propulsion strength  $\alpha$  vs. substrate's shear modulus  $G$



**Figure:** Phase diagram for propulsion strength  $\alpha$  vs. substrate's shear modulus  $G$ . ● denotes non-moving states, ■ steady moving (gliding) states, ◆ stick-slip motion, ★ wandering bipedal and ▼, ▲ breathing and bipedal modes, respectively.

# Stick-slip motion



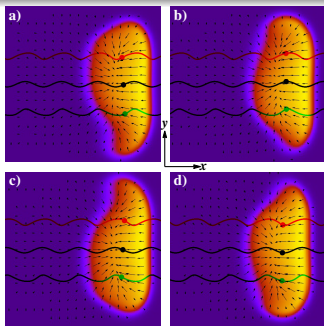
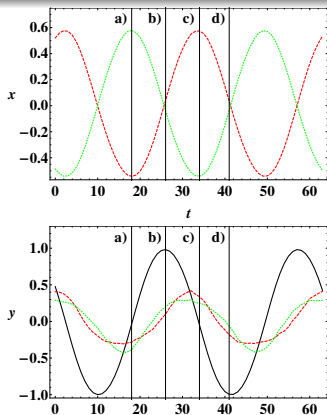
- top panel:  $y$ -component of center of mass (c.o.m.) of upper (red) and lower (green) half of cell
- $x$ -component does not show oscillations
- overall c.o.m. (black line) moves in a straight line
- compare with experiment<sup>a</sup>

<sup>a</sup>K. Keren et al. Nature **453**, 475 (2008).

Figure: Cell shape and substrate displacement field.



# Bipedal motion

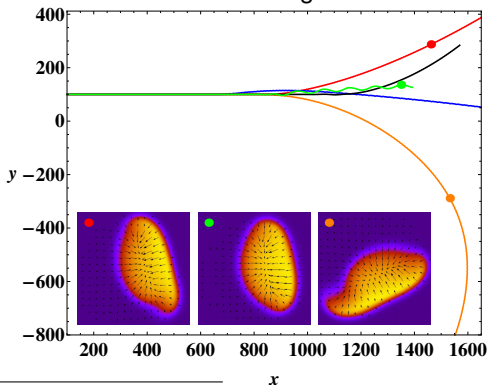


- anti-phase oscillations of c.o.m.  $x$ - components of upper (red) and lower (green) cell half
- in-phase oscillations of  $y$ - components
- c.o.m. (black) also oscillates [compare with experiment 1<sup>8</sup> 2](#)

<sup>8</sup>EL Barnhart, GM Allen, F Jülicher, JA Theriot, Biophys. J. **98**, 933 (2010).

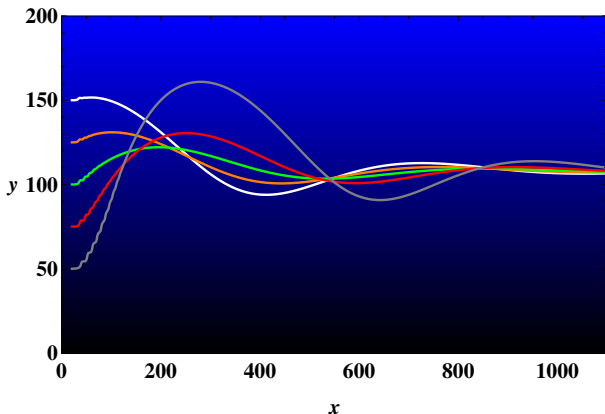
# Wandering bipedal

- instability in the propagation direction
- similar behavior found in a simple model for deformable self-propelled particles<sup>9</sup>:
  - drift bifurcation leads from stationary to moving states
  - 2nd bifurcation leads from straight motion to circular motion



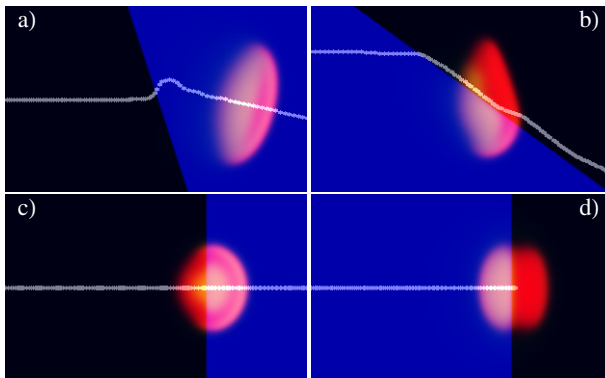
<sup>9</sup>T. Ohta, T. Ohkuma, PRL **102**, 154101 (2009).

# Durotaxis (cell migration in a stiffness gradient)



**Figure:** A linear gradient in substrate's stiffness  $G$  in the  $y$ -direction from  $G = 0$  (black) at the bottom to  $G = 0.4$  (blue) at the top. The curves show center of mass trajectories for different initial positions. They converge to an optimal value of  $G$ .

# Stiffness step



**Figure:** Examples for the behavior of cells colliding with a step in the substrate stiffness (blue:  $G = 0.4$ , black:  $G = 0.05$ ). The center of mass trajectories are shown in white. Top row:  $\alpha = 4 = 2\beta$ , bottom row:  $\alpha = 4, \beta = 1.5$ . Other parameters:  $U_c^2 = 0.25$ .

# Cell-cell interaction with multiple phase fields

- phase fields  $\rho_i$  for  $N$  cells

$$\partial_t \rho_i + \alpha \mathbf{A} \mathbf{p} \cdot \nabla \rho_i = D_\rho \Delta \rho_i - \frac{\partial}{\partial \rho_i} V(\rho_i) - \frac{\partial}{\partial \rho_i} W(\rho_1, \dots, \rho_N), \quad i = 1, \dots, N.$$

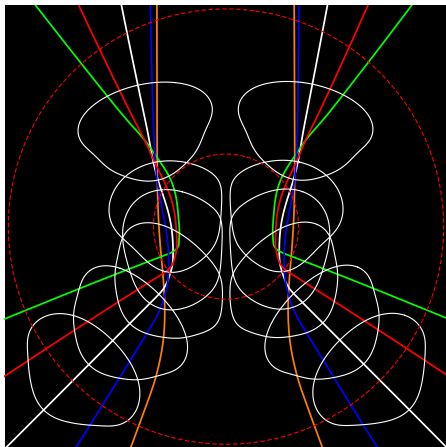
- $V$  : self-interaction  $\frac{\partial}{\partial \rho_i} V(\rho_i) = \rho_i (\rho_i - \delta_i) (\rho_i - 1)$
- $W$  : volume (steric) interaction avoids interpenetration of cells

$$W(\rho_1, \dots, \rho_N) = \sum_{j,k} W_2(\rho_j, \rho_k)$$

- two cell pair potential  $W_2(\rho_1, \rho_2) = \frac{\lambda}{2} \rho_1^m \rho_2^n$ 
  - large and positive if the two cells overlap
  - zero for no overlap
  - $W_2$  does not depend on  $m, n$  in the sharp interface limit  $D_\rho \rightarrow 0$
  - for  $D_\rho > 0$  perturbations could lead to  $\rho_i < 0 \Rightarrow$  choose even exponents  $m = n = 2$  to avoid attraction
- all other fields are shared between cells. [Video](#). [Experiment](#).<sup>10</sup>

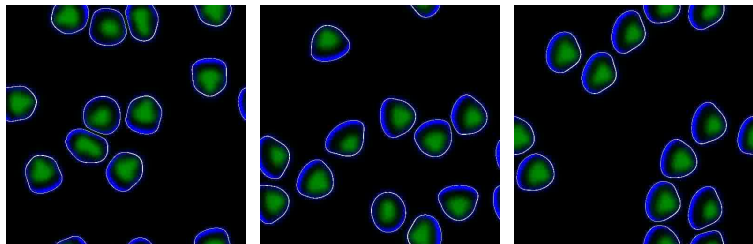
<sup>10</sup><http://cellix.imba.oeaw.ac.at/>

# Alignment mechanism responsible for collective motion



**Figure:** The angle of incidence of two cells colliding in a symmetric fashion is larger than their exit angles. White: phase field contours with  $\rho = 0.5$ . Colored: trajectories of colliding cell for different angles of incidence. [See video.](#)

# Unidirectional collective motion



**Figure:** Initially, cells move uncorrelated. The alignment mechanism leads to an unidirectional collective motion towards the top left corner. Time is increasing from left to right. [Video](#). [Experiment from Phys. Rev. E 74, 061908 \(2006\)](#).

# Coexistence of moving and stationary cells

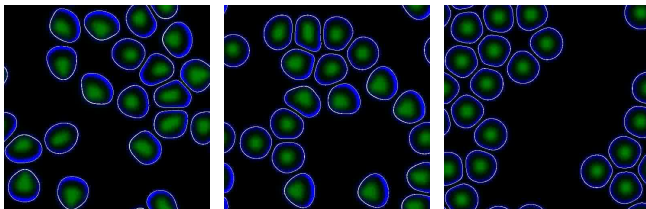


Figure: Initially moving cells gather in stationary clusters. [See video.](#)

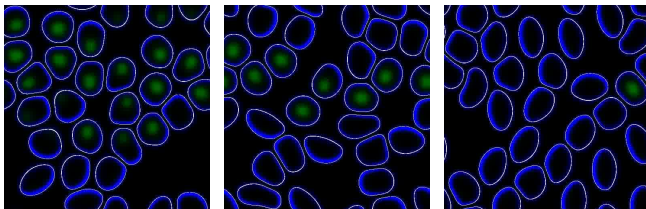


Figure: Initially, some cells are moving while some are stationary. Cell-cell collisions set the stationary cells into motion. [See video.](#)

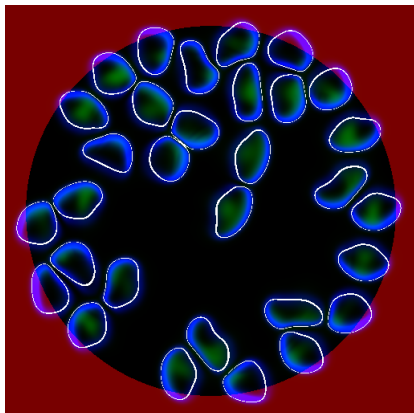
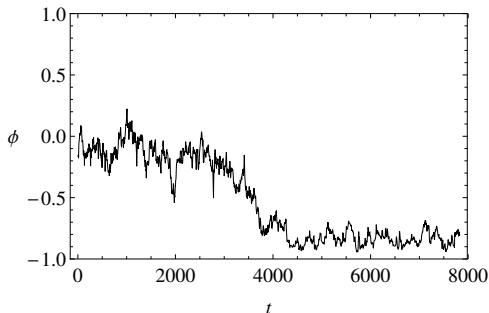


# Collective rotational motion

order parameter  $\phi$

$$\phi(t) = \frac{1}{N} \sum_{i=1}^N \hat{\mathbf{e}}_{\theta}(t) \cdot \hat{\mathbf{v}}_i(t)$$

normalized velocity vector  $\hat{\mathbf{v}}_i(t) = \frac{\mathbf{v}_i(t)}{|\mathbf{v}_i(t)|}$   
for each cell  $i$  is projected onto the unit  
vector  $\hat{\mathbf{e}}_{\theta}$  tangential to a circle



**Figure:** Clockwise rotational motion in a confined medium. Adhesion is larger inside. [Video. Experiment \(Phys. Rev. E 74, 061908 \(2006\)\).](#)

# Adhesion between cells

- keratocytes are responsible for wound healing  $\Rightarrow$  can build cell monolayers
- cell boundaries located at  $\nabla\rho_i$
- adhesion = interaction between cell boundaries:  $\nabla\rho_i \cdot \sum_{j \neq i} \nabla\rho_j$

$$\partial_t \rho_i + \alpha \mathbf{A} \mathbf{p} \cdot \nabla \rho_i + \underbrace{\kappa \nabla \rho_i \cdot \sum_{j \neq i} \nabla \rho_j}_{\text{cell-cell adhesion}} = D_\rho \Delta \rho_i - \frac{\partial}{\partial \rho_i} V(\rho_i) - \frac{\partial}{\partial \rho_i} W(\rho_1, \dots, \rho_N)$$

- multiple cells with cell-cell adhesion
- increasing adhesion strength  $\kappa$  should yield a transition to tissue (= cells sticking firmly together) but gives numerical instabilities instead
- other possibilities:<sup>11</sup>

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<sup>11</sup>Study on multicellular systems using a phase field model, M. Nonomura, PloS one 7, e33501 (2012).

# Summary

- phenomenological model for crawling cells based on a reaction-diffusion system
- cells exhibit different modes of movement accompanied by shape changes similar to experiments
  - stick-slip motion
  - bipedal motion
- migration of cells is sensitive to mechanical properties of substrate
- collective motion of multiple cells modeled with interacting phase fields

- introduce different adhesion terms to model tissue
- fit model parameters to specific cell types
- avoid breakup of cells
- derive model equations in a more fundamental way as e.g. in  
12

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<sup>12</sup>Generic theory of active polar gels: a paradigm for cytoskeletal dynamics, K. Kruse, J.F. Joanny, F. Jülicher, J. Prost, K. Sekimoto, Eur. Phys. J. E **16**, 5 (2005)

# For Further Reading I



J. Löber, F. Ziebert, and I. S. Aranson.

Modeling crawling cell movement on soft engineered substrates.

Soft Matter **10**, 1365 (2014).



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PLOS ONE, **8**, e64511.

## For Further Reading II

-  B. A. Camley, Y. Zhao, B. Li, H. Levine, and W.-J. Rappel.  
Phys. Rev. Lett. **111**, 158102 (2013).
-  D. Shao, H. Levine, and W.-J. Rappel, Proc. Natl. Acad. Sci.  
U.S.A. **109**, 6851 (2012).
-  D. Shao, W.-J. Rappel, and H. Levine.  
Phys. Rev. Lett. **105**, 108104 (2010).

## For Further Reading III

-  Toward a thermodynamically consistent picture of the phase-field model of vesicles: Curvature energy.  
D. Jamet, and C. Misbah, Phys. Rev. E **78**, 031902 (2008).
-  Thermodynamically consistent picture of the phase-field model of vesicles: Elimination of the surface tension.  
D. Jamet, and C. Misbah, Phys. Rev. E **78**, 041903 (2008).
-  Towards a thermodynamically consistent picture of the phase-field model of vesicles: Local membrane incompressibility.  
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-  Phase-field approach to three-dimensional vesicle dynamics.  
T. Biben, K. Kassner, and C. Misbah, Phys. Rev. E **72**, 041921 (2005).