# Collective cell motion in endothelial monolayers

A. Szabó<sup>1,5,6</sup>, R. Ünnep<sup>1</sup>, E. Méhes<sup>1</sup>, W.O. Twal<sup>2</sup>, W.S. Argraves<sup>2</sup>, Y. Cao<sup>3</sup>, A. Czirók<sup>1,4</sup>

<sup>1</sup>Eötvös University, Budapest, Hungary <sup>2</sup>Medical University of South Carolina, Charleston, SC, USA <sup>3</sup>Karolinska Institute, Stockholm, Sweden <sup>4</sup>University of Kansas Medical Center, Kansas City, KS, USA <sup>5</sup>Centrum Wiskunde en Informatica, Amsterdam, The Netherlands <sup>6</sup>NISB, NCSB



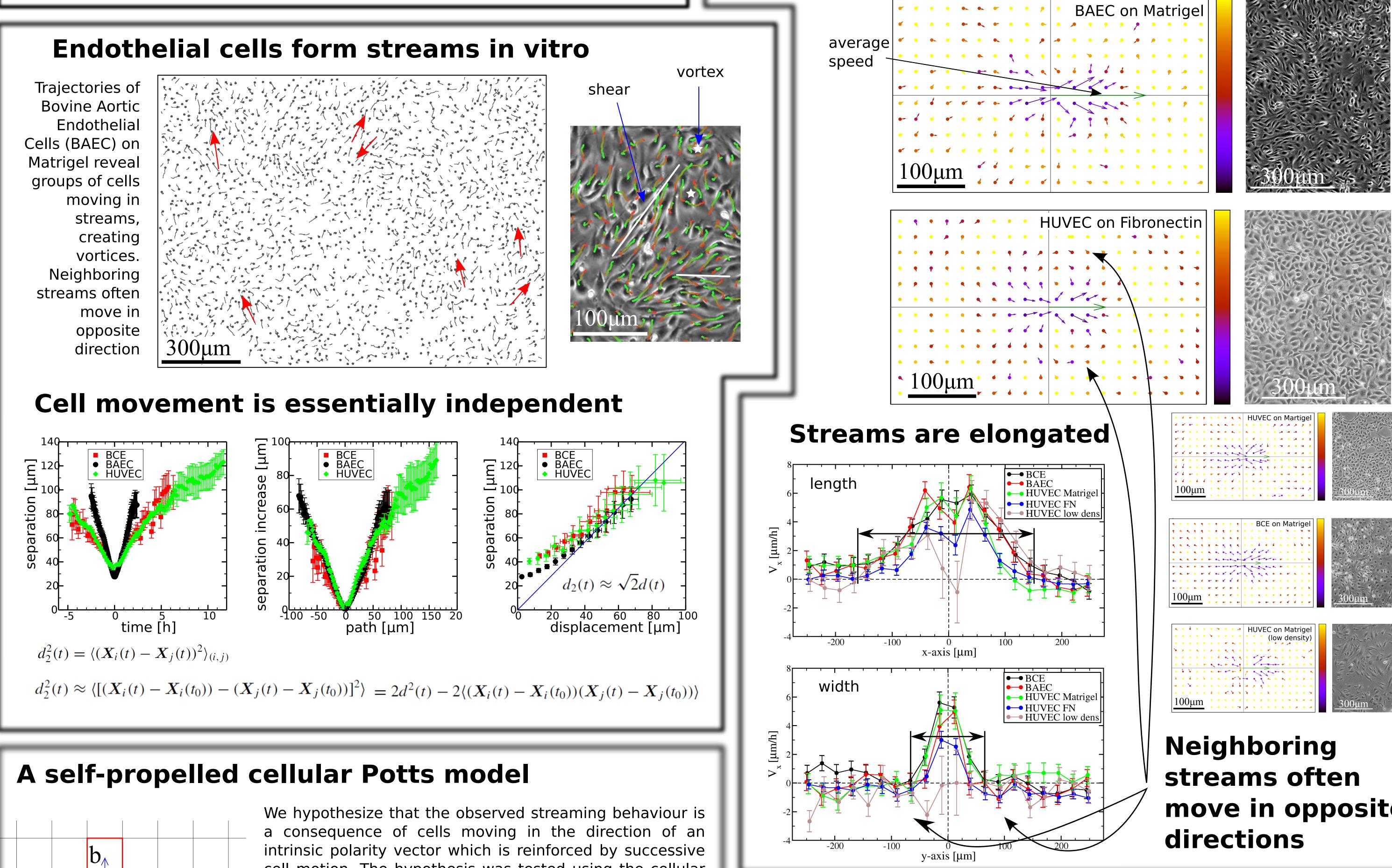
#### Abstract

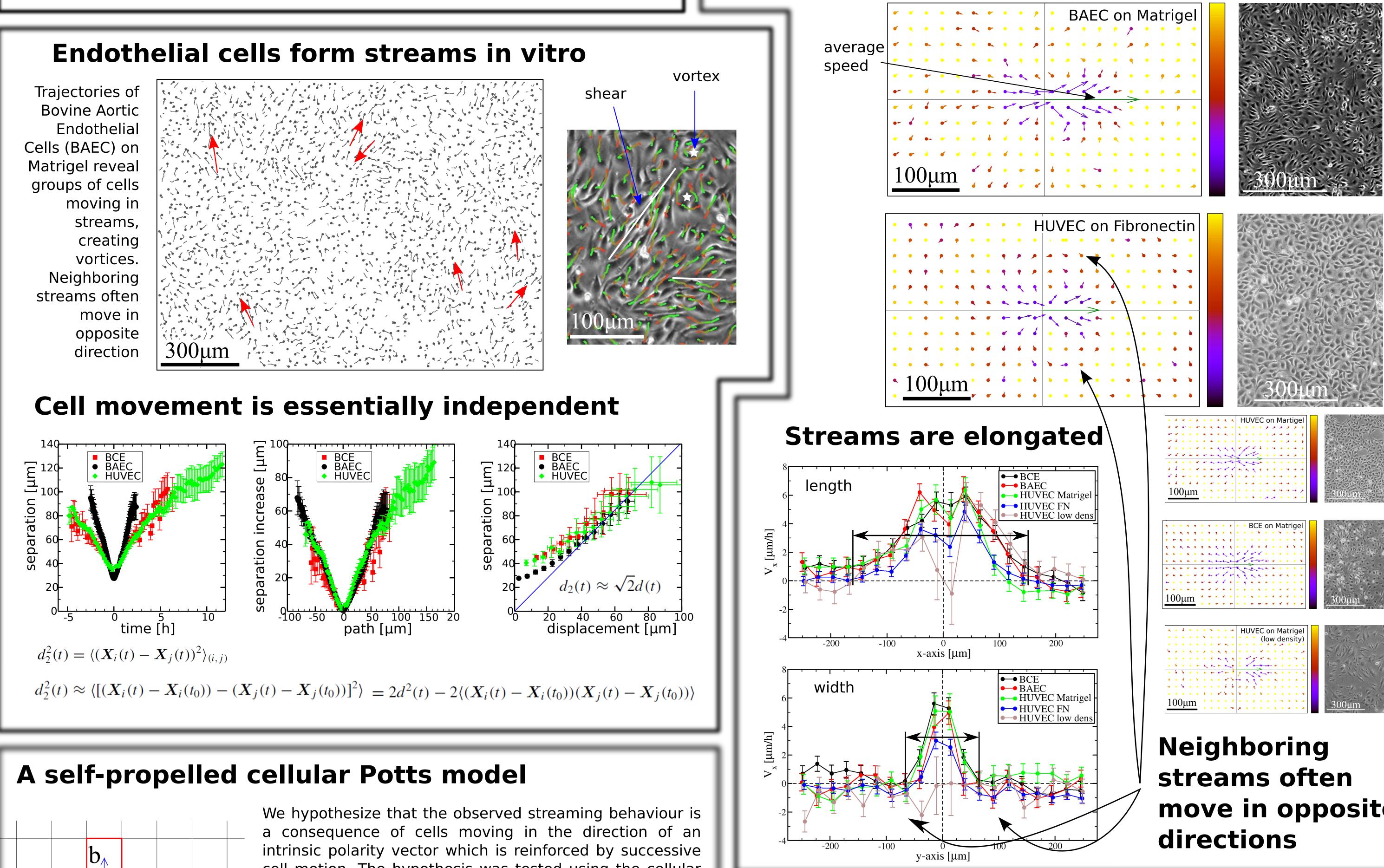
E · L · T · E KUTATÓEGYETEM

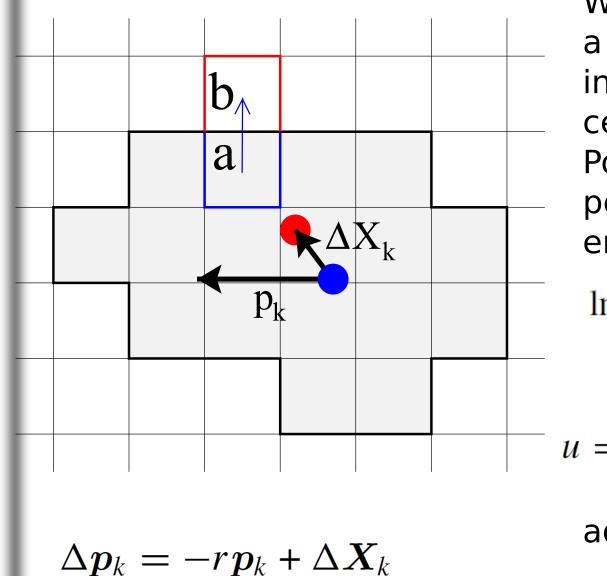
Random collective motion in monolayer cultures of endothelial cells is observed and described using statistical methods. Trajectory analysis reveal vortices and neighboring cell streams in opposing direction creating shear lines between cells. To understand the observed behavior, we implement active cell motility in the cellular Potts model using a compass-like cell polarity approach. For spontaneous directed motility we assume a positive feedback between cell displacements and cell polarity. The model behavior is compatible with the experimental results: both the speed and persistence of cell motion decrease in monolayer cultures, transient cell chains move together as groups, and velocity correlations extend over several cell diameters.

## Averaging vector fields around moving cells

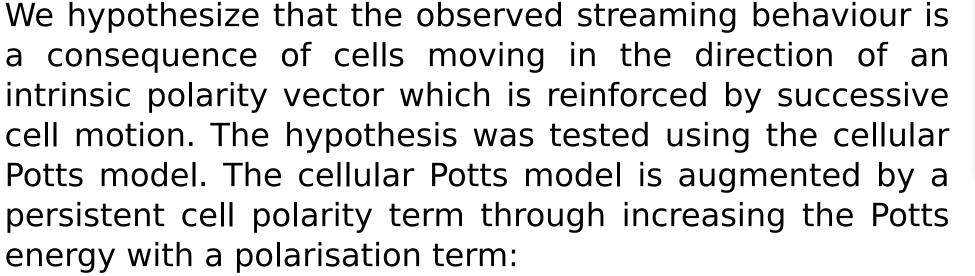
The average flow field around moving cells is a measure sensitive to the local cell movement patterns. For a given configuration of cell positions and velocities, this procedure assigns reference systems co-aligned with the movement of each cell and averages the velocity vectors observed at similar locations (e.g. immediately in front, behind, left and right). The vectors of the flow field diminish in a hypothetical ensemble of statistically independent cells, as they are averages of independent random vectors. The value of SEM relative to the vectors is depicted in color showing SEM=0 with black and SEM equal to the length of the vector with yellow.







polarity is updated with cell displacements



$$n \ p(a \to b) = \min[0, -\Delta u(a \to b) + w(a \to b)]$$

$$= \sum_{\langle x, x' \rangle} J_{\sigma(x), \sigma(x')} + \lambda \sum_{i=1}^{N} \delta A_i^2 \qquad w(a \to b) = P \sum_{k=\sigma(a), \sigma(b)} \frac{\Delta X_k(a \to b) p_k}{|p_k|}$$
  
dhesion and volume constraint probability of movement enhanced in

the direction of the polarity vector

The model reproduces the observed collective behavior:

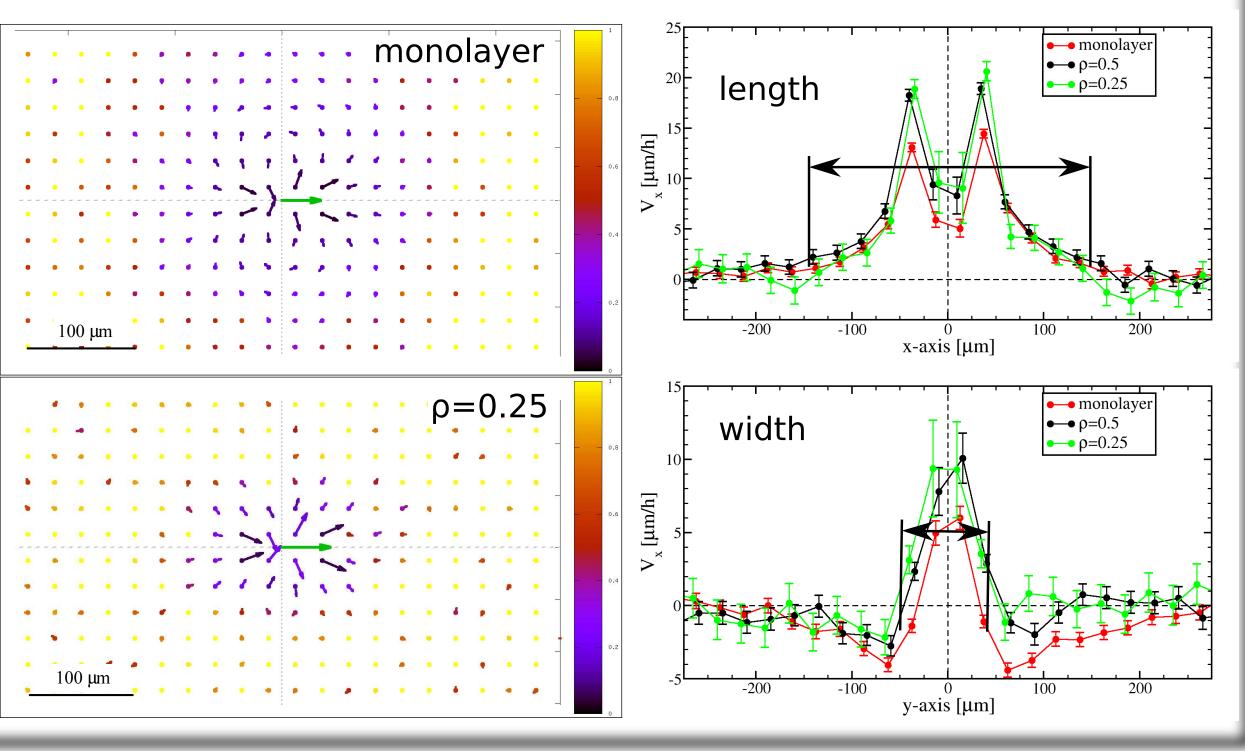
elongated streams form at random positions

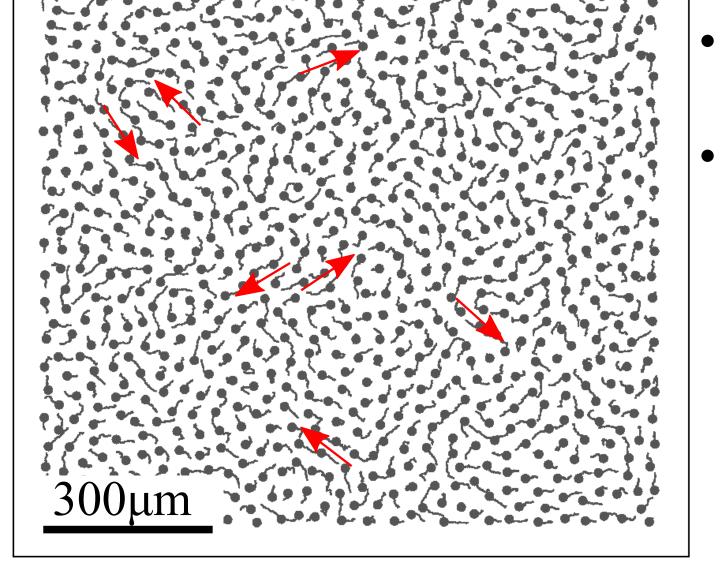
vortex

shear

move in opposite

### Streams in the model





neighboring streams often move in opposite directions, creating shear lines and vortices

, at low density the streams are still apparent, but shear is reduced

### Acknowledgements

We are grateful to Roeland MH Merks for generously sharing his simulation code with us, and to Charles D Little and Tamás Vicsek for stimulating discussions. This work was supported by the NIH (R01 HL87136) and the Hungarian Science Fund (OTKA K72664).

A Szabó, R Ünnep, E Méhes, WO Twal, WS Argraves, Y Cao, A. Czirók (2010) Collective cell motion in endothelial monolayers. *Physical Biology* 7:046007





