

Various aspects of reaction-diffusion problems

Tomáš Vejchodský

Centre for Mathematical Biology
Mathematical Institute



CMB Group Meeting, Oxford, 10 June, 2013

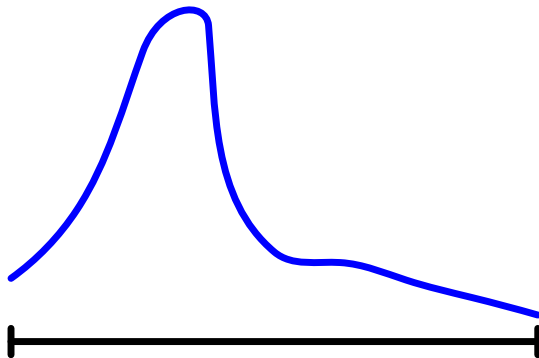
Numerical analysis – Finite Element Method

- ▶ Mesh adaptivity
- ▶ A posteriori error estimates
- ▶ Discrete maximum principles

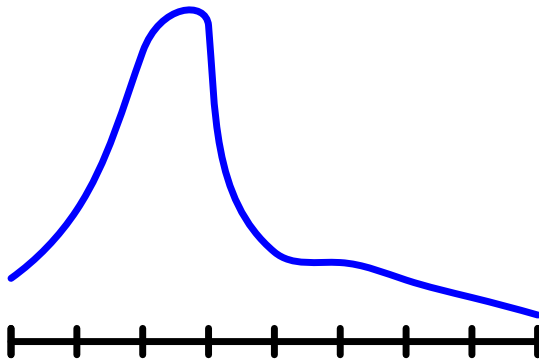
Mathematical biology

- ▶ Circadian rhythms
- ▶ Skin patterns formation

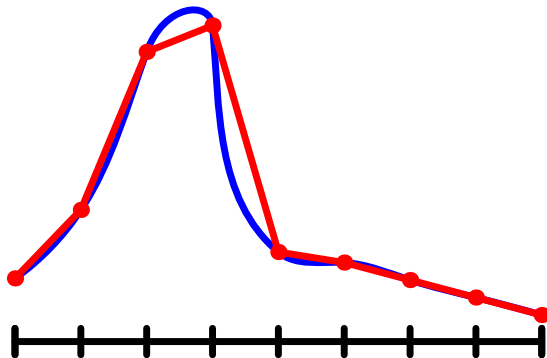
Mesh adaptivity



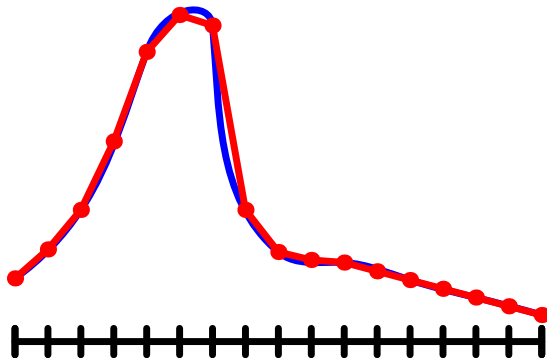
Mesh adaptivity



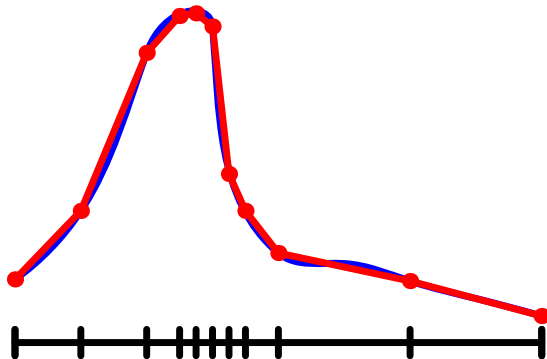
Mesh adaptivity



Mesh adaptivity



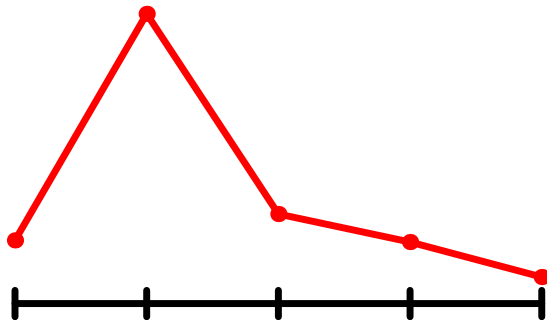
Mesh adaptivity



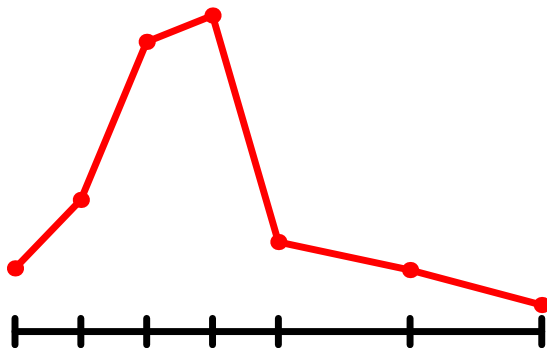
?



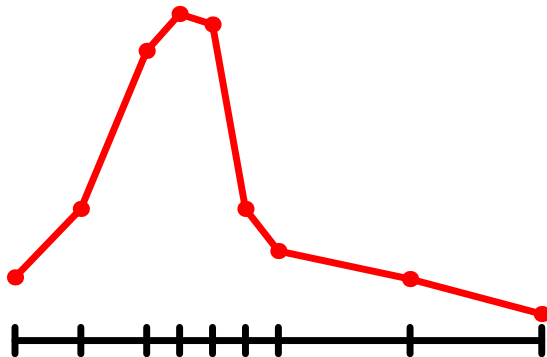
Mesh adaptivity



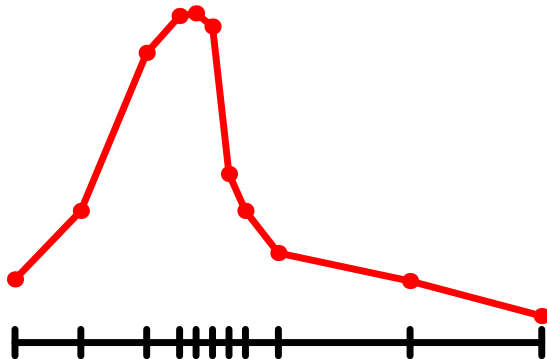
Mesh adaptivity



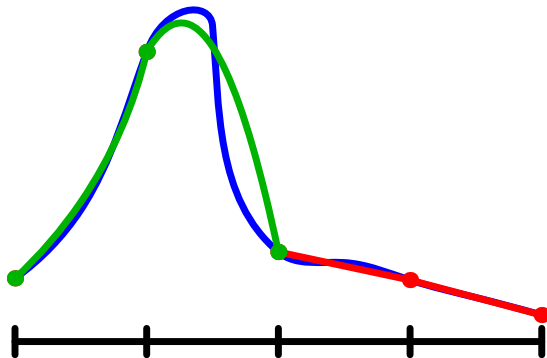
Mesh adaptivity



Mesh adaptivity



Mesh adaptivity – *hp* version



Error indicators \times Error estimators

Properties

- ▶ Efficiency and reliability
- ▶ Guaranteed upper bound
- ▶ Guaranteed lower bound
- ▶ Asymptotic exactness
- ▶ Robustness
- ▶ Locality

Approaches

- ▶ Explicit residual
- ▶ Implicit residual – Dirichlet
- ▶ Implicit residual – Neumann
- ▶ Hierarchical
- ▶ Postprocessing
- ▶ Complementarity
- ▶ Quantity of interest

Discrete maximum principles

$$-\Delta u = f \text{ in } \Omega = (0, 4) \times (0, 2), \quad u = 0 \text{ on } \partial\Omega$$

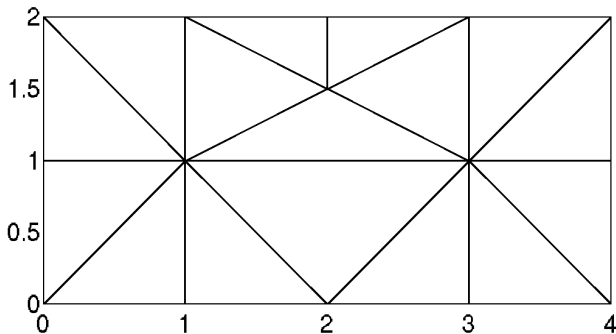
Conservation of nonnegativity: $f \geq 0 \Rightarrow u \geq 0$

Discrete maximum principles

$$-\Delta u = f \text{ in } \Omega = (0, 4) \times (0, 2), \quad u = 0 \text{ on } \partial\Omega$$

Conservation of nonnegativity: $f \geq 0 \Rightarrow u \geq 0$

$$f(x_1, x_2) = \begin{cases} 1 & \text{for } x_1 < 1 \\ 0 & \text{for } x_1 \geq 1 \end{cases} \quad u_h \text{ by linear FEM}$$



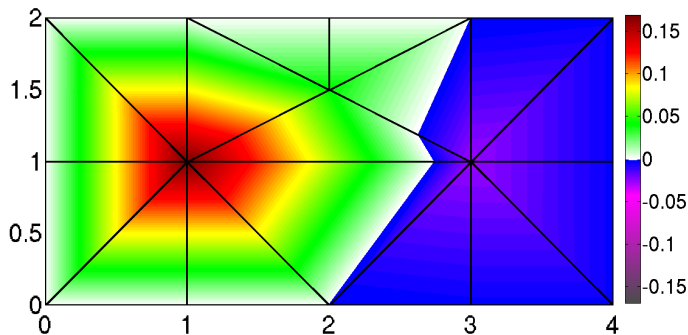
Brandts, Korotov, Křížek, Šolc, SIAM Review 51 (2009), 317–335

Discrete maximum principles

$$-\Delta u = f \text{ in } \Omega = (0, 4) \times (0, 2), \quad u = 0 \text{ on } \partial\Omega$$

Conservation of nonnegativity: $f \geq 0 \Rightarrow u \geq 0$

$$f(x_1, x_2) = \begin{cases} 1 & \text{for } x_1 < 1 \\ 0 & \text{for } x_1 \geq 1 \end{cases} \quad u_h \text{ by linear FEM}$$



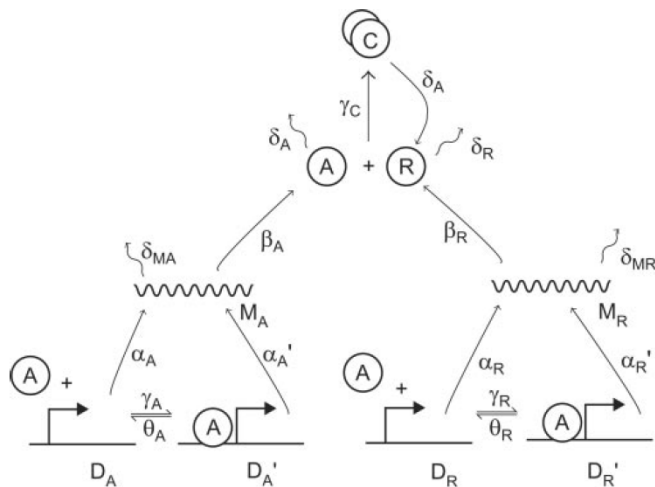
Negative values 10 \times magnified.

Marie Curie Intra-European Fellowship for Career Development

Scope:

- ▶ Analytical and computational methods for reaction-diffusion systems [Cotter, Vejchodsky, Erban, 2013]
- ▶ Models with and without stochastic effects [Erban, Chapman, Kevrekidis, Vejchodsky, 2009]
- ▶ Circadian rhythms – spatial aspects
- ▶ Skin pattern formation – unilateral regulation

Circadian rhythms – chemical reactions



$$\begin{aligned} \alpha_A &= 50 \text{ h}^{-1} \\ \alpha'_A &= 500 \text{ h}^{-1} \\ \alpha_R &= 0.01 \text{ h}^{-1} \\ \alpha'_R &= 50 \text{ h}^{-1} \\ \beta_A &= 50 \text{ h}^{-1} \\ \beta_R &= 5 \text{ h}^{-1} \\ \gamma_A &= 1 \text{ mol}^{-1} \text{ h}^{-1} \\ \gamma_R &= 1 \text{ mol}^{-1} \text{ h}^{-1} \\ \gamma_C &= 2 \text{ mol}^{-1} \text{ h}^{-1} \\ \delta_A &= 1 \text{ h}^{-1} \\ \delta_R &= 0.2 \text{ h}^{-1} \\ \delta_{M_A} &= 10 \text{ h}^{-1} \\ \delta_{M_R} &= 0.5 \text{ h}^{-1} \\ \theta_A &= 50 \text{ h}^{-1} \\ \theta_R &= 100 \text{ h}^{-1} \end{aligned}$$

[Vilar et al, 2002]

Circadian rhythms – equations

Law of mass action:

$$d\bar{D}_A/dt = \theta_A \bar{D}'_A - \gamma_A \bar{D}_A \bar{A}$$

$$d\bar{D}'_A/dt = -\theta_A \bar{D}'_A + \gamma_A \bar{D}_A \bar{A}$$

$$d\bar{D}_R/dt = \theta_R \bar{D}'_R - \gamma_R \bar{D}_R \bar{A}$$

$$d\bar{D}'_R/dt = -\theta_R \bar{D}'_R + \gamma_R \bar{D}_R \bar{A}$$

$$d\bar{M}_A/dt = \alpha'_A \bar{D}'_A + \alpha_A \bar{D}_A - \delta_{M_A} \bar{M}_A$$

$$d\bar{M}_R/dt = \alpha'_R \bar{D}'_R + \alpha_R \bar{D}_R - \delta_{M_R} \bar{M}_R$$

$$d\bar{A}/dt = \beta_A \bar{M}_A + \theta_A \bar{D}'_A + \theta_R \bar{D}'_R \\ - \bar{A}(\gamma_A \bar{D}_A + \gamma_R \bar{D}_R + \gamma_C \bar{R} + \delta_A)$$

$$d\bar{R}/dt = \beta_R \bar{M}_R - \gamma_C \bar{A} \bar{R} + \delta_A \bar{C} - \delta_R \bar{R}$$

$$d\bar{C}/dt = \gamma_C \bar{A} \bar{R} - \delta_A \bar{C}$$

Initial conditions:

$$\bar{D}_A = \bar{D}_R = 1 \text{ mol}$$

$$\bar{D}'_A = \bar{D}'_R = \bar{M}_A = \bar{M}_R = \bar{A} = \bar{R} = \bar{C} = 0 \text{ mol}$$

Circadian rhythms – equations

Law of mass action:

$$d\bar{D}_A/dt = \theta_A - (\theta_A + \gamma_A\bar{A})\bar{D}_A$$

$$\bar{D}'_A = 1 - \bar{D}_A$$

$$d\bar{D}_R/dt = \theta_R - (\theta_R + \gamma_R\bar{A})\bar{D}_R$$

$$\bar{D}'_R = 1 - \bar{D}_R$$

$$d\bar{M}_A/dt = \alpha'_A + (\alpha_A - \alpha'_A)\bar{D}_A - \delta_{M_A}\bar{M}_A$$

$$d\bar{M}_R/dt = \alpha'_R + (\alpha_R - \alpha'_R)\bar{D}_R - \delta_{M_R}\bar{M}_R$$

$$d\bar{A}/dt = \beta_A\bar{M}_A + \theta_A(1 - \bar{D}_A) + \theta_R(1 - \bar{D}_R) \\ - \bar{A}(\gamma_A\bar{D}_A + \gamma_R\bar{D}_R + \gamma_C\bar{R} + \delta_A)$$

$$d\bar{R}/dt = \beta_R\bar{M}_R - \gamma_C\bar{A}\bar{R} + \delta_A\bar{C} - \delta_R\bar{R}$$

$$d\bar{C}/dt = \gamma_C\bar{A}\bar{R} - \delta_A\bar{C}$$

Circadian rhythms – equations

Law of mass action:

$$dD_A/dt = \theta_A - (\theta_A + \gamma_A A)D_A$$

$$D'_A = 1 - D_A$$

$$dD_R/dt = \theta_R - (\theta_R + \gamma_R A)D_R$$

$$D'_R = 1 - D_R$$

$$\partial M_A/\partial t = \alpha'_A + (\alpha_A - \alpha'_A)D_A - \delta_{M_A}M_A + d_{M_A}\partial^2 M_A/\partial x^2$$

$$\partial M_R/\partial t = \alpha'_R + (\alpha_R - \alpha'_R)D_R - \delta_{M_R}M_R + d_{M_R}\partial^2 M_R/\partial x^2$$

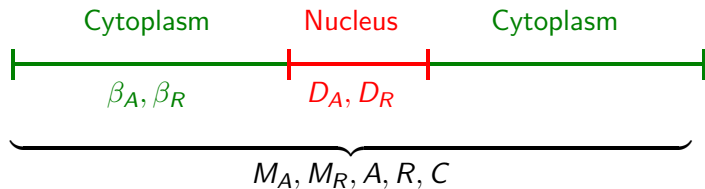
$$\begin{aligned} \partial A/\partial t = & \beta_A M_A + \theta_A(1 - D_A) + \theta_R(1 - D_R) \\ & - A(\gamma_A D_A + \gamma_R D_R + \gamma_C R + \delta_A) + d_A\partial^2 A/\partial x^2 \end{aligned}$$

$$\partial R/\partial t = \beta_R M_R - \gamma_C A R + \delta_A C - \delta_R R + d_R\partial^2 R/\partial x^2$$

$$\partial C/\partial t = \gamma_C A R - \delta_A C + d_C\partial^2 C/\partial x^2$$

No flux boundary conditions

Concentration: $D_A = \bar{D}_A/\nu$, $D_R = \bar{D}_R/\nu$, ... $\nu = 1$ cell



Cell size:

- ▶ $L_{\text{cell}} = 10\text{--}100 \mu\text{m}$

Diffusivities:

- ▶ Proteins:

$$d_A = d_R = d_C = 20\,000 \mu\text{m}^2\text{h}^{-1} = 20\,000/L_{\text{cell}}^2 \text{ cell}^2\text{h}^{-1}$$

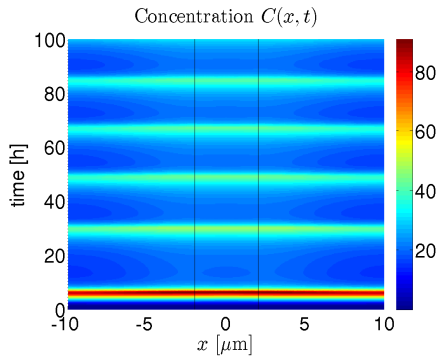
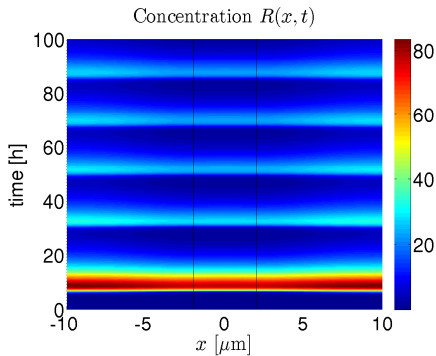
(measurements [Nenninger 2010]: $\approx 14\,400\text{--}36\,000 \mu\text{m}^2\text{h}^{-1}$)

- ▶ mRNA:

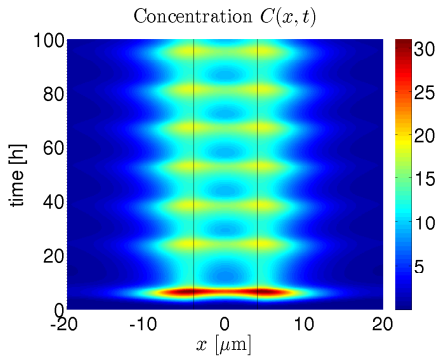
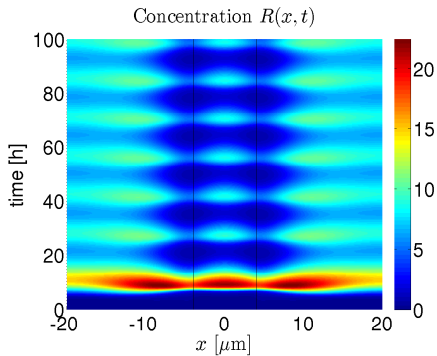
$$d_{M_A} = d_{M_R} = d_A/\sqrt[3]{10}$$

(mRNA is roughly $10\times$ bigger than protein)

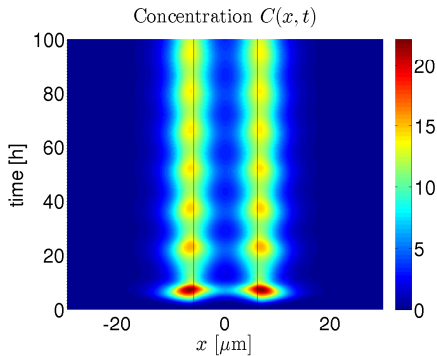
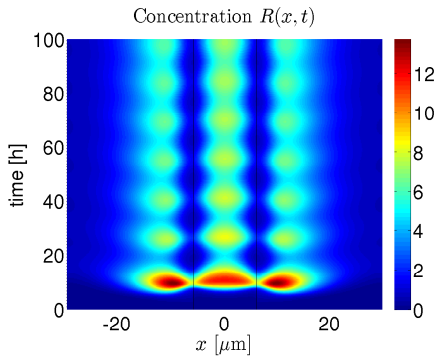
$$L_{\text{cell}} = 20 \mu\text{m}$$



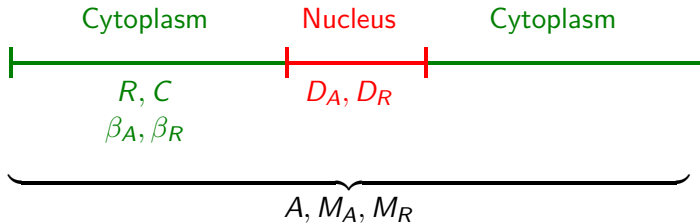
$$L_{\text{cell}} = 40 \mu\text{m}$$



$$L_{\text{cell}} = 60 \mu\text{m}$$

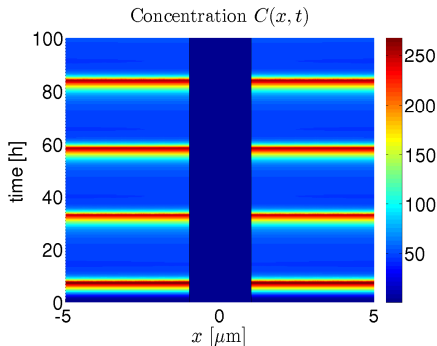
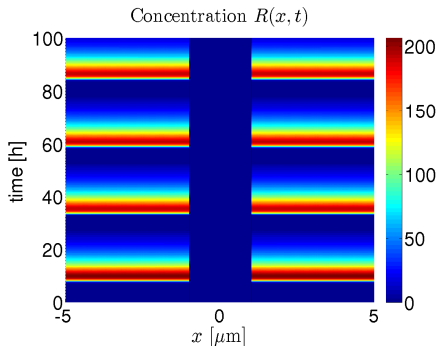


R and C in cytoplasm only



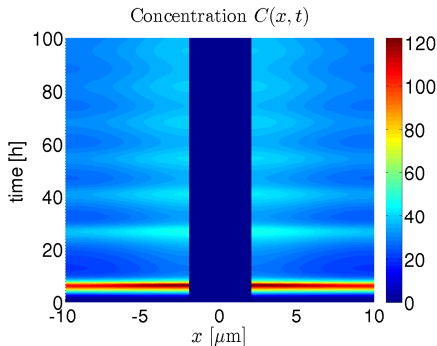
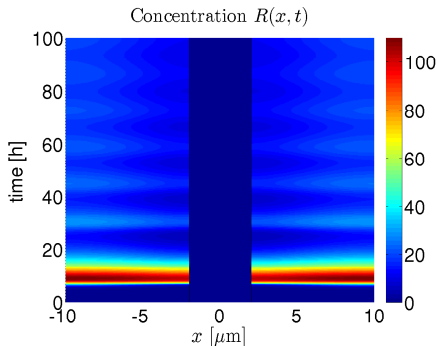
R and C in cytoplasm only – results

$$L_{\text{cell}} = 10 \mu\text{m}$$



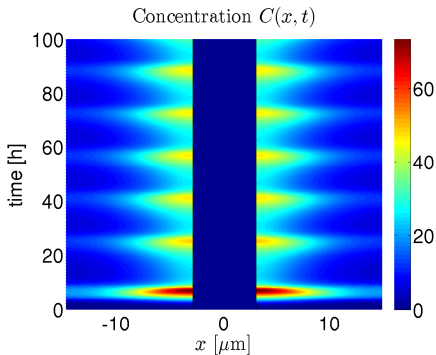
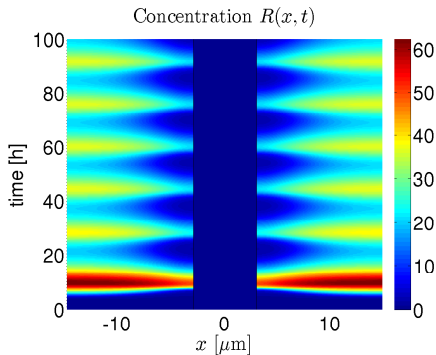
R and C in cytoplasm only – results

$$L_{\text{cell}} = 20 \mu\text{m}$$



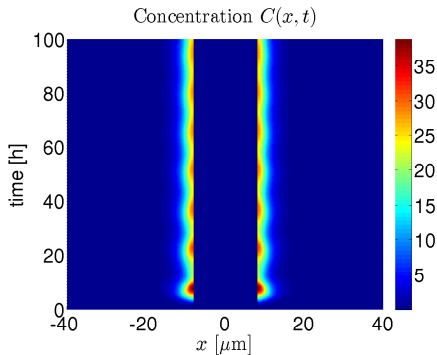
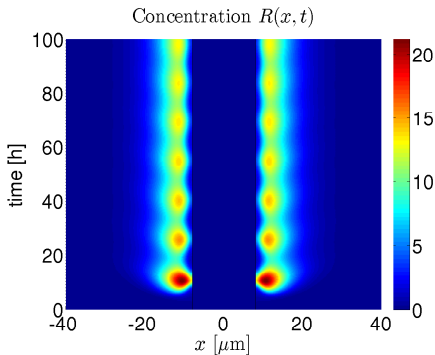
R and C in cytoplasm only – results

$$L_{\text{cell}} = 30 \mu\text{m}$$



R and C in cytoplasm only – results

$$L_{\text{cell}} = 80 \mu\text{m}$$



Reaction–diffusion system:

$$\left. \begin{aligned} \frac{\partial u}{\partial t} &= \delta_1 \Delta u + f(u, v) \\ \frac{\partial v}{\partial t} &= \delta_2 \Delta v + g(u, v) \end{aligned} \right\} \text{in } \Omega$$
$$\left. \begin{aligned} \frac{\partial u}{\partial n} &= 0 \\ \frac{\partial v}{\partial n} &= 0 \end{aligned} \right\} \text{on } \partial\Omega$$

Patterns for $\frac{\delta_1}{\delta_2} < 1$

Idea: add a unilateral regulation to the Turing's mechanism

Reaction–diffusion system with Signorini b.c.:

$$\left. \begin{aligned} \frac{\partial u}{\partial t} &= \delta_1 \Delta u + f(u, v) \\ \frac{\partial v}{\partial t} &= \delta_2 \Delta v + g(u, v) \end{aligned} \right\} \text{in } \Omega \quad \left. \begin{aligned} \frac{\partial u}{\partial n} &= 0 \\ v &\geq 0, \frac{\partial v}{\partial n} \geq 0, v \frac{\partial v}{\partial n} = 0 \end{aligned} \right\} \text{on } \partial\Omega$$

Patterns even for $\frac{\delta_1}{\delta_2} \approx 1$ [Kučera, Väh, 2012]

Idea: add a unilateral regulation to the Turing's mechanism

Reaction–diffusion system with unilateral source:

$$\left. \begin{aligned} \frac{\partial u}{\partial t} &= \delta_1 \Delta u + f(u, v) \\ \frac{\partial v}{\partial t} &= \delta_2 \Delta v + g(u, v) + \gamma v^- \end{aligned} \right\} \text{in } \Omega \quad \left. \begin{aligned} \frac{\partial u}{\partial n} &= 0 \\ \frac{\partial v}{\partial n} &= 0 \end{aligned} \right\} \text{on } \partial\Omega$$

Idea: add a unilateral regulation to the Turing's mechanism

Reaction–diffusion system with unilateral source:

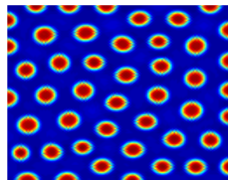
$$\left. \begin{aligned} \frac{\partial u}{\partial t} &= \delta_1 \Delta u + f(u, v) \\ \frac{\partial v}{\partial t} &= \delta_2 \Delta v + g(u, v) + \gamma v^- \end{aligned} \right\} \text{in } \Omega \quad \left. \begin{aligned} \frac{\partial u}{\partial n} &= 0 \\ \frac{\partial v}{\partial n} &= 0 \end{aligned} \right\} \text{on } \partial\Omega$$

Numerical experiments

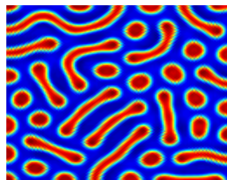
$$f(u, v) = \alpha u + v - r_2 uv - \alpha r_3 uv^2$$

$$g(u, v) = -\alpha u + \beta v + r_2 uv + \alpha r_3 uv^2$$

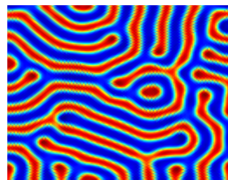
[Liu, Liaw, Maini, 2006]



$$\beta = -0.97$$
$$\gamma = 0.00$$



$$\beta = -0.89$$
$$\gamma = 0.08$$



$$\beta = -0.80$$
$$\gamma = 0.17$$

- ▶ Philip K. Maini
- ▶ Radek Erban
- ▶ Simon Cotter
- ▶ Shuohao Liao – Higher-dimensional Fokker-Planck equation

- ▶ Milan Kučera
- ▶ Filip Jaroš
- ▶ Martin Väh

Circadian rhythms

- ▶ Analysis of the spatial model
- ▶ Stochastic spatial model

Skin pattern formation

- ▶ Implementation of Signorini boundary conditions
- ▶ Another dynamics (Thomas system)

Marie Curie Fellowship, StochDetBioModel



EUROPEAN
COMMISSION

The research leading to these results has received funding from the People Programme (Marie Curie Actions) of the European Union's Seventh Framework Programme (FP7/2007-2013) under REA grant agreement no. 328008.

Thank you for your attention

Tomáš Vejchodský

Centre for Mathematical Biology
Mathematical Institute



CMB Group Meeting, Oxford, 10 June, 2013