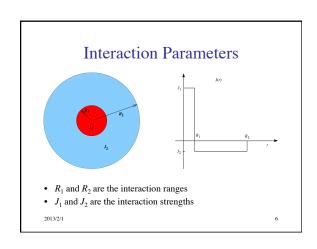


Activation & Inhibition in Pattern Formation • Color patterns typically have a characteristic length scale • Independent of cell size and animal size • Achieved by: - short-range activation ⇒ local uniformity - long-range inhibition ⇒ separation



CA Activation/Inhibition Model

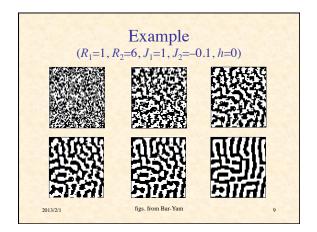
- Let states $s_i \in \{-1, +1\}$
- and h be a bias parameter
- and r_{ii} be the distance between cells i and j
- Then the state update rule is:

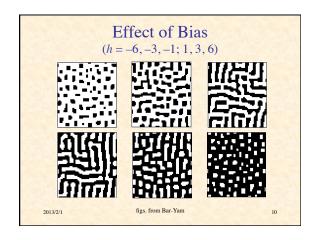
$$s_i(t+1) = \text{sign}\left[h + J_1 \sum_{r_{ij} < R_1} s_j(t) + J_2 \sum_{R_1 \le r_{ij} < R_2} s_j(t)\right]$$

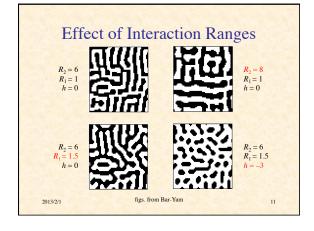
2013/2/1

Demonstration of NetLogo
Program for Activation/Inhibition
Pattern Formation:
Fur

RunAICA.nlogo







Differential Interaction Ranges

- How can a system using strictly local interactions discriminate between states at long and short range?
- E.g. cells in developing organism
- Can use two different *morphogens* diffusing at two different rates
 - activator diffuses slowly (short range)
 - inhibitor diffuses rapidly (long range)

2013/2/1

12

Digression on Diffusion

• Simple 2-D diffusion equation:

$$\dot{A}(x,y) = D\nabla^2 A(x,y)$$

• Recall the 2-D Laplacian:

$$\nabla^2 A(x,y) = \frac{\partial^2 A(x,y)}{\partial x^2} + \frac{\partial^2 A(x,y)}{\partial y^2}$$

- The Laplacian (like 2nd derivative) is:
 - positive in a local minimum
- negative in a local maximum

13

Reaction-Diffusion System

$$\frac{\partial A}{\partial t} = D_{A} \nabla^{2} A + f_{A}(A, I) \text{ reaction}$$

$$\frac{\partial I}{\partial t} = D_{I} \nabla^{2} I + f_{A}(A, I) + f_{A}(A, I) \text{ reaction}$$

$$\frac{\partial}{\partial t} \begin{pmatrix} A \\ I \end{pmatrix} = \begin{pmatrix} D_{A} & 0 \\ 0 & D_{I} \end{pmatrix} \begin{pmatrix} \nabla^{2} A \\ \nabla^{2} I \end{pmatrix} + \begin{pmatrix} f_{A}(A, I) \\ f_{I}(A, I) \end{pmatrix}$$

$$\dot{\mathbf{c}} = \mathbf{D} \nabla^{2} \mathbf{c} + \mathbf{f}(\mathbf{c}), \text{ where } \mathbf{c} = \begin{pmatrix} A \\ I \end{pmatrix}$$
2013/2/1

General Reaction-Diffusion System

$$\frac{\partial c_i}{\partial t} = \sum_{\alpha} k_{\alpha} \mathbf{v}_{i\alpha} \left(\prod_{k=1}^n c_k^{m_{k\alpha}} \right) - \nabla \cdot \mathbf{j}_i$$

where $\mathbf{j}_i = \vec{\mu}_i c_i - \mathbf{div} \ \mathbf{D}_i c_i$ (flux)

where k_{α} = rate constant for reaction α

and $v_{i\alpha}$ = stoichiometric coefficient

and $m_{k\alpha}$ = a non-negative integer

and $\vec{\mu}_i$ = drift vector

and \mathbf{D}_i = diffusivity matrix

where **div** $\mathbf{D}c = \sum_{i} \mathbf{e}_{j} \sum_{k} D_{jk} \frac{\partial c}{\partial x_{k}}$

2013/2/1

Framework for Complexity

- change = source terms + transport terms
- source terms = local coupling = interactions local to a small region
- transport terms = spatial coupling
 - = interactions with contiguous regions
 - = advection + diffusion
 - advection: non-dissipative, time-reversible
 - diffusion: dissipative, irreversible

013/2/1

Continuous-time Activator-Inhibitor System

- Activator A and inhibitor I may diffuse at different rates in x and y directions
- Cell becomes more active if activator + bias exceeds inhibitor
- · Otherwise, less active

$$\frac{\partial A}{\partial t} = d_{Ax} \frac{\partial^2 A}{\partial x^2} + d_{Ay} \frac{\partial^2 A}{\partial y^2} + k_A (A + B - I)(1 - A)$$

$$\frac{\partial I}{\partial t} = d_{1x} \frac{\partial^2 I}{\partial x^2} + d_{1y} \frac{\partial^2 I}{\partial y^2} + k_1 (A + B - I)(1 - I)$$

2013/2/1

17

NetLogo Simulation of Reaction-Diffusion System

- 1. Diffuse activator in X and Y directions
- 2. Diffuse inhibitor in X and Y directions
- 3. Each patch performs:

stimulation = bias + activator – inhibitor + noise if stimulation > 0 then

set activator and inhibitor to 100

else

set activator and inhibitor to 0

2013/2/1

18

Demonstration of NetLogo Program for Activator/Inhibitor Pattern Formation

Run Pattern.nlogo

3/2/1

Demonstration of NetLogo Program for Activator/Inhibitor Pattern Formation with Continuous State Change

Run Activator-Inhibitor.nlogo

2013/2/1 20

Turing Patterns

- Alan Turing studied the mathematics of reaction-diffusion systems
- Turing, A. (1952). The chemical basis of morphogenesis. *Philosophical Transactions of the Royal Society* **B 237**: 37–72.
- The resulting patterns are known as *Turing* patterns

2013/2/1 21

A Key Element of Self-Organization

- · Activation vs. Inhibition
- · Cooperation vs. Competition
- · Amplification vs. Stabilization
- Growth vs. Limit
- Positive Feedback vs. Negative Feedback
 - Positive feedback creates
 - Negative feedback shapes

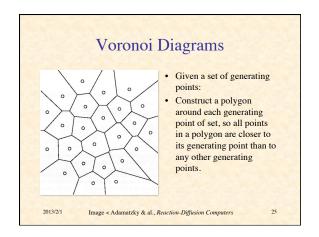
013/2/1 22

Reaction-Diffusion Computing

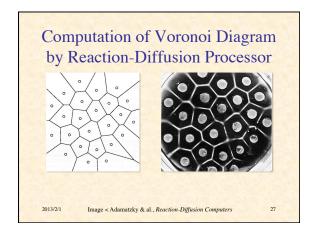
- Has been used for image processing
 - diffusion ⇒ noise filtering
 - reaction ⇒ contrast enhancement
- Depending on parameters, RD computing
 - restore broken contours
 - detect edges
 - improve contrast

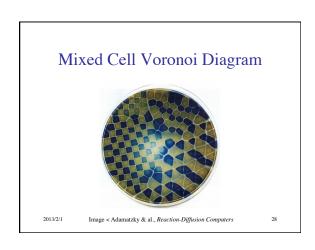
013/2/1 23

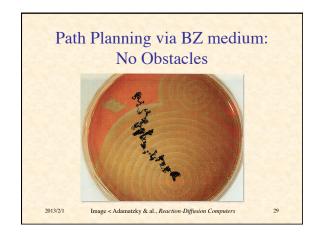
Image Processing in BZ Medium A C (A) boundary detection, (B) contour enhancement, (C) shape enhancement, (D) feature enhancement Image < Adamatzky, Comp. in Nonlinear Media & Autom. Coll. 24

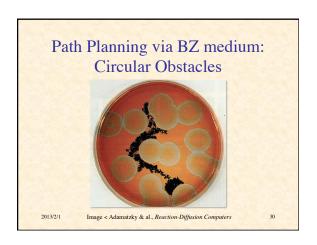


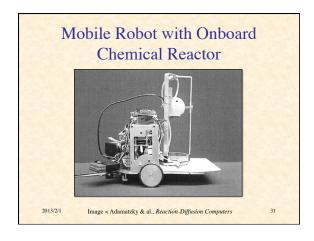
Some Uses of Voronoi Diagrams Collision-free path planning Determination of service areas for power substations Nearest-neighbor pattern classification Determination of largest empty figure

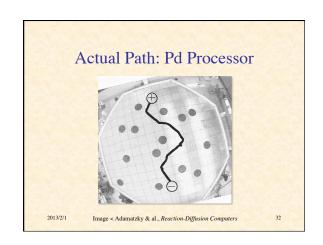


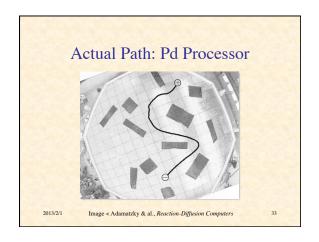


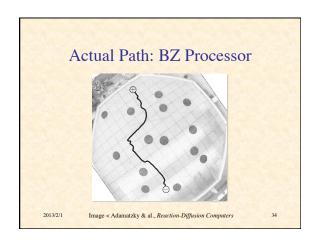












Bibliography for Reaction-Diffusion Computing 1. Adamatzky, Adam. Computing in Nonlinear Media and Automata Collectives. Bristol: Inst. of Physics Publ., 2001. 2. Adamatzky, Adam, De Lacy Costello, Ben, & Asai, Tetsuya. Reaction Diffusion Computers. Amsterdam: Elsevier, 2005.