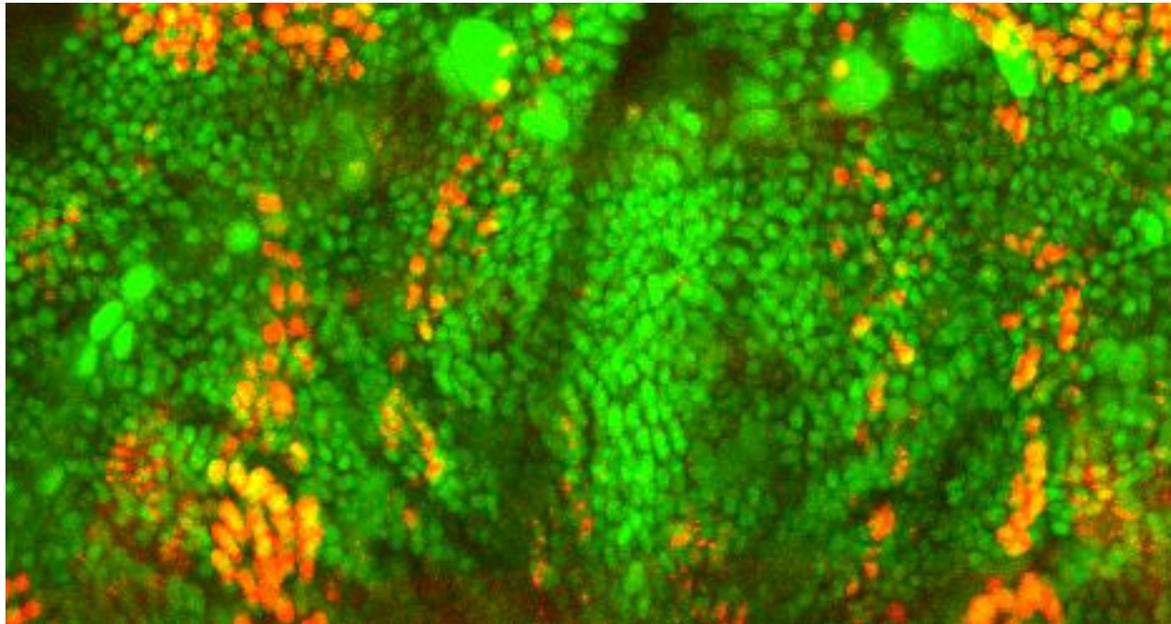


# Robust computing over networks : Lessons from nature

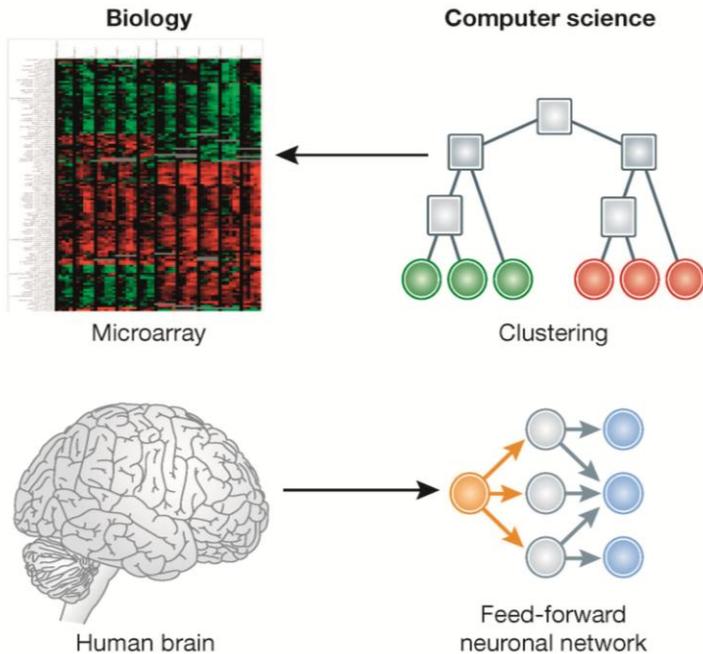
Ziv Bar-Joseph

Lane Center for Computational Biology & Machine Learning  
School of Computer Science  
Carnegie Mellon University

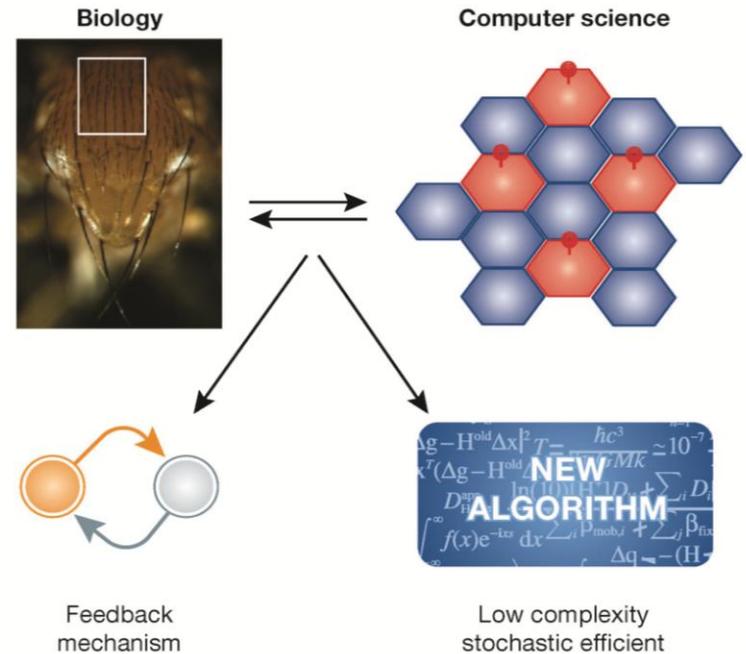


# Bi-directional studies

## A Traditional studies



## B Computational thinking

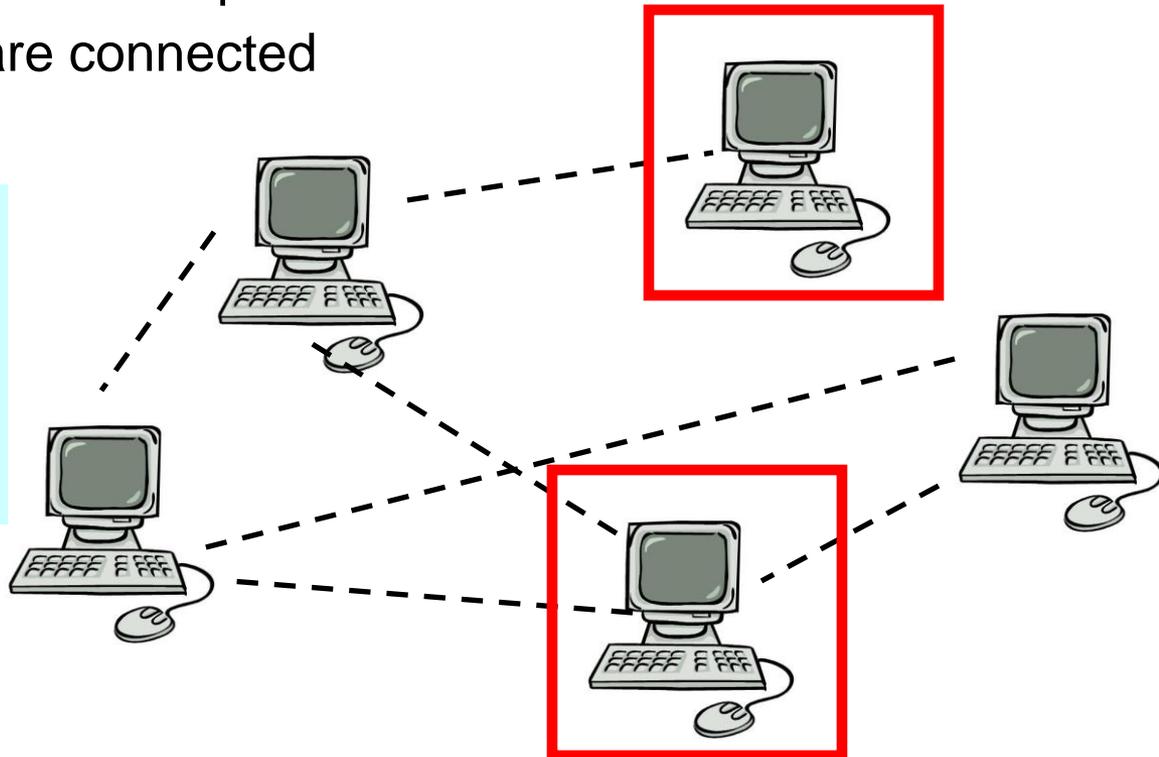


MIS and fly development

# Maximal Independent Set (MIS)

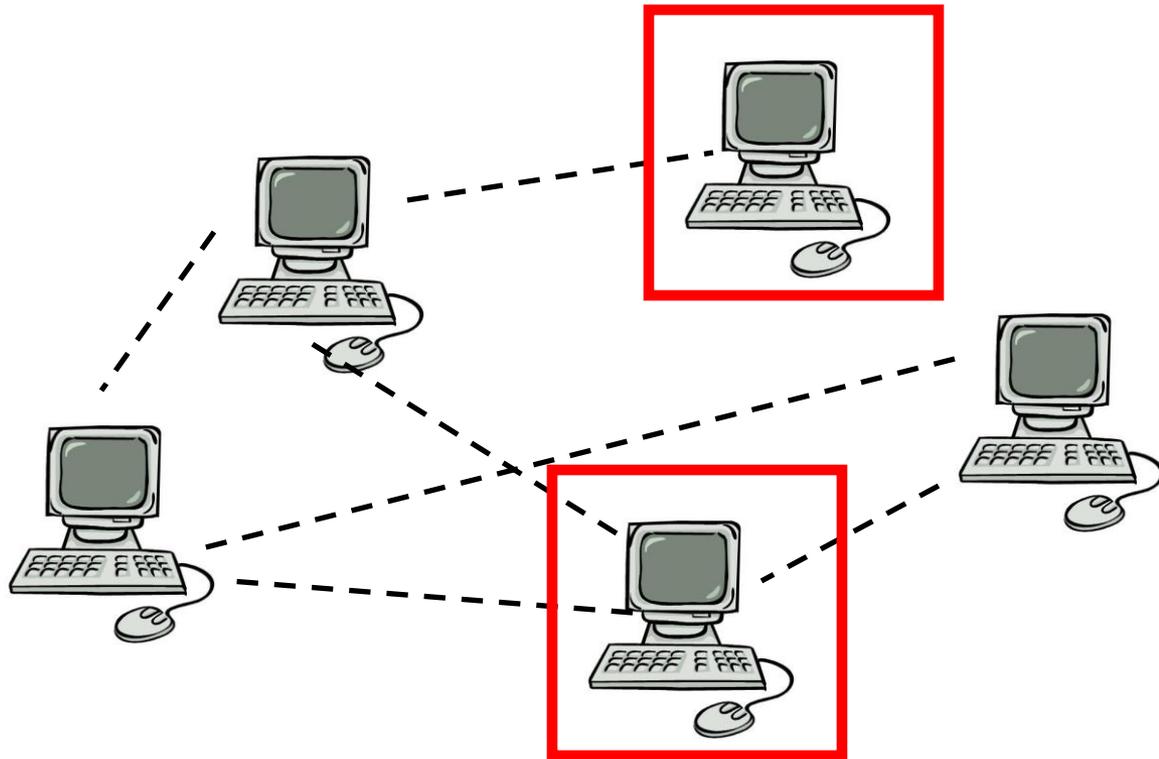
- A fundamental problem in distributed computing:
  - Often required to establish wireless networks
  - Used for routing messages, grouping sensors etc.
- For a set of nodes select a subset  $A$  such that:
  - Each processor is connected to a processor in  $A$
  - No two processors in  $A$  are connected

“it is difficult to see how this problem can be solved in substantially fewer stages such as  $O(\sqrt{n}) \dots$ ” (Valiant 1982)



# Maximal Independent Set (MIS)

- Fast algorithms (e.g. Luby and Alon et al) exist for distributively selecting the MIS set but:
  - They Assume nodes know the status of their neighbors and also the topology of the graph (which is changing)
  - Use large messages



# Algorithm for MIS

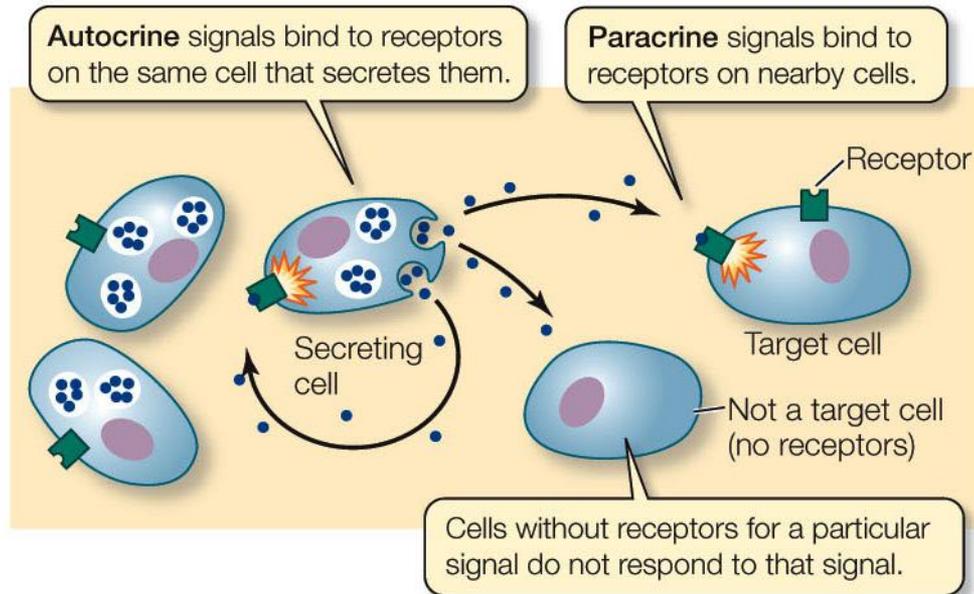
Each process needs to know how many neighbors it has

- Algorithm (proceed in rounds)
  - Each processor flips a coin with probability  $1/d$  where  $d$  is the number of its neighbors
  - If result is 0, do nothing
  - If result is 1, send to all other processors
    - If no collisions, Leader; all processes exit
    - Otherwise process with highest number of neighbors wins and becomes leader

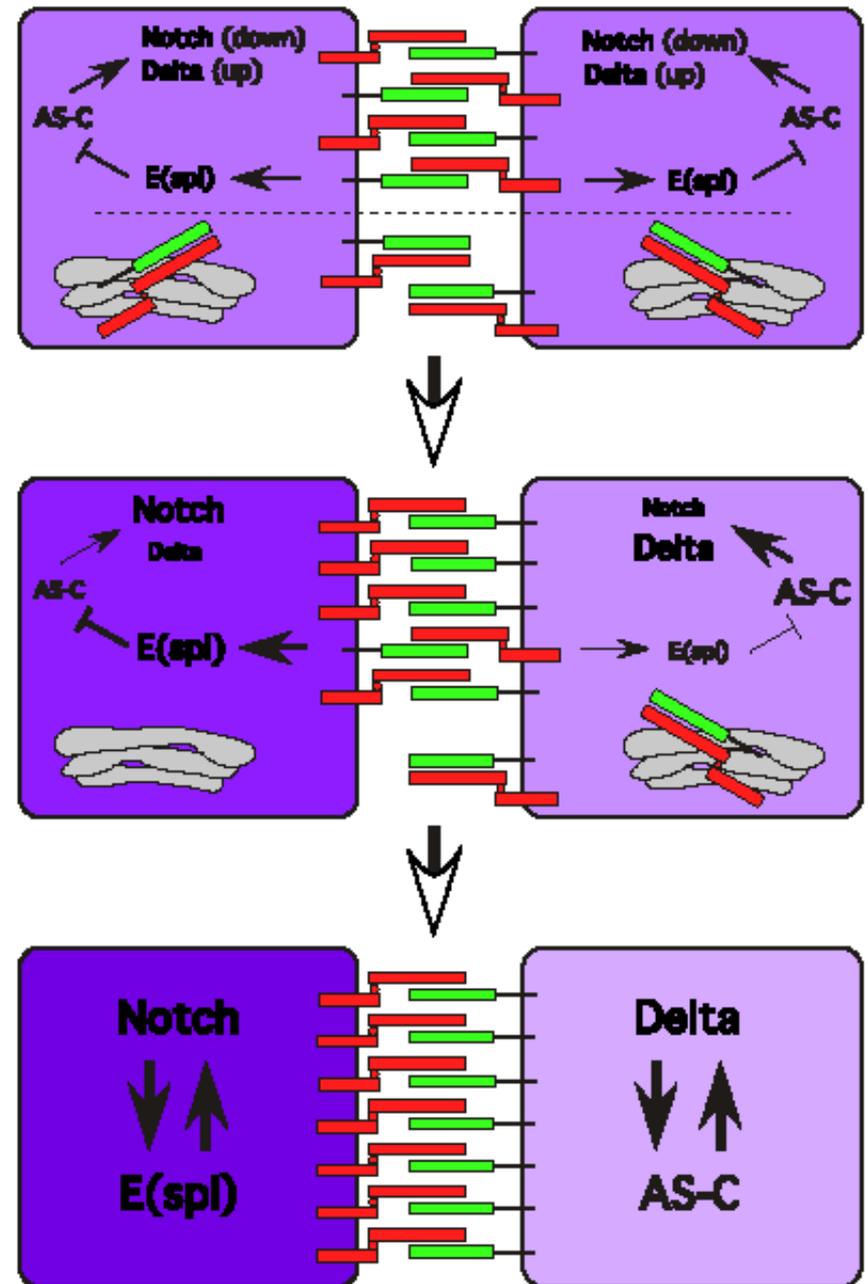
Each process needs to know how many neighbors its neighbors have

Luby, *SIAM J. Comput.* 1986  
Alon et al *J. Algorithms* 1986

# Cell signaling

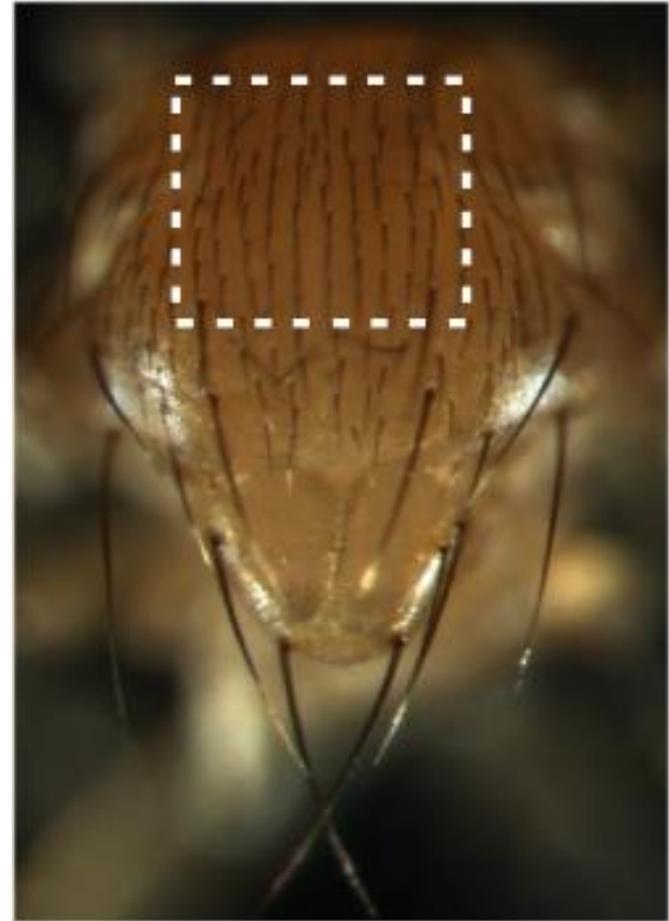


# Lateral inhibition: Notch-Delta signaling

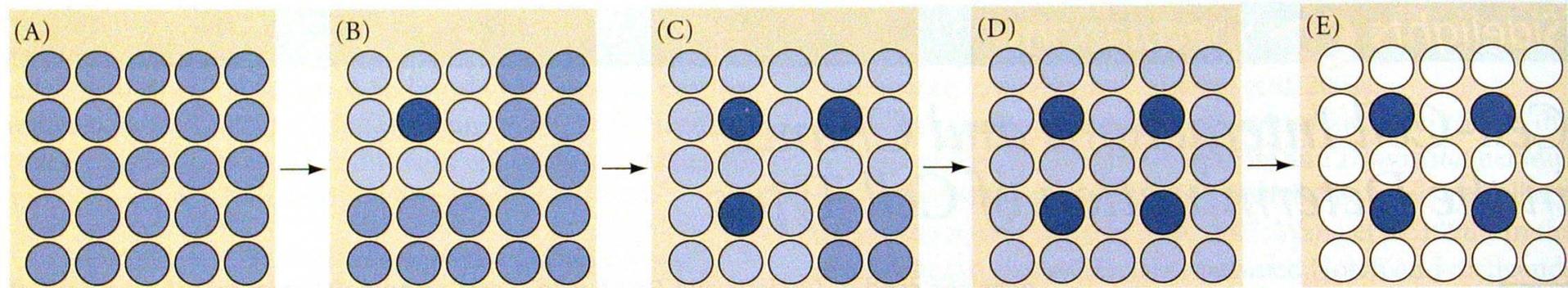


# SOP selection in flies

- When the fly's nervous system develops several cells are selected as sensory organ precursors (SOPs)
- These cells are later attached to the fly's sensory bristles
- Similar to the MIS requirements, in a highly accurate process each cell in a predefined cluster is either:
  - Selected as a SOP
  - Laterally inhibited by a neighboring SOP so it cannot become a SOP



# Generation of regular patterns of cells from equivalence groups



# Similarities between MIS and SOP selection

- Both are performed using a stochastic processes
  - Proven for MIS, experimentally validated for SOP
- Both are constrained by time
  - A cell that is not inhibited by certain time becomes a SOP
- Both only send messages if a node (cell) decides to join *A*
  - Reduces communication in computational systems, based on *cis* interactions for cells

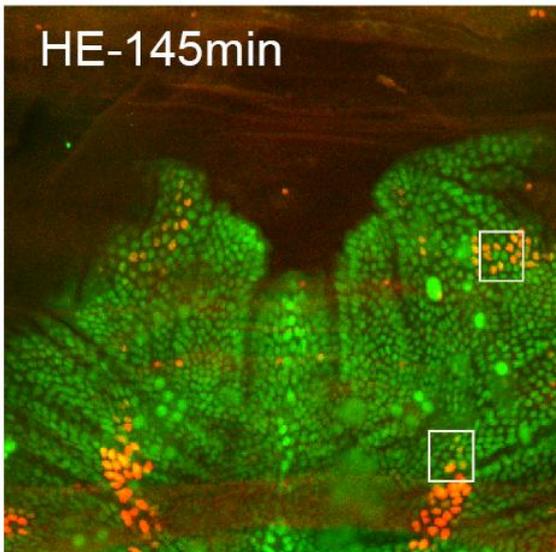
# Differences between SOP and MIS selection

- In SOP selection cells do not know the status of their neighbors and the overall topology
- Messages in SOP selection are binary

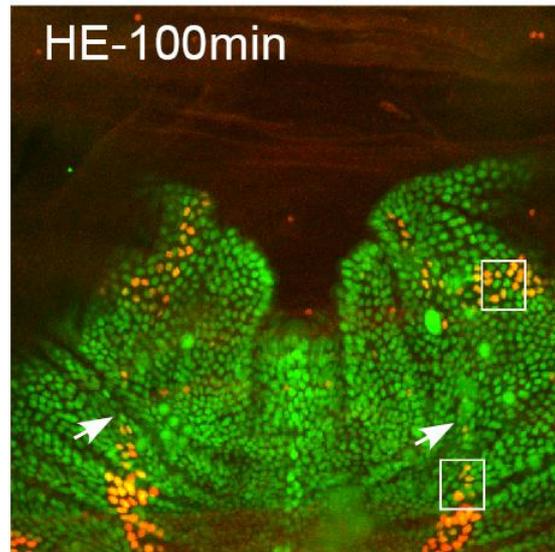
Can we improve current algorithms for MIS by understating how the biological process is performed?

# Movie

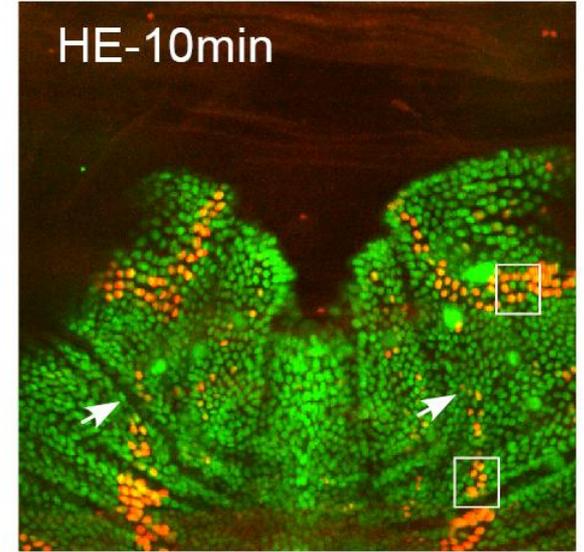
a1



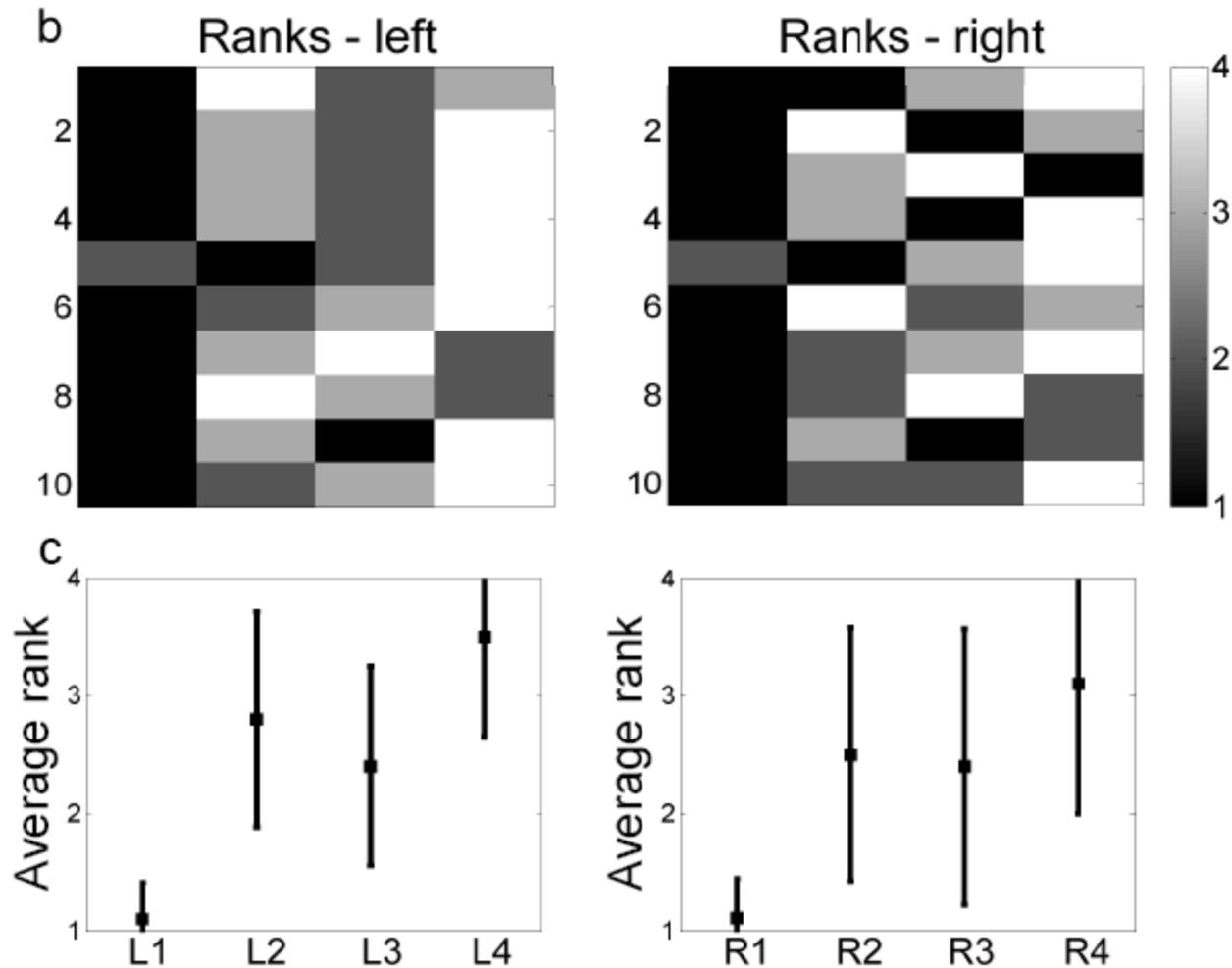
a2



a3

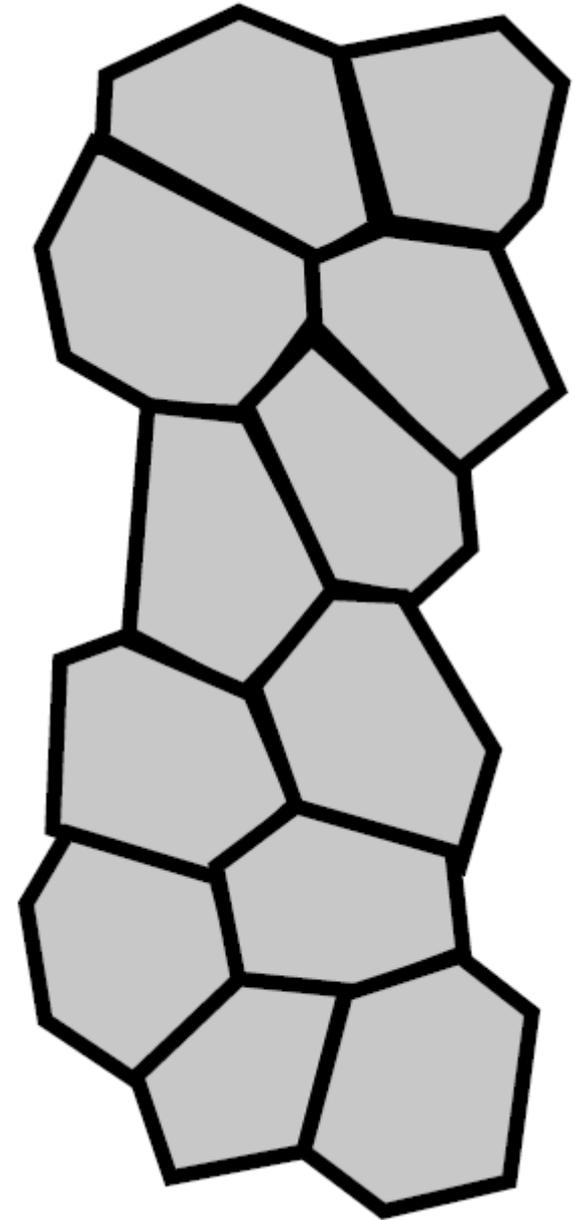


# Observation 1: SOP selection is stochastic



# Simulations

- 2 by 6 grid (also tried 2 by 7)
- Each cell touches all adjacent and diagonal neighbors



# Simulations

- All models assume a cell becomes a SOP by accumulating the protein Delta until it passes some threshold

## Four different models:

### 1. Accumulation

- Accumulating Delta based on a Gaussian distribution

### 2. Fixed Accumulation

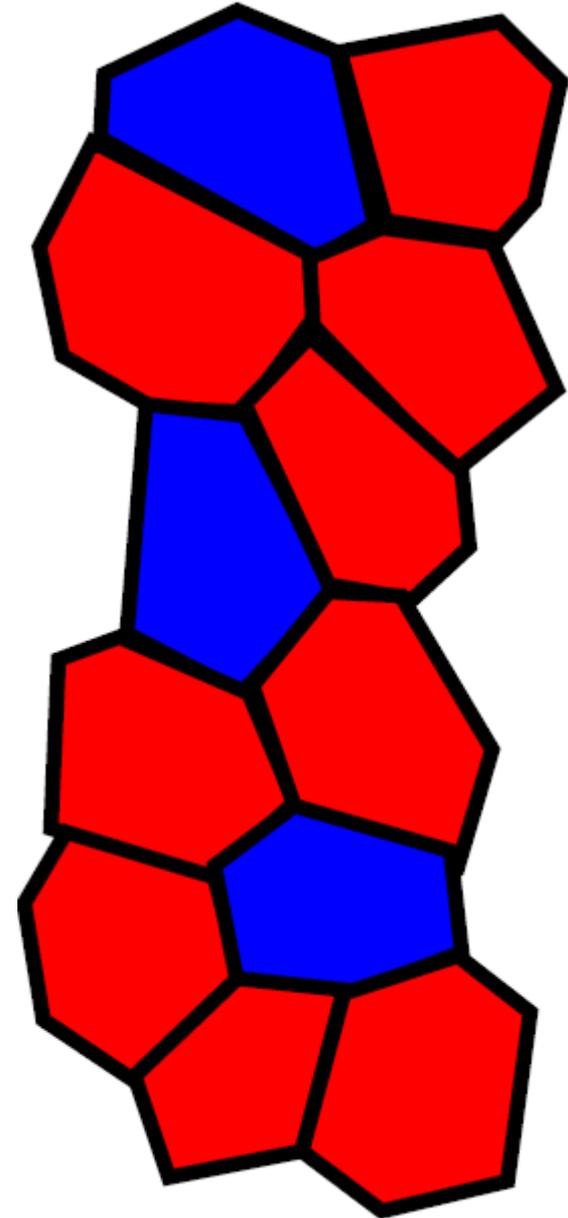
- Randomly select an accumulation rate only once

### 3. Rate Change

- Increase accumulation probability as time goes by by using feedback loop

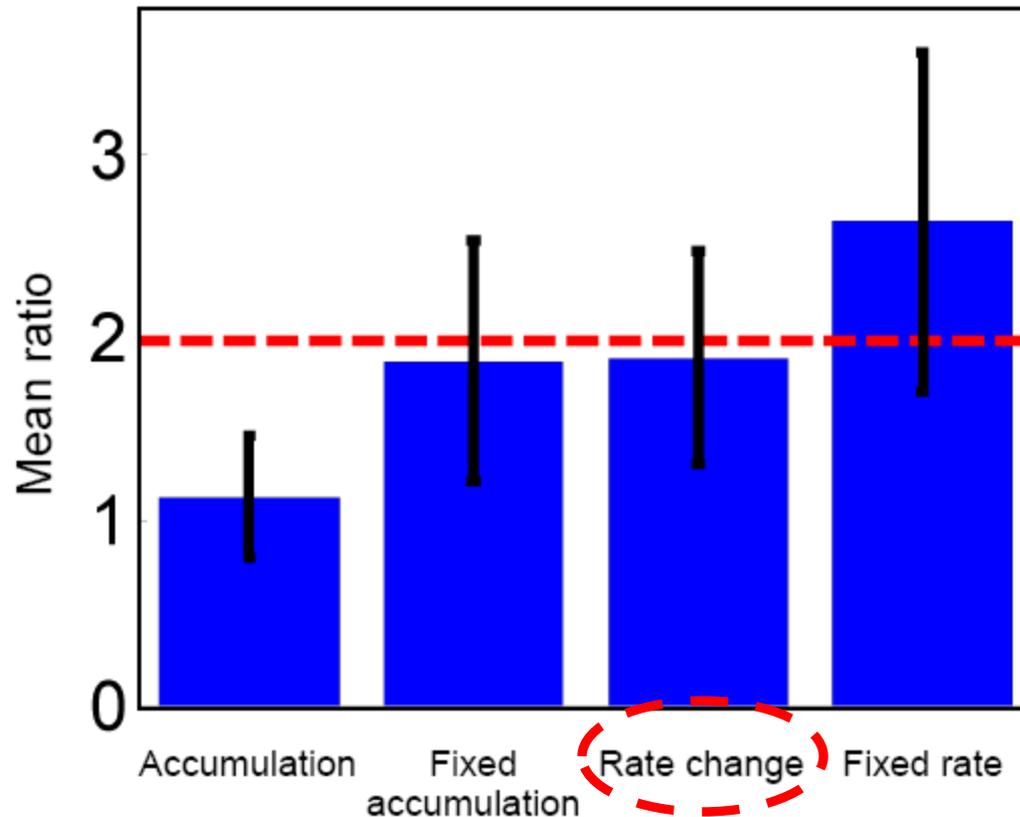
### 4. Fixed rate

- Fix accumulation probability, use the same probability in all rounds



# Observation 2: Comparing the time of experimental and simulated selection

Ratio between selection time differences



# MIS Algorithm (revised)

MIS Algorithm (n,D) // n – upper bound on number of nodes  
D - upper bound on number of neighbors

- $p = 1/D$
  - round = round +1
  - if round > log(n)
    - $p = p * 2$  ; round = 0 // we start a new phase
  - Each processor flips a coin with probability  $p$ 
    - If result is 0, do nothing
    - If result is 1, send to all other processors
      - If no collisions, Leader; all processes exit
      - Otherwise
-

# Why does it work?

- Can show that by phase  $i$  there are no processes with more  $n/2^i$  neighbors
- Overall running time is  $O(\log(n) \log(D))$  where  $D$  is an upper bound on the number of neighbors
- For grids this is as fast as the best known algorithm for this problem.
- Message complexity is also extremely low:  $O(n)$

## Can be extended to:

1. Continuous probability increases
2. Unsynchronized settings
3. Cases where no upper bound is given on degree
4. No collision detection

# (several) open problems

- Can we learn how to protect important nodes from the way cells rely on redundancy?
- Can we improve coordination among agents (or robots) based on bacterial quorum sensing?
- What other specific biological problems can gain from the information processing prescriptive, and how insights into these will improve computational algorithms?

# Thanks



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