TEL AUIU UNIVERSITY 👾 אוניברסיטת תל-אביב

Ki-net – 17.01.13 Complex Collective Navigation in Bacteria-

Inspired Models

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Bacteria tools

Chemotaxis



• Physical interactions

Communication

[Miller and Bassler 2001, Annual Reviews Microbiology]



Modeling Swarms

• <u>Chemotaxis</u>:



run and tumble

low concentration

high concentration

run time is a function of the temporal gradient

Model:

The angle depends on the temporal gradient



Modeling Swarms

Simple interactions



Navigation in complex terrain

The environment is constant in time and spatially noisy



 $\cos(\omega_1 x + \varphi_1) \cdot \cos(\omega_2 x + \varphi_2)$



Bacteria-Inspired Models

Adaptable Interactions

[Shklarsh el al. 2011, Plos Computational Biology] Elad Schneidman, Gil Ariel, Eshel Ben-Jacob

Cargo Carrying Swarms

[Shklarsh el al. 2012, Interface Focus] Alin Finkelshtein, Gil Ariel, Oren Kalisman, Colin Ingham, Eshel Ben-Jacob

Collective Dynamics

Gil Ariel, Adi Shklarsh, Oren Kalisman, Colin Ingham, Eshel Ben-Jacob











Adaptable interactions

$\vec{v}_i(t+\Delta t) = \vec{d}_i(t) + w_i(t) \cdot \vec{v}_i(t)$

adaptable weights

$w_i(t) = \begin{bmatrix} 1 & sign(grad_i) > 0 \\ 0 & else \end{bmatrix}$

We consider adaptable interactions as a biologicallyinspired design principle that will dynamically modify the behavior of the system in response to changes in the environment.

Simulations



independent agents

interacting agents

adaptable interactions















Distribution of search paths













Cargo Carrying Swarms Beads Conidia





Bacteria swarms carry cargo and react to external forces



[Ingham et al. 2011, PNAS]





Rope Model:







Rope Model:





Simulation of the rope model







Group path length







Simulation of the extended model



Lubricating fluid edge is modeled as a discrete dual phase field Edge blocks motion of agents Enough close agents move the edge

Bacteria movement dynamics





Intermediate level: movement dynamics of a group of

lubricating bacteria in an envelope

[pictures by Ingham and Ben-Jacob]

Bacteria dynamics in a branch



3 lane dynamics



collective chemotaxis







branch repulsion



[movie by Gil Ariel]

Group navigation by collective organization

The point of return predicts the direction



Modeling branch dynamics

Identifying the key mechanisms underlying the dynamics





represents local cluster of bacteria coarse graining of bacteria branch

a dynamic boundary represents lubricating fluid limit

Self propelled agents



Dynamic envelope





A phenomenological expression to the speed of the boundary

$$\frac{d\gamma_n(s)}{dt} \simeq \hat{v}_i \times v_i \times \nabla n$$

Simulation

Moving branch with three lanes



Static branch with two lanes



Agreement with experiments – velocity, vorticity, and order parameter



How can local interaction rules control the direction of the branch?

agents which become closer to the food source increase their speed

An uneven accumulation of cells on the side of the branch tip closer to the food source results in shifting of the tip to the other side



Summary

• Adaptable interactions improve

efficiency in a navigation task

Complex behavior in cargo

carrying swarms

Navigation by self organization

in bacteria branches





Links:

- <u>https://sites.google.com/site/adishklarsh/</u>
- "Smart Swarms of Bacteria-Inspired Agents with Performance Adaptable Interactions"
 Shklarsh *et al.* PLoS Computational Biology
 "Collective Navigation of Cargo-Carrying Swarms"
- Shklarsh et al. Interface Focus