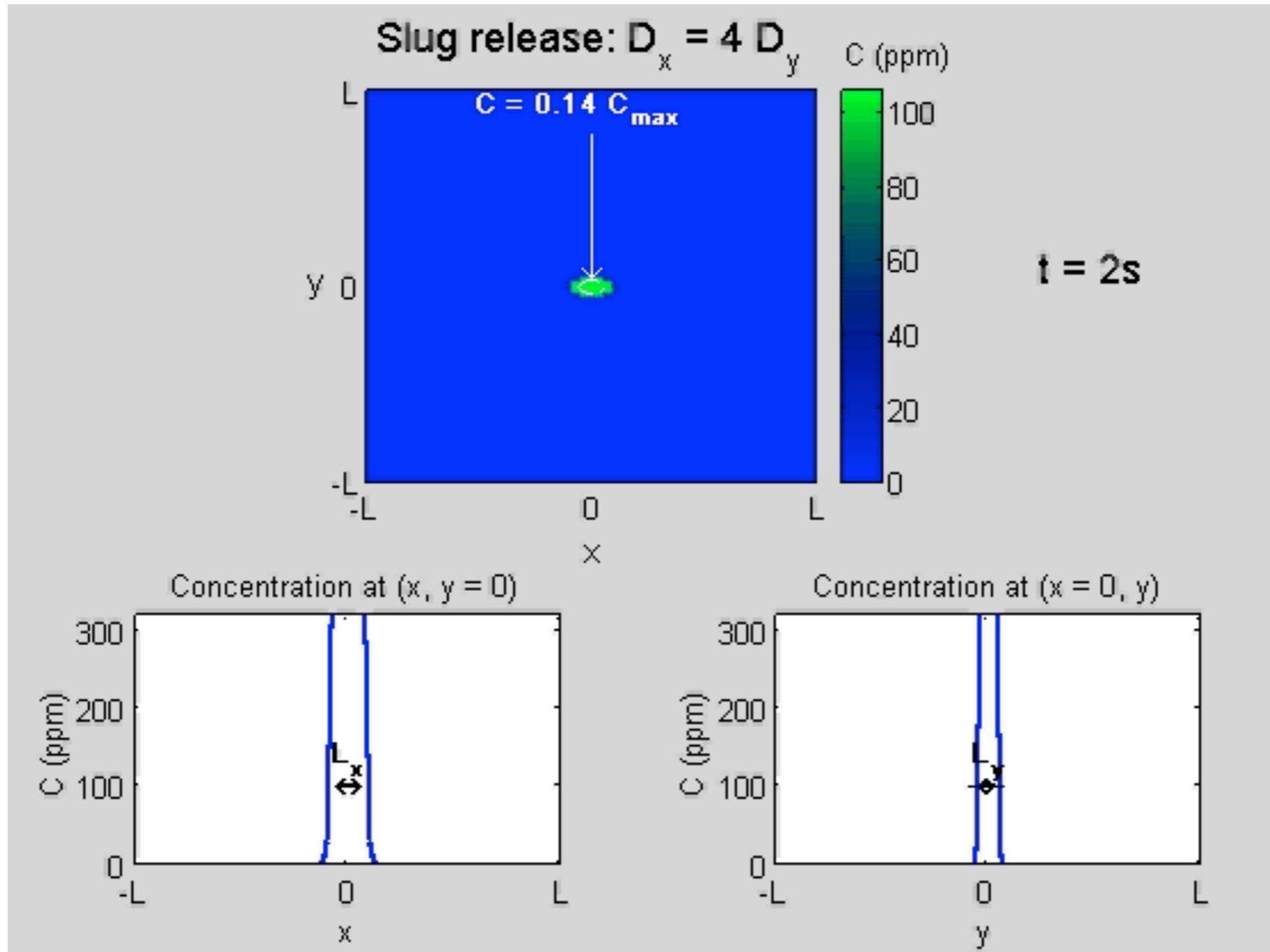


Beyond diffusion: Pattern formation

18.354 - L07

2D diffusion equation



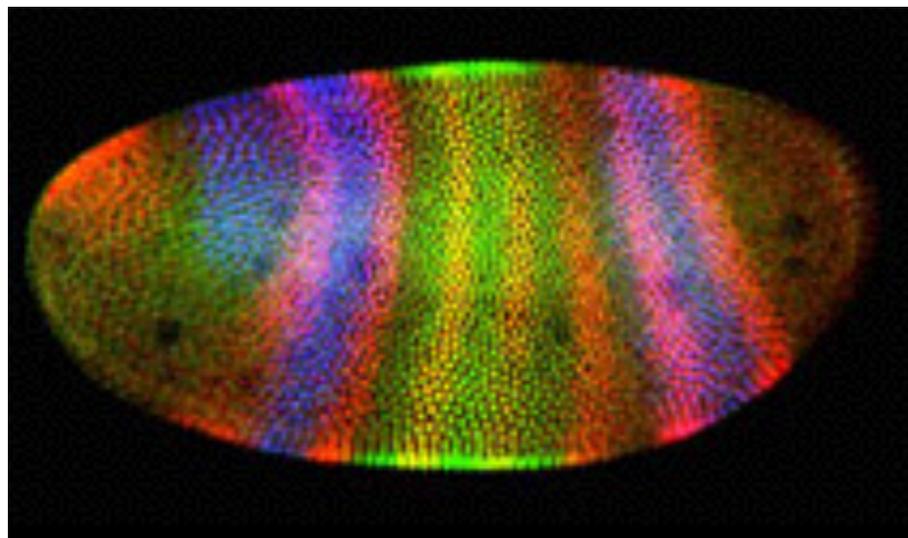
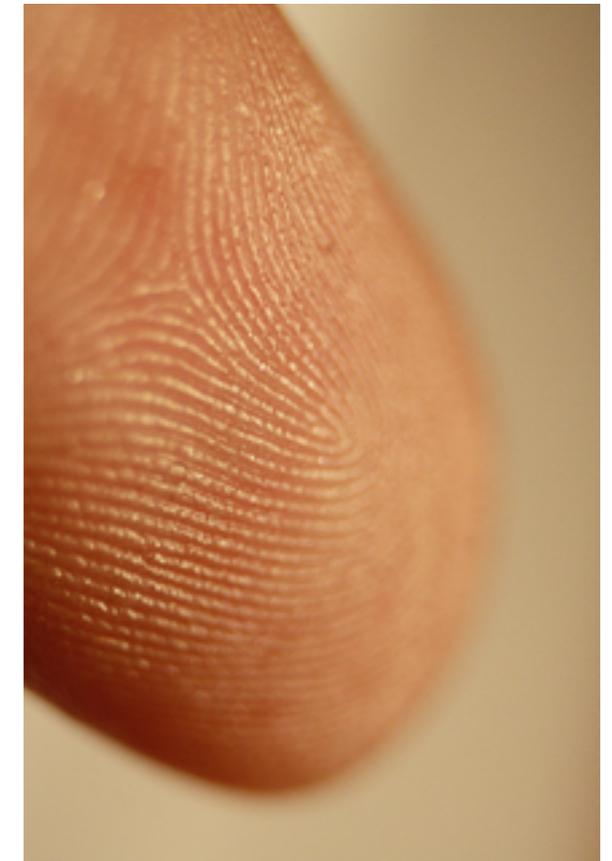
Pattern Formation and Dynamics in Nonequilibrium Systems



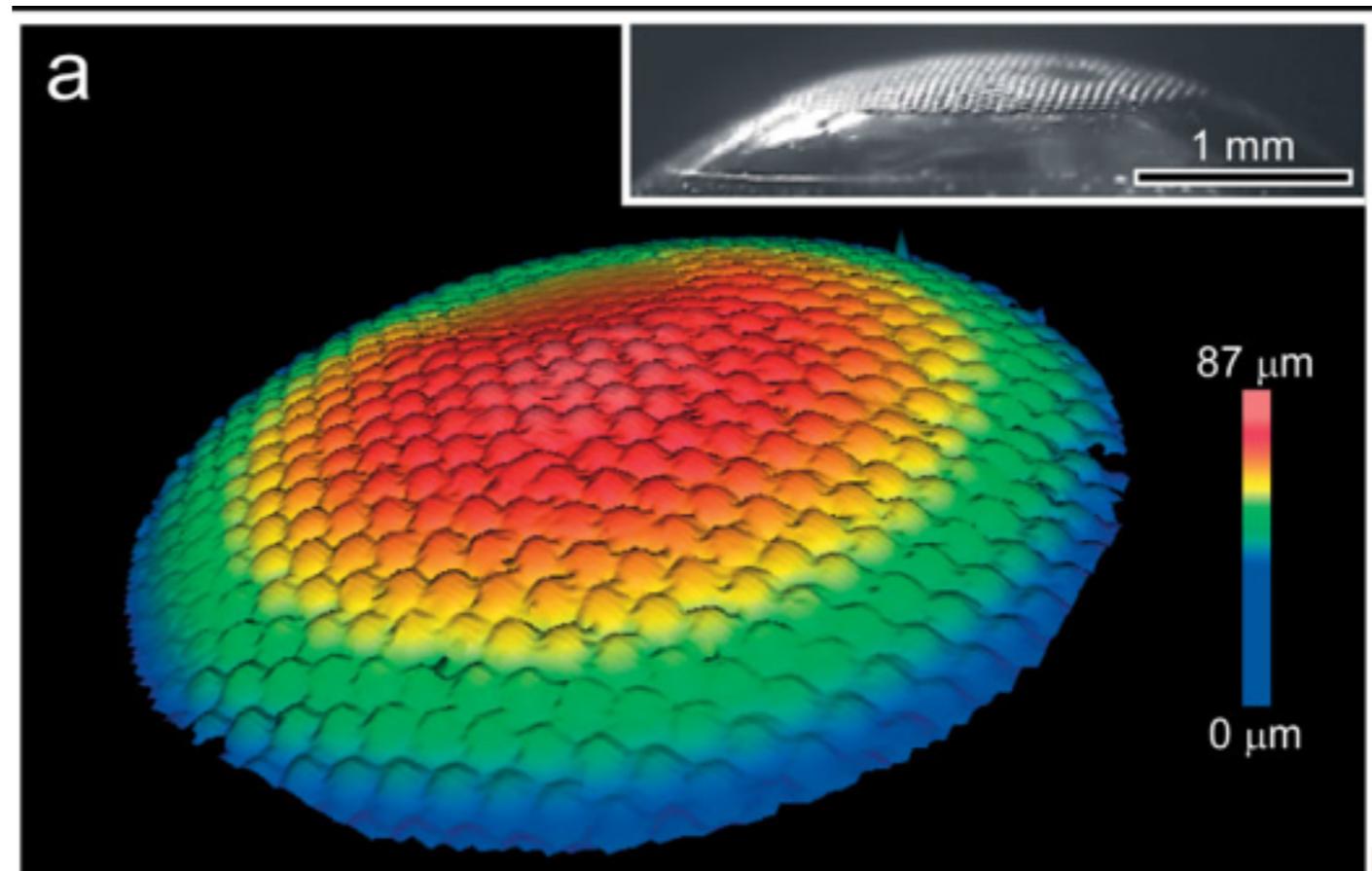
Michael Cross and Henry Greenside

CAMBRIDGE

Natural & artificial patterns



Paddock (2001) Biotechniques



Chan & Crosby (2006) Advanced Materials



Plankton

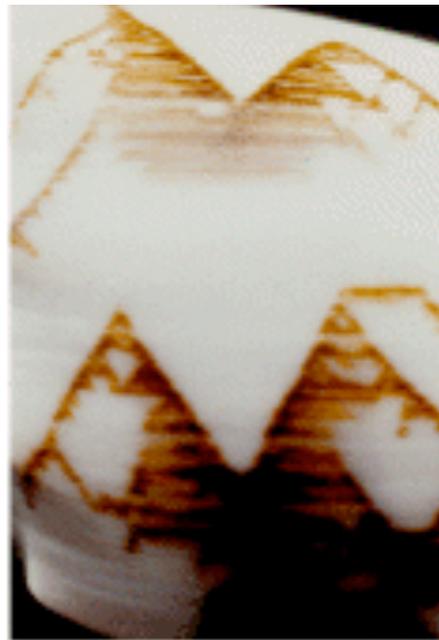
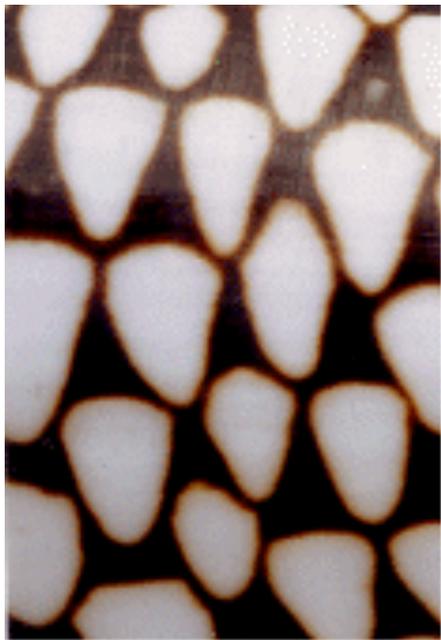


ESA cost of Ireland

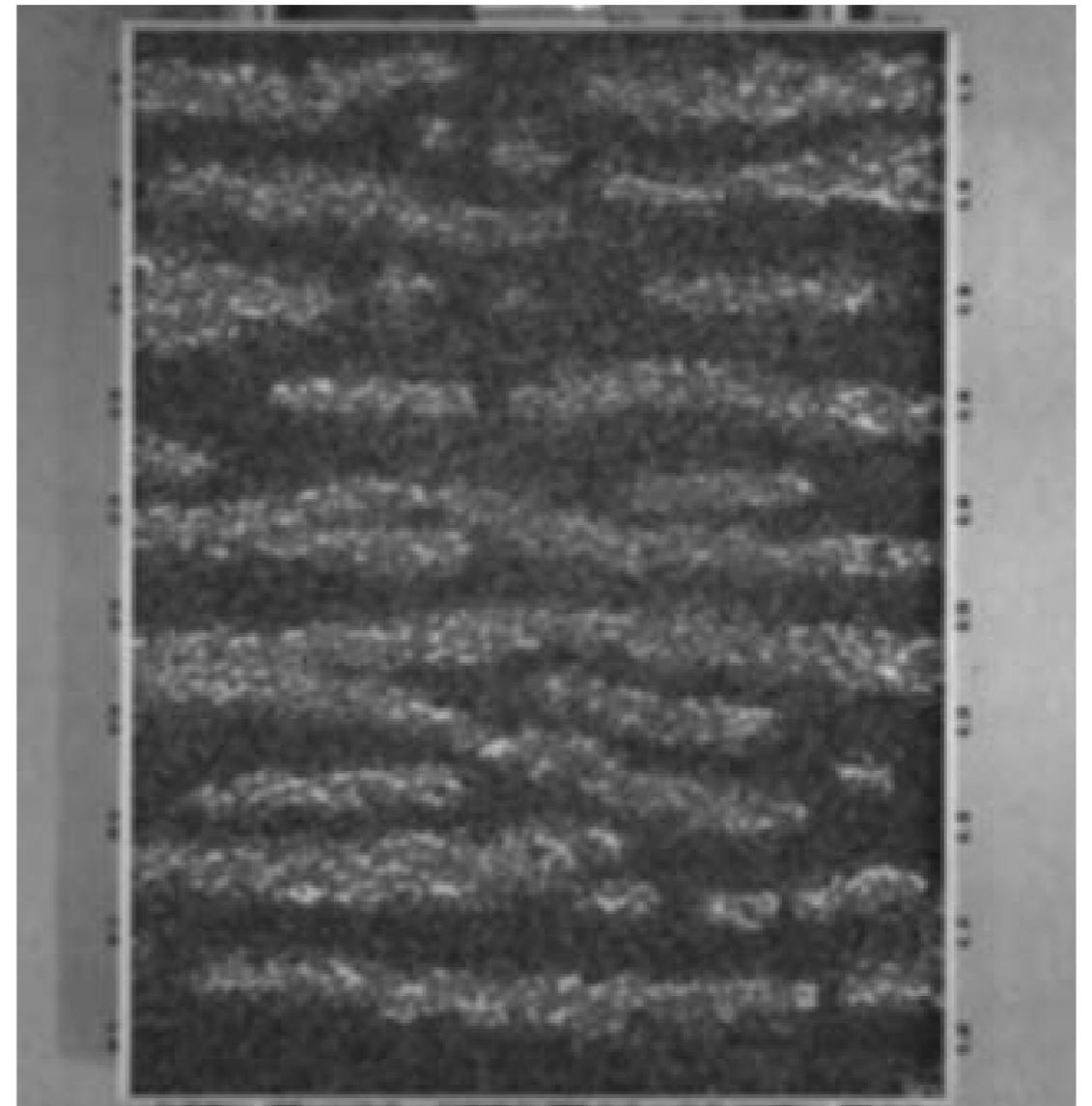
Animals



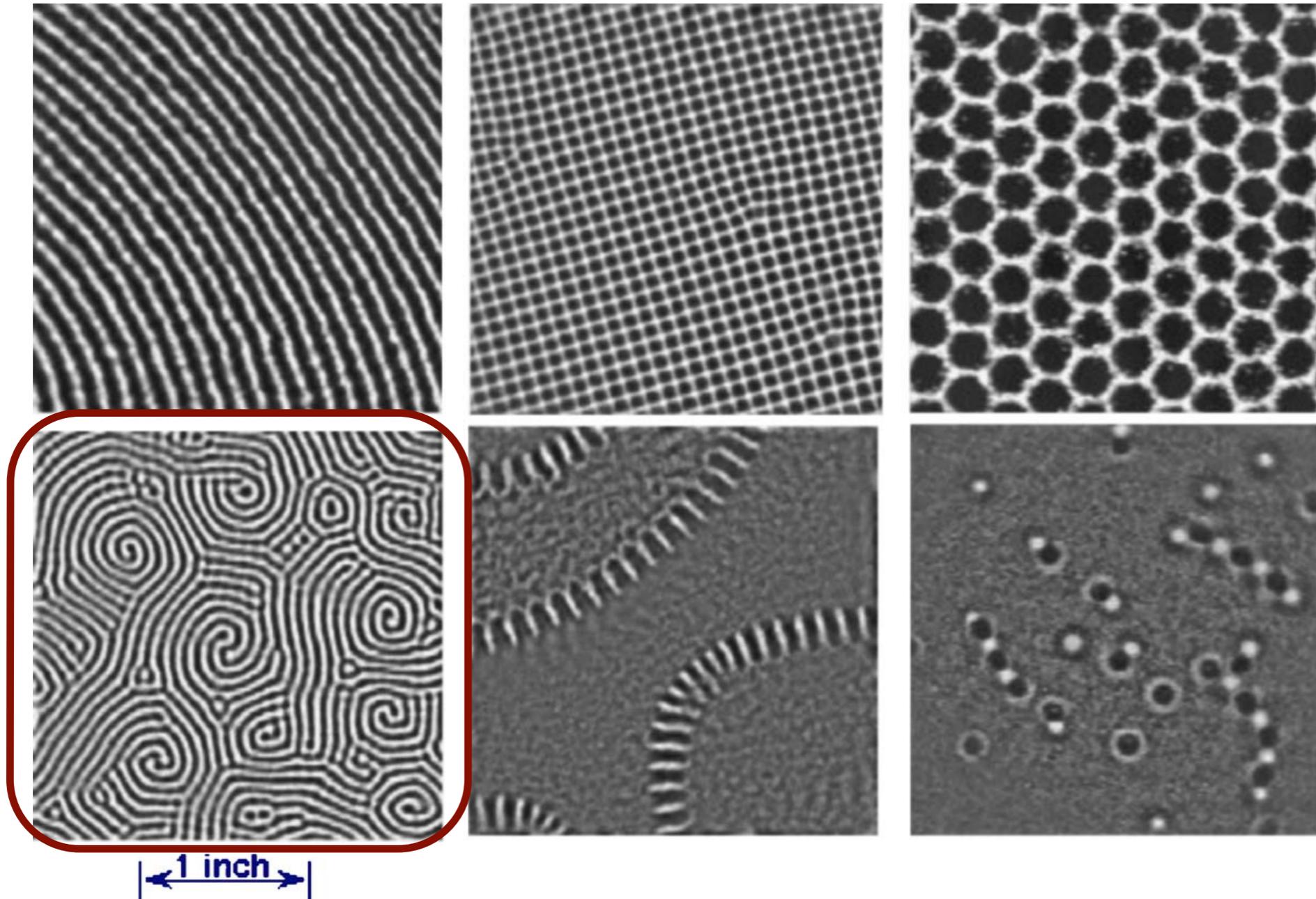
Shells



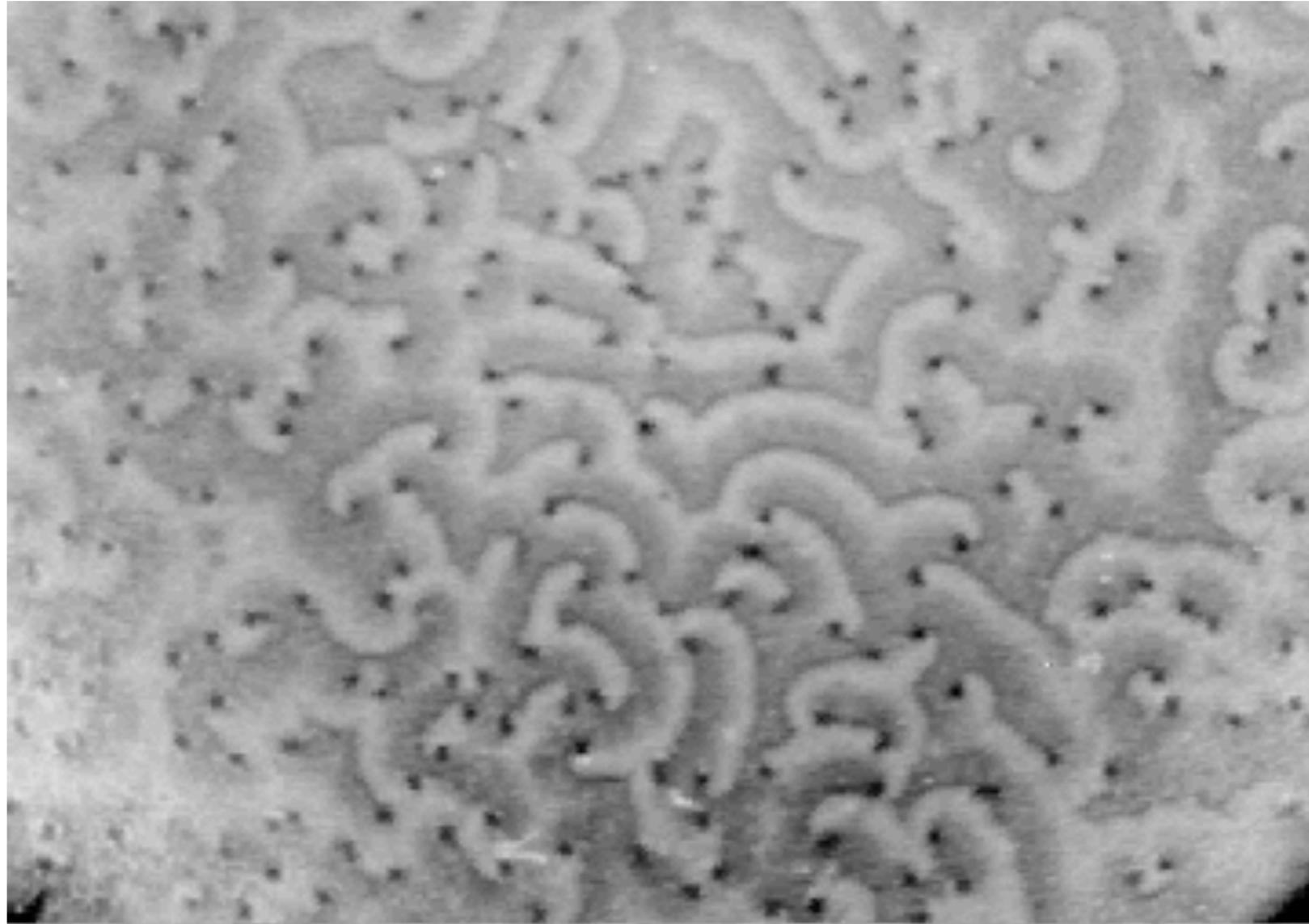
Compare: vibrated granular media



Vibrated granular media

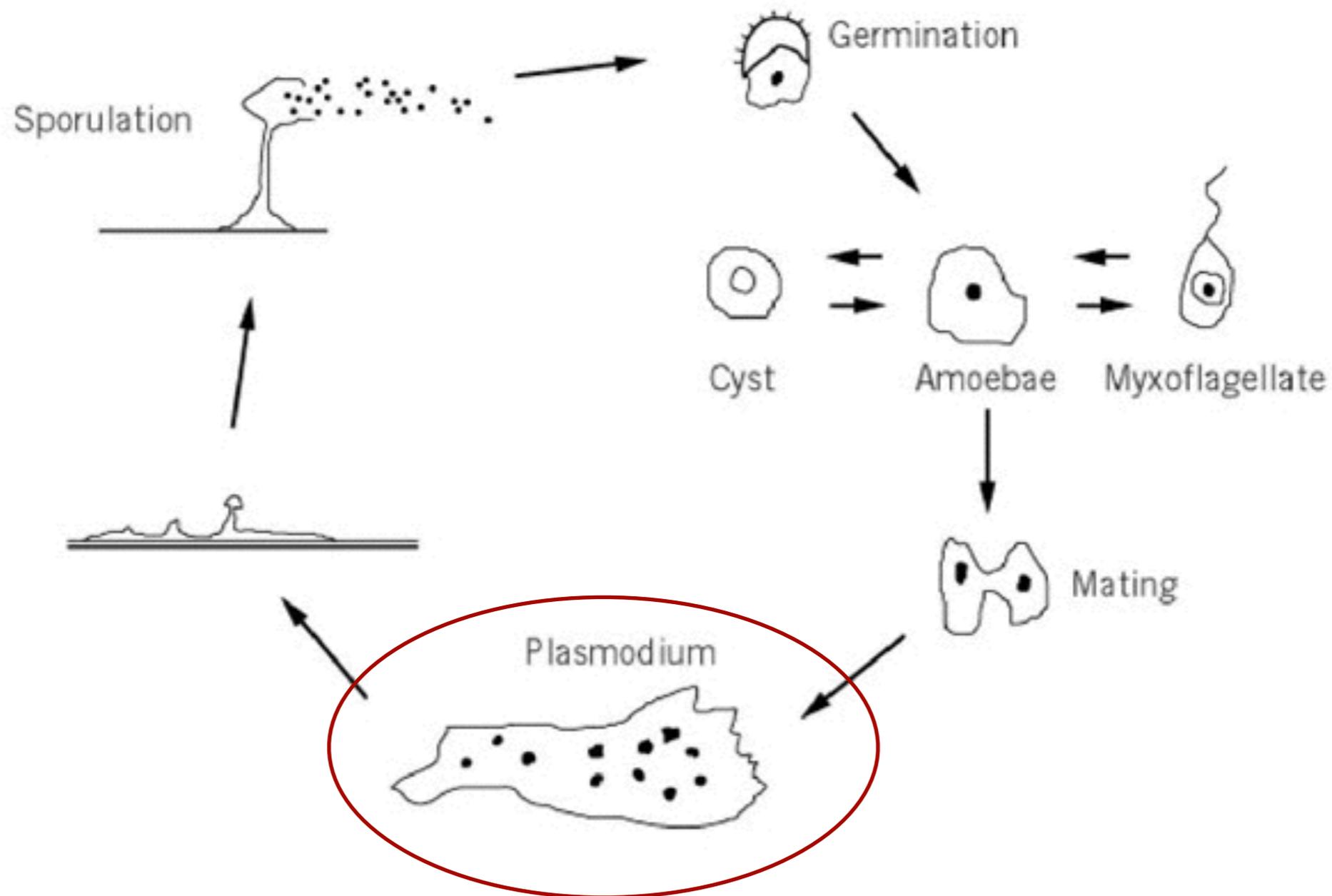


Slime mold



aggregation of a starving slime mold (credit: Florian Siegert)

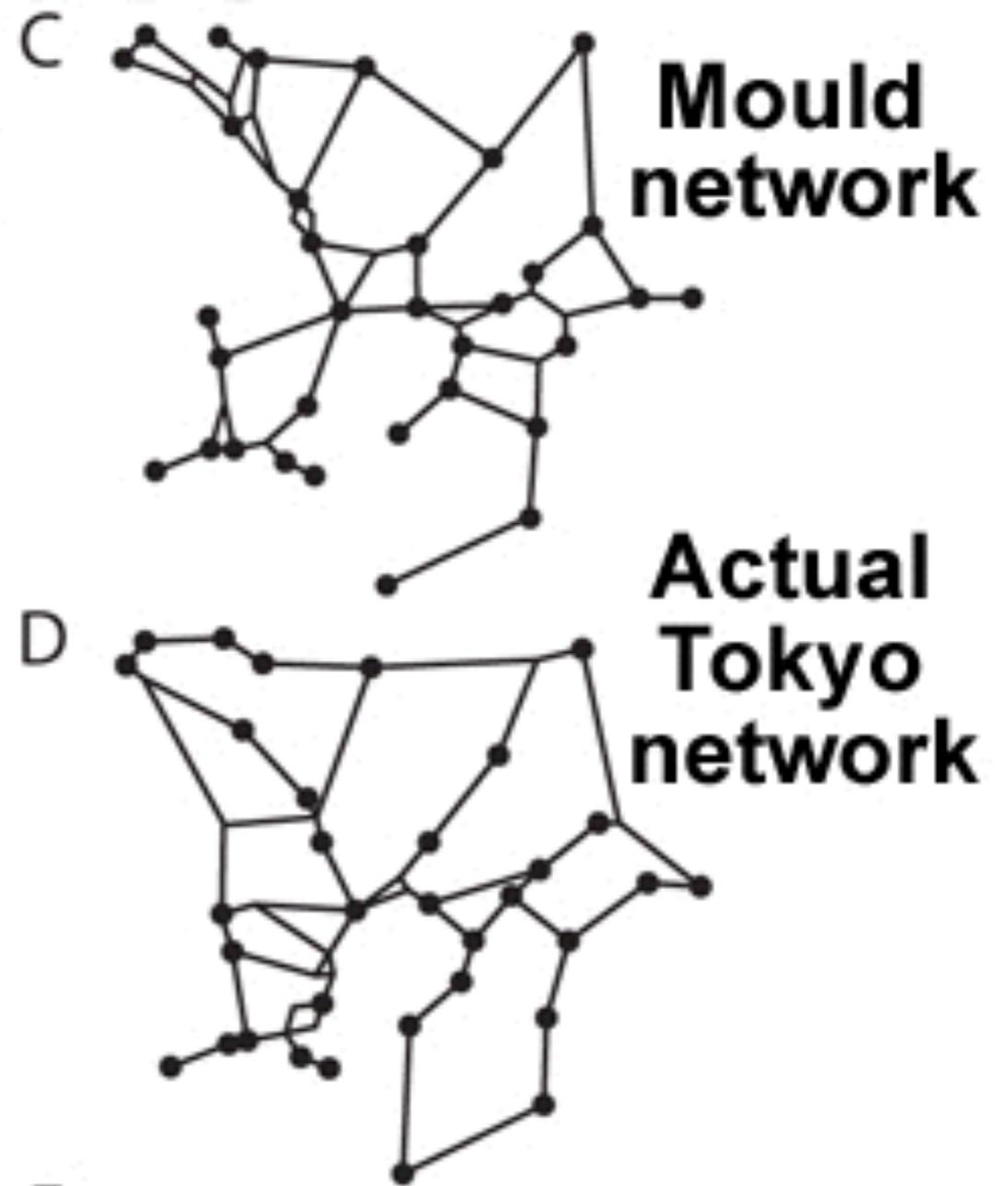
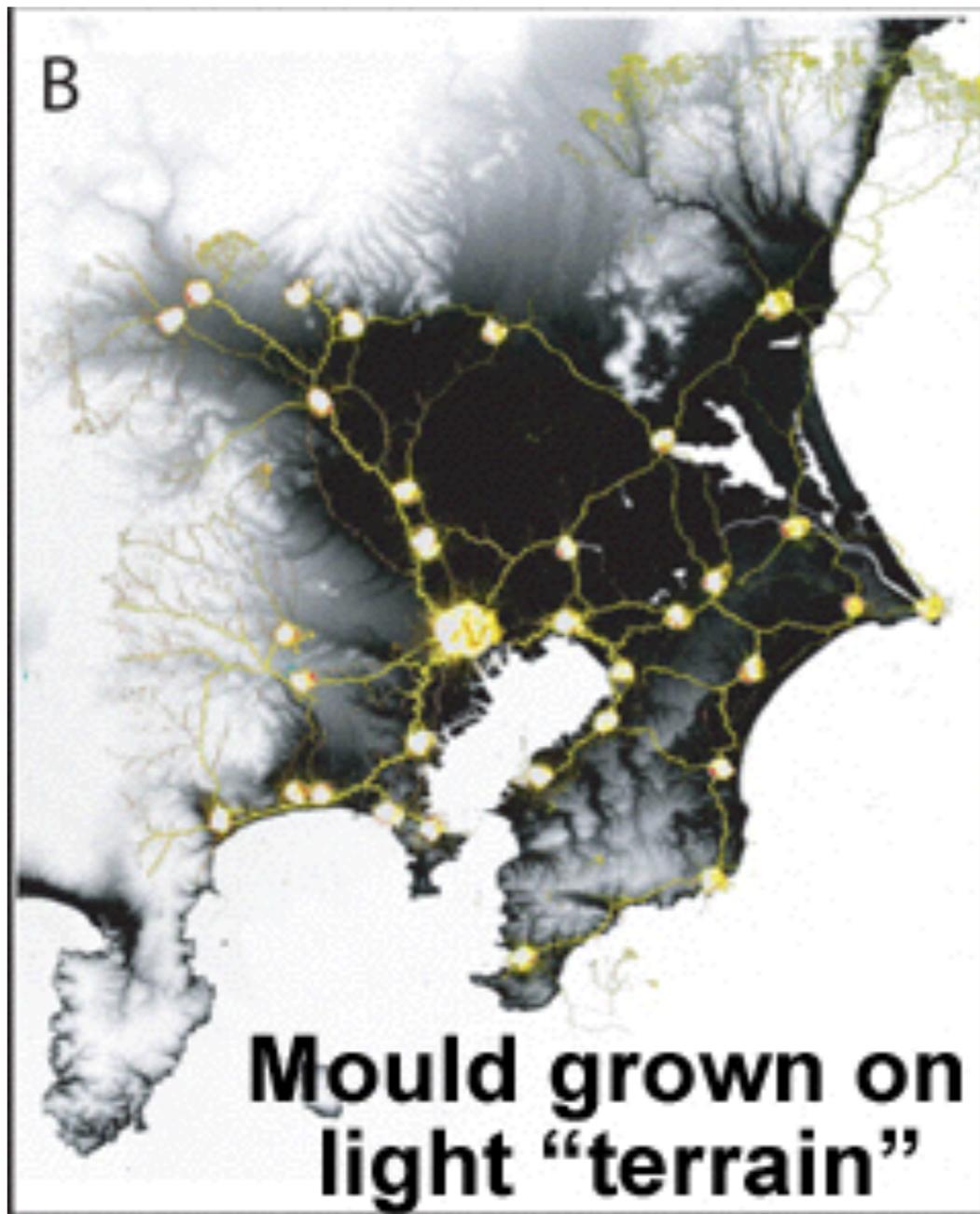
Physarum developmental cycle



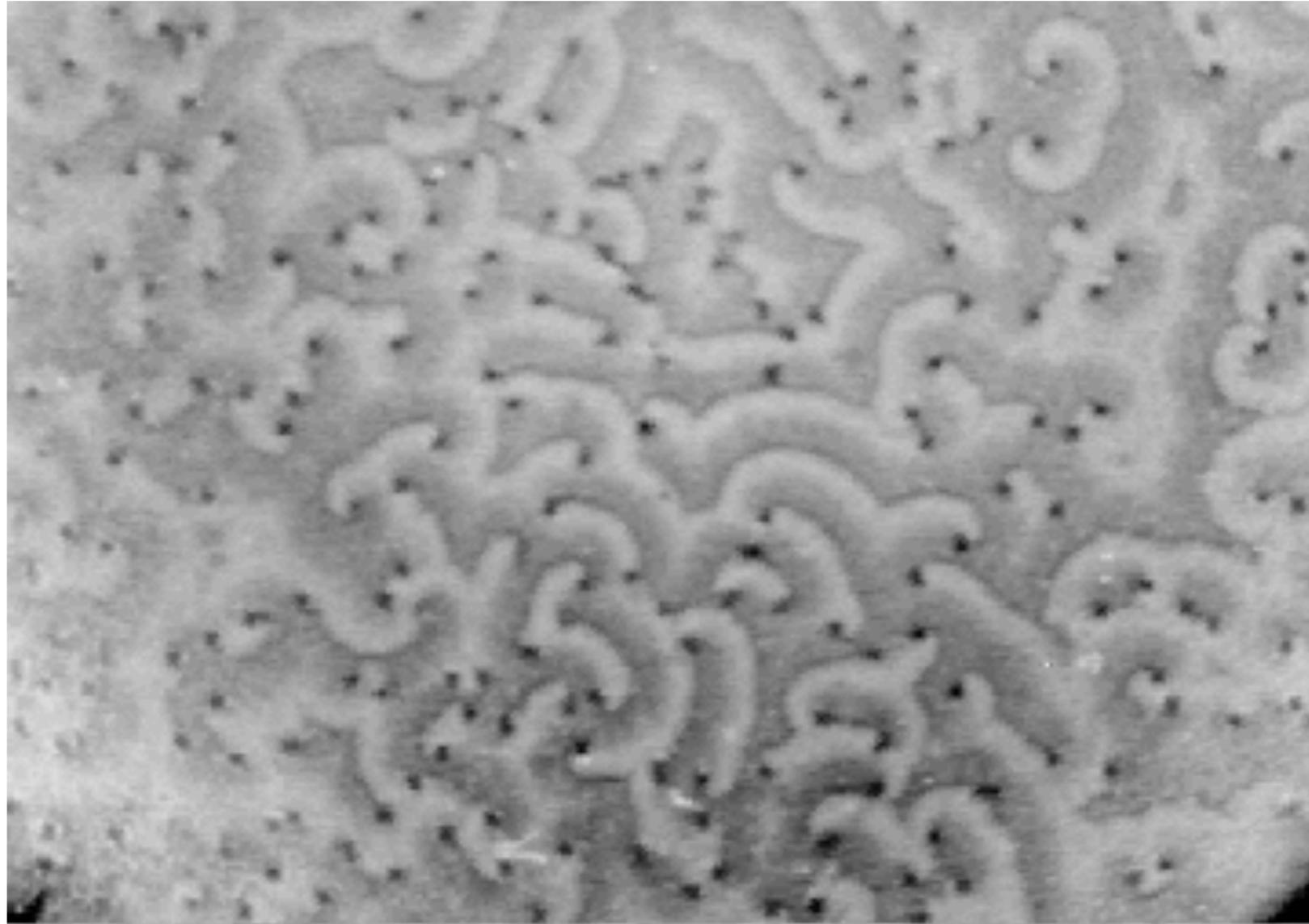
Slime mold (*Physarum plasmodium*)



Tero *et al* (2010) Science



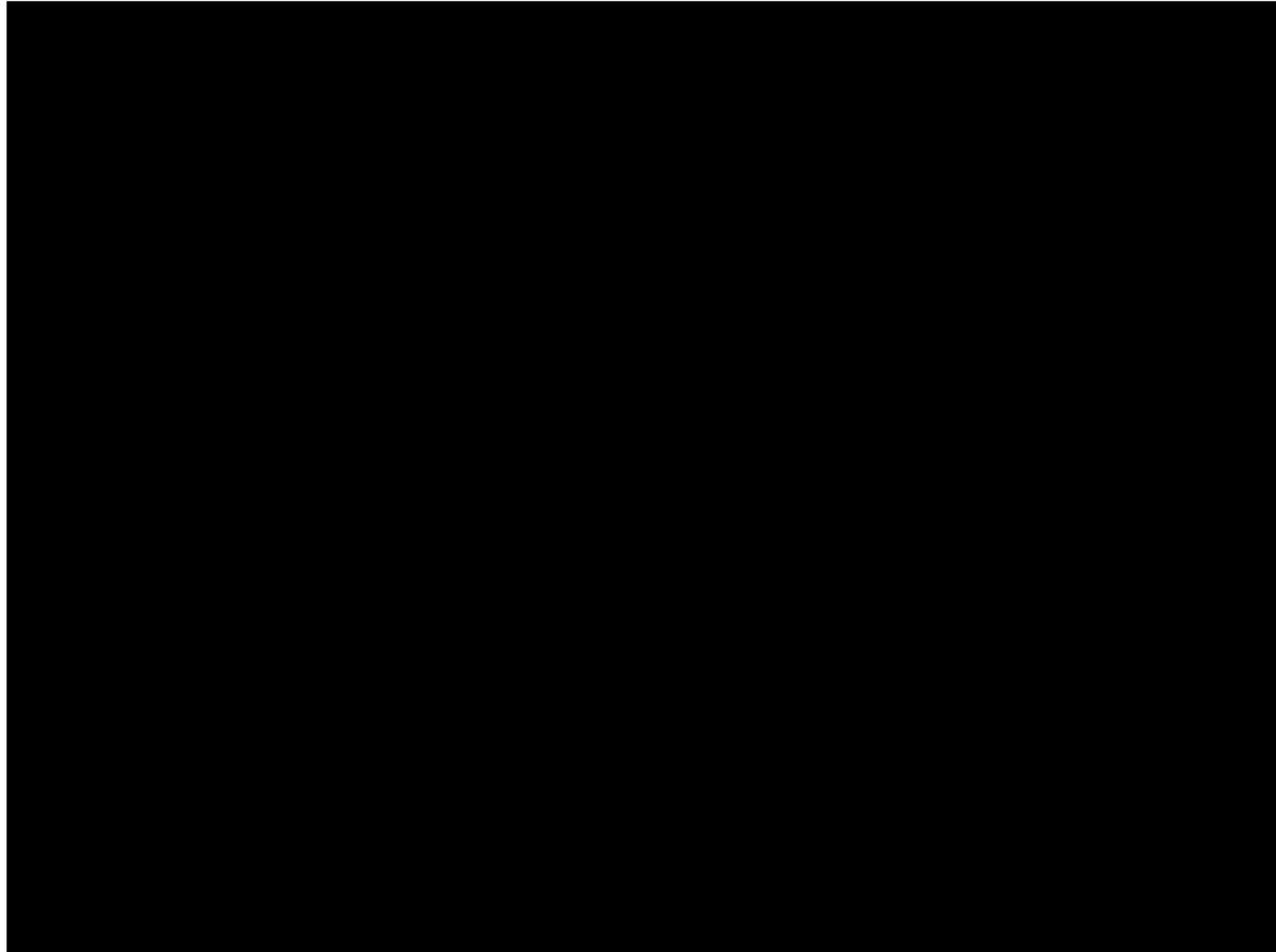
Slime mold



aggregation of a starving slime mold (credit: Florian Siegert)

Belousov-Zhabotinsky reaction

Movie credit: Stephen Morris



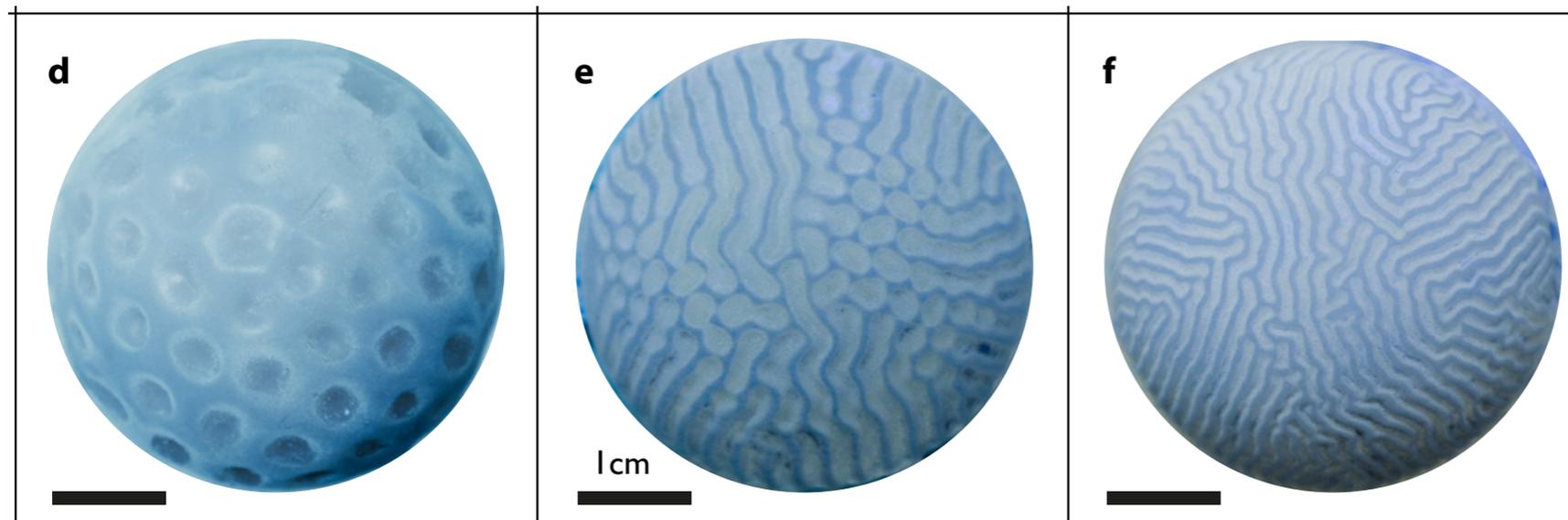
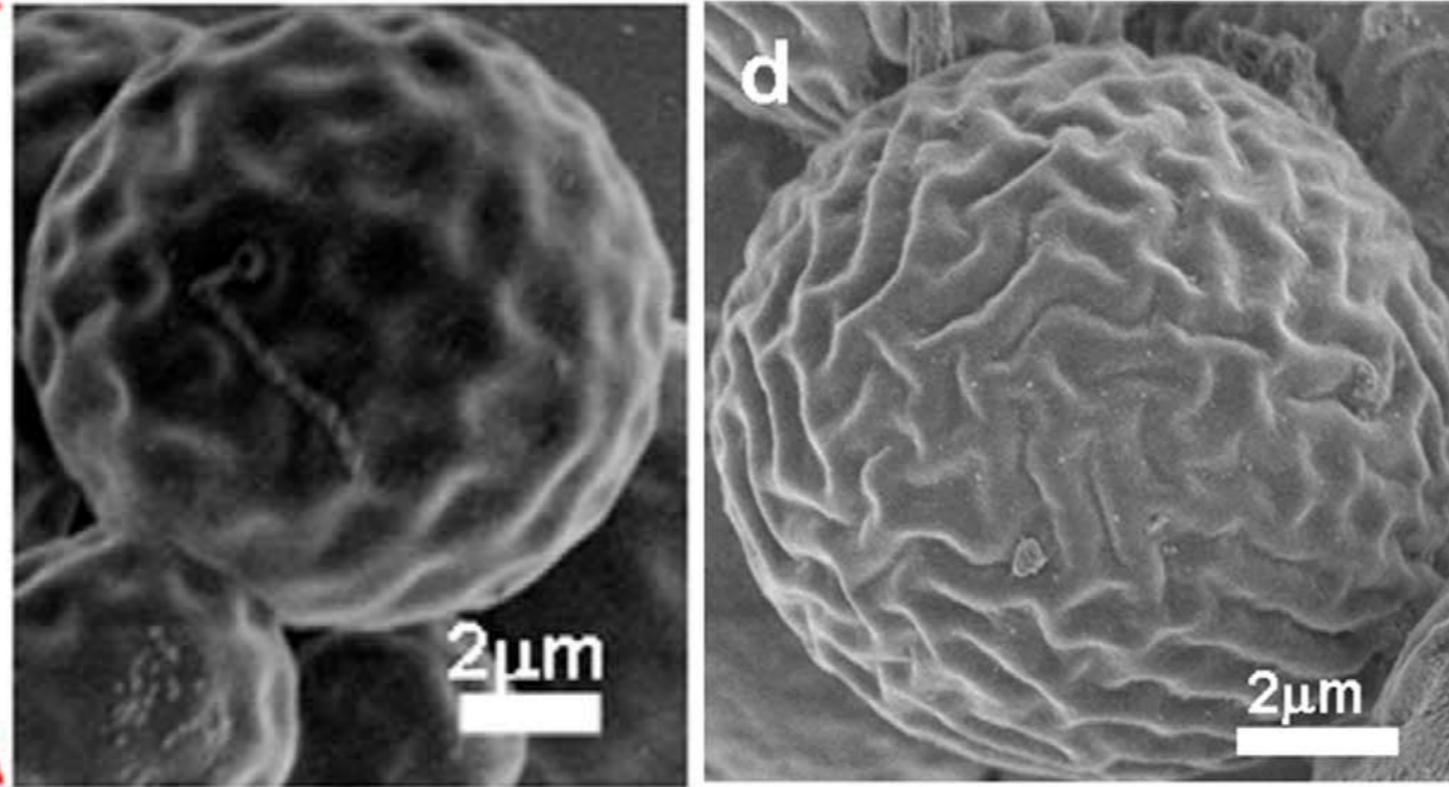
Mix of potassium bromate, cerium(IV) sulfate, malonic acid and citric acid in dilute sulfuric acid.

The ratio of concentration of the cerium(IV) and cerium(III) ions oscillates.

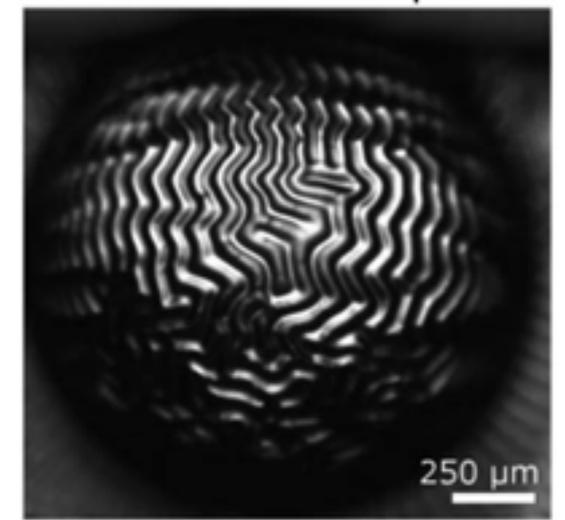
This is due to the cerium(IV) ions being reduced by malonic acid to cerium(III) ions, which are then oxidized back to cerium(IV) ions by bromate(V) ions.

Surface buckling

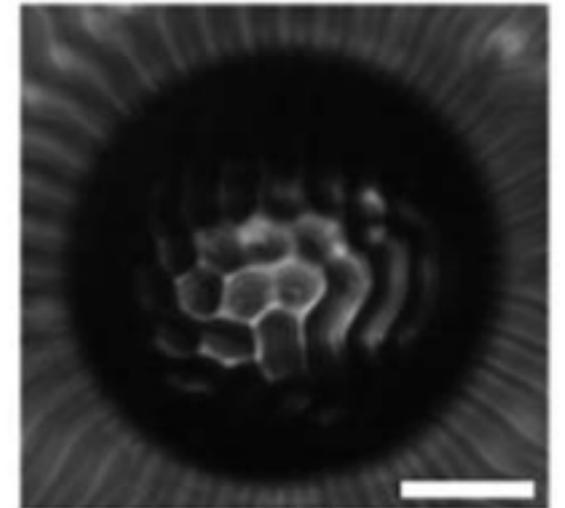
Yin et al (2014) Sci Rep



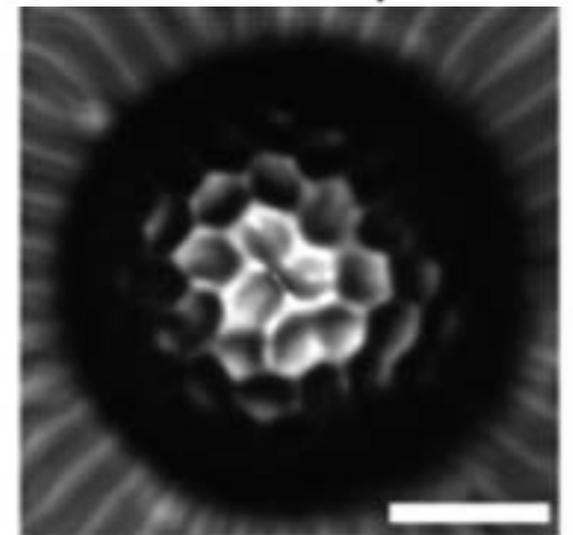
Breid & Crosby (2013) Soft Matter



$R = 805 \mu\text{m}$



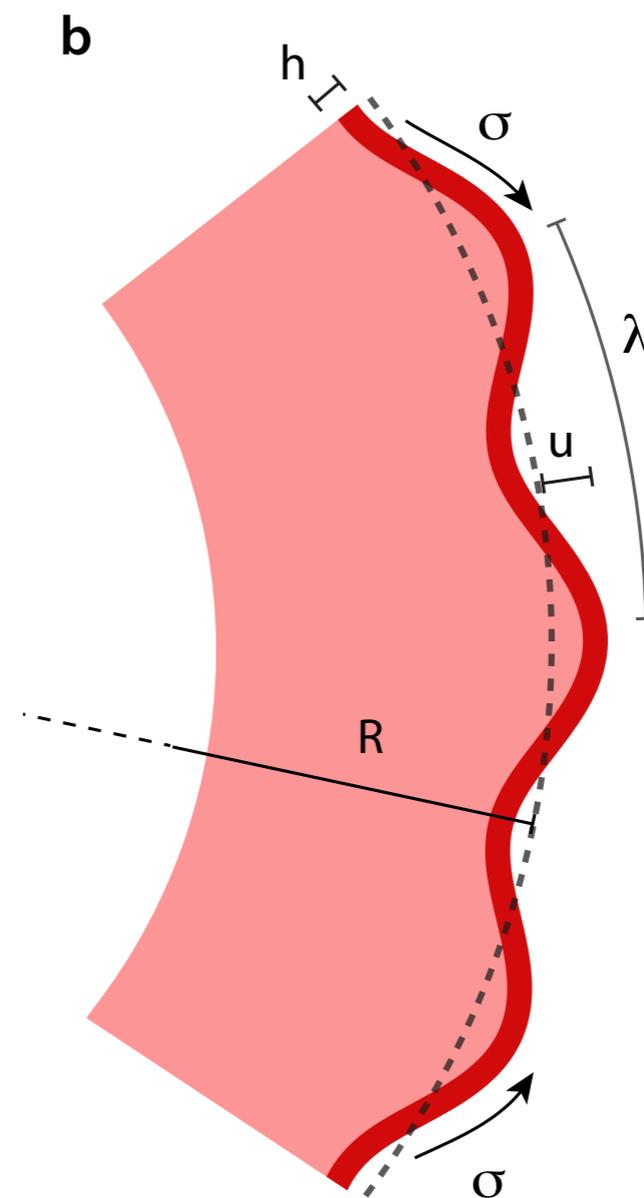
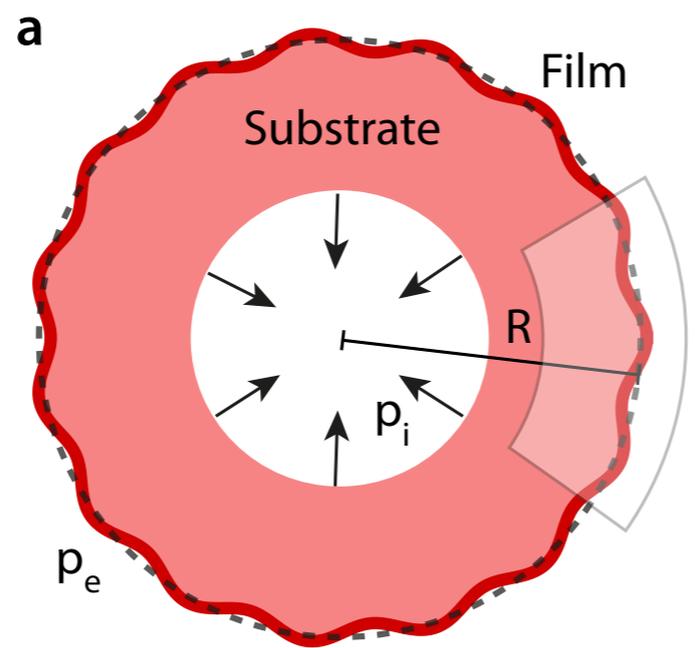
$R = 428 \mu\text{m}$



$R = 381 \mu\text{m}$

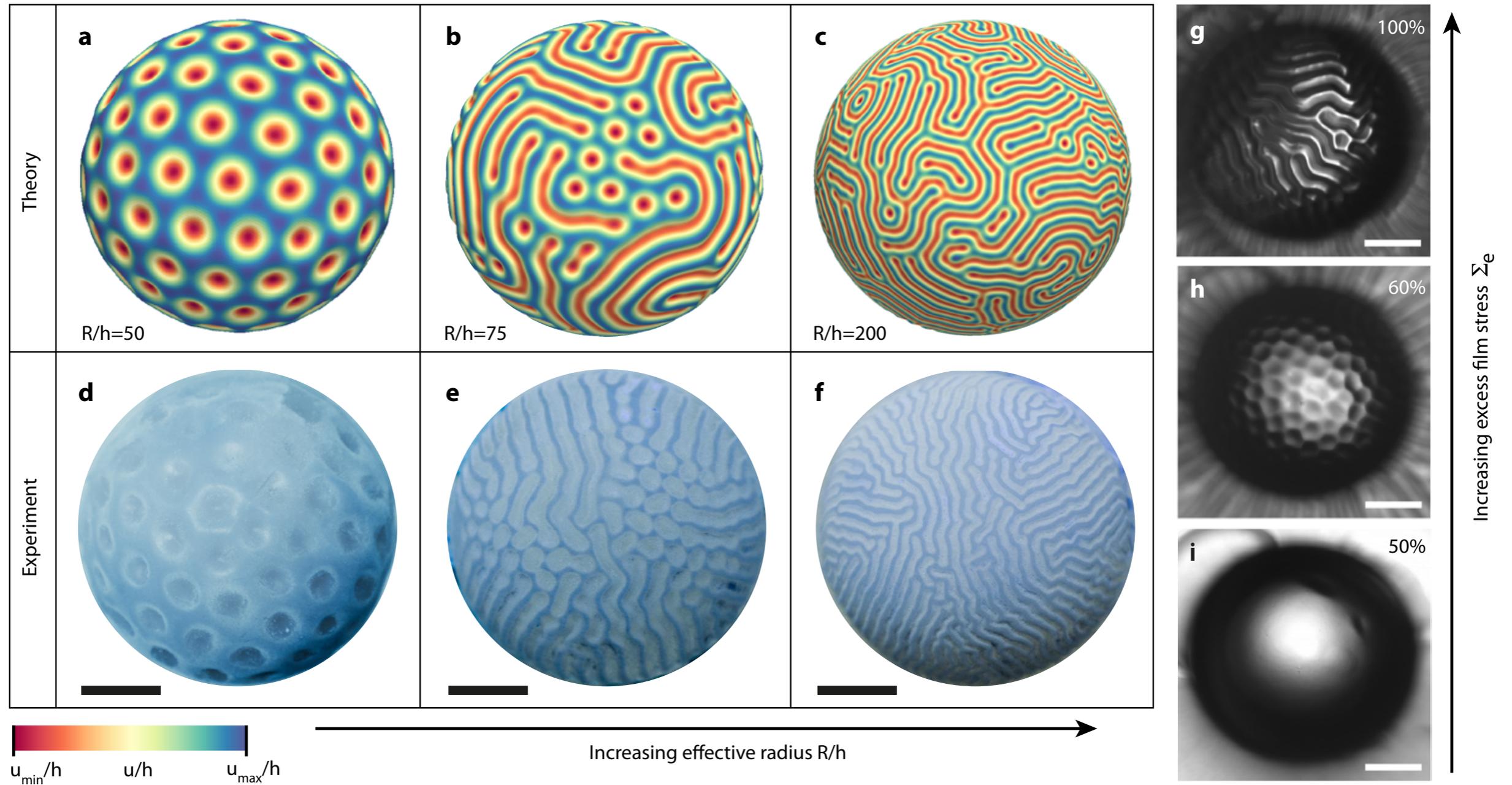
Stoop, Lagrange, Terwagne, Reis & Dunkel, 2014





Reis lab @ MIT

Theory vs. experiment



Stoop et al., Nature Materials, 2015



Higher-order PDE models & symmetry-breaking

Mathematical description

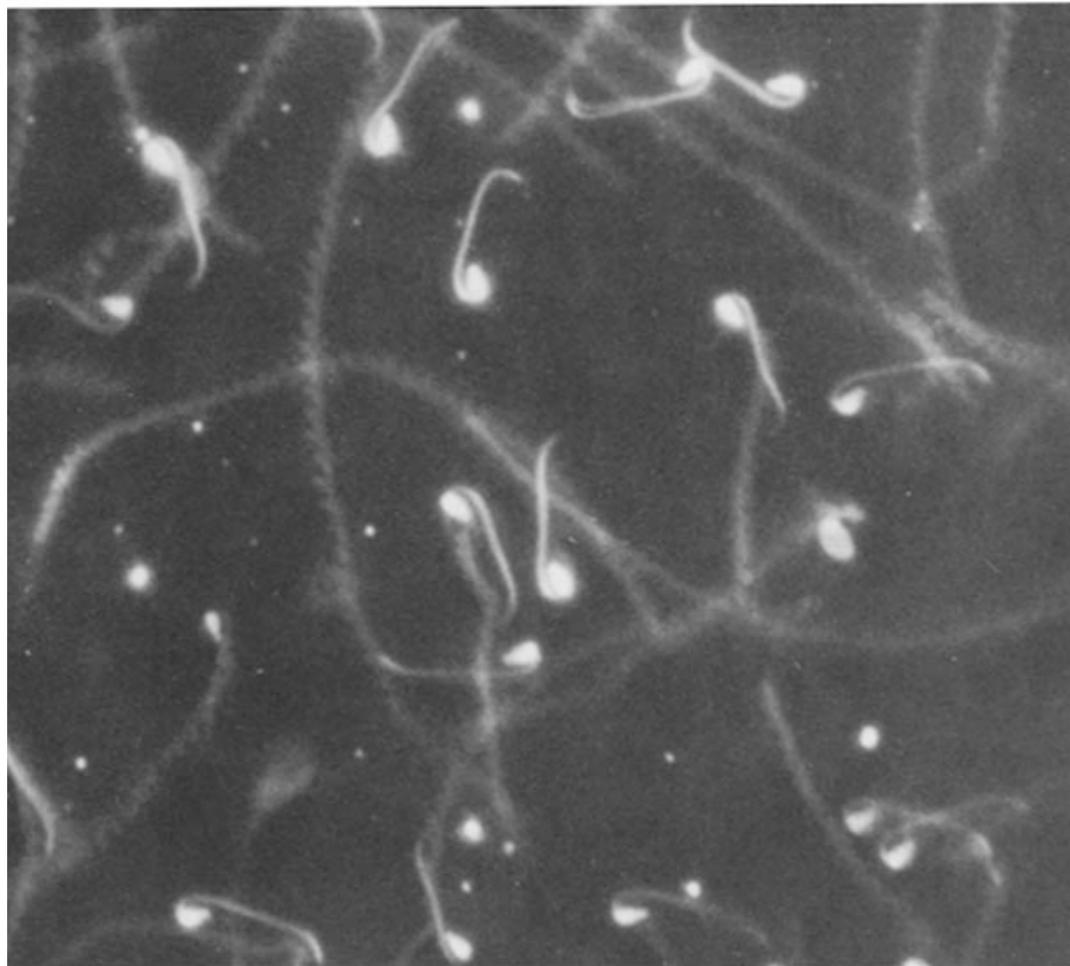
1. higher-than-second-order PDEs
2. many coupled second-order PDEs

Broken reflection-symmetry at surfaces

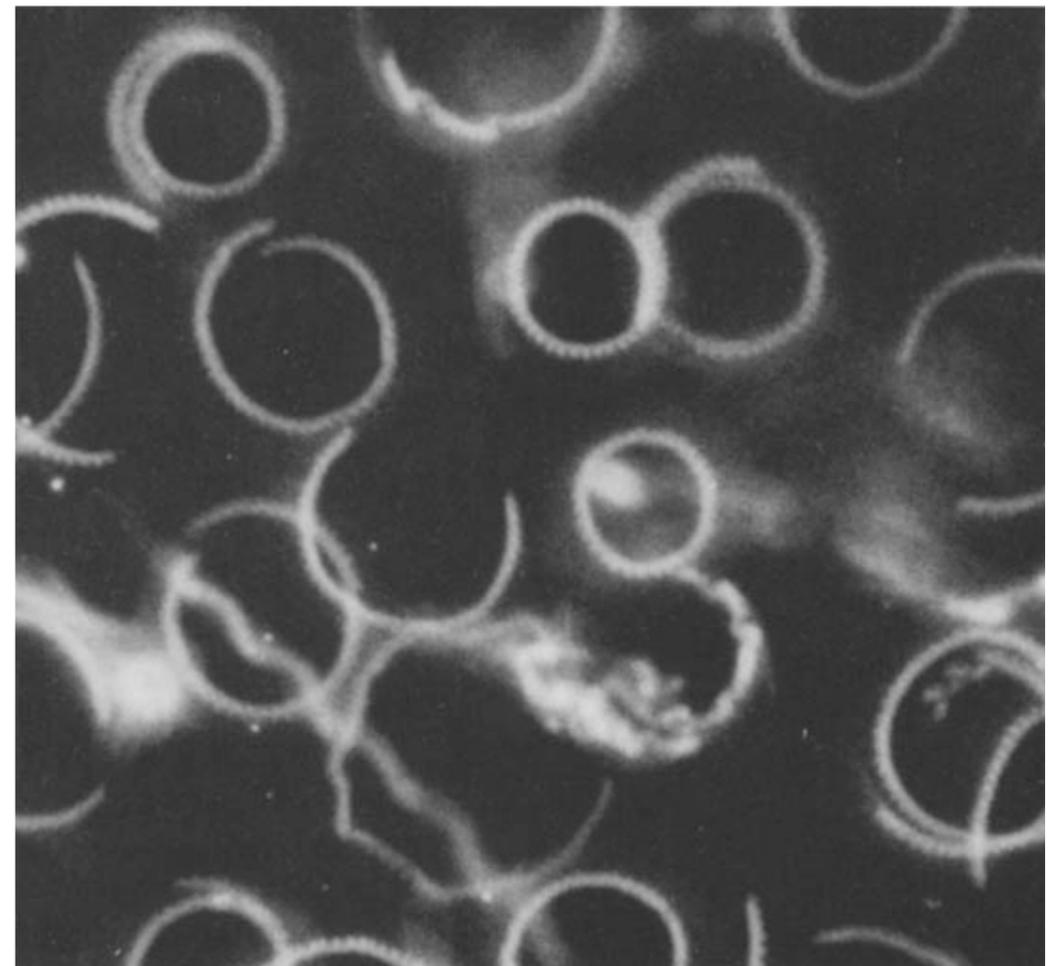
Gibbons (1980) JCB

Sea urchin sperm

in bulk (dilute)



near surface (dilute)



similar for bacteria (*E. coli*): Di Luzio et al (2005) Nature

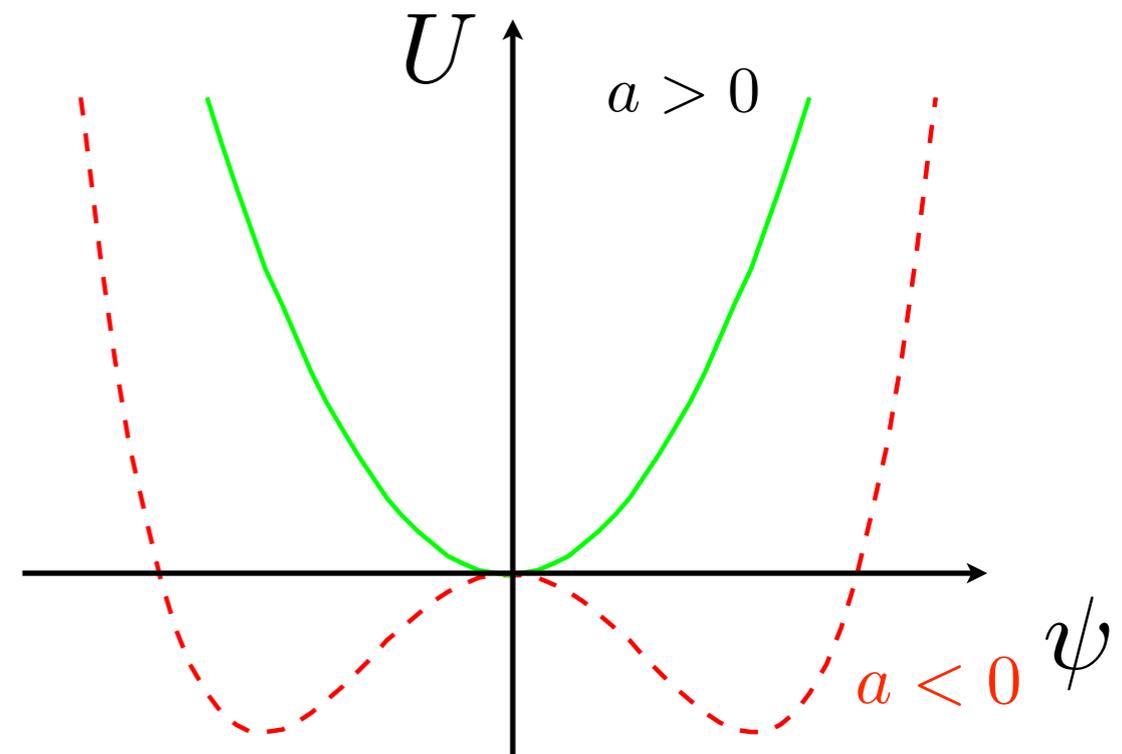
2d Swift-Hohenberg model

$$\partial_t \psi = -U'(\psi) + \gamma_0 \nabla^2 \psi - \gamma_2 (\nabla^2)^2 \psi$$

$$U(\psi) = \frac{a}{2}\psi^2 + \cancel{\frac{b}{3}\psi^3} + \frac{c}{4}\psi^4$$

reflection-symmetry

$$\psi \mapsto -\psi$$



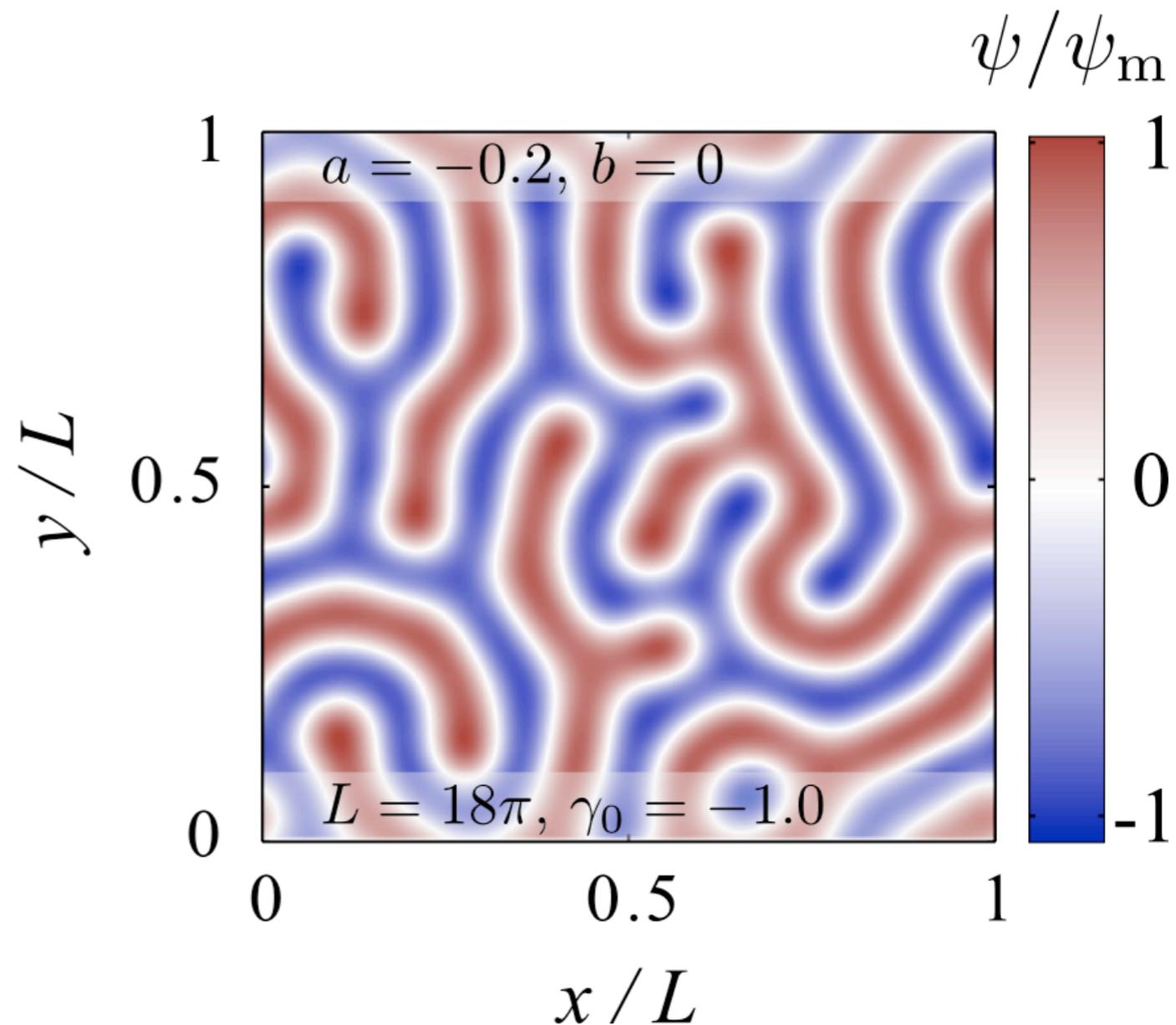
2d Swift-Hohenberg model

$$\partial_t \psi = -U'(\psi) + \gamma_0 \nabla^2 \psi - \gamma_2 (\nabla^2)^2 \psi$$

$$U(\psi) = \frac{a}{2} \psi^2 + \cancel{\frac{b}{3} \psi^3} + \frac{c}{4} \psi^4$$

reflection-symmetry

$$\psi \mapsto -\psi$$



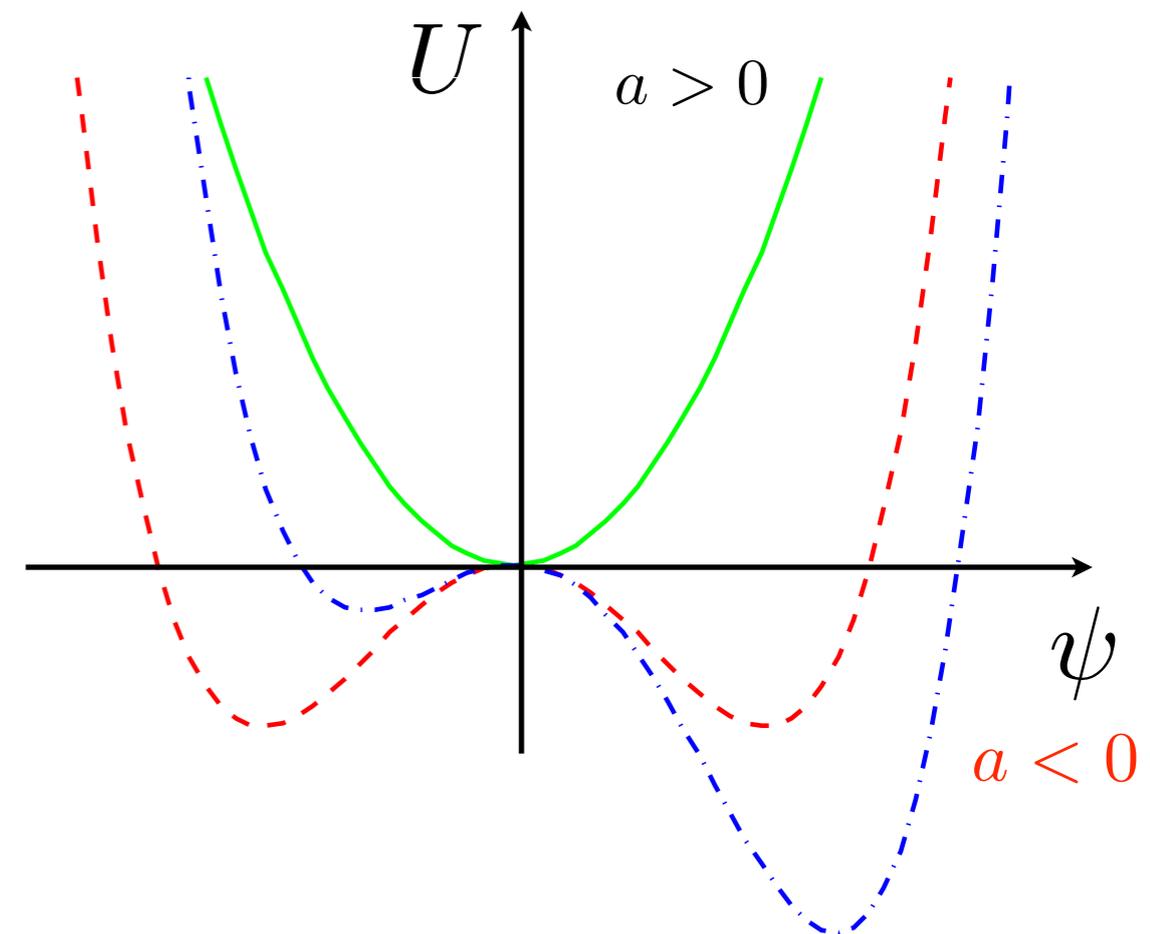
2d Swift-Hohenberg model

$$\partial_t \psi = -U'(\psi) + \gamma_0 \nabla^2 \psi - \gamma_2 (\nabla^2)^2 \psi$$

$$U(\psi) = \frac{a}{2} \psi^2 + \frac{b}{3} \psi^3 + \frac{c}{4} \psi^4$$

broken
reflection-symmetry

$$\psi \not\mapsto -\psi$$



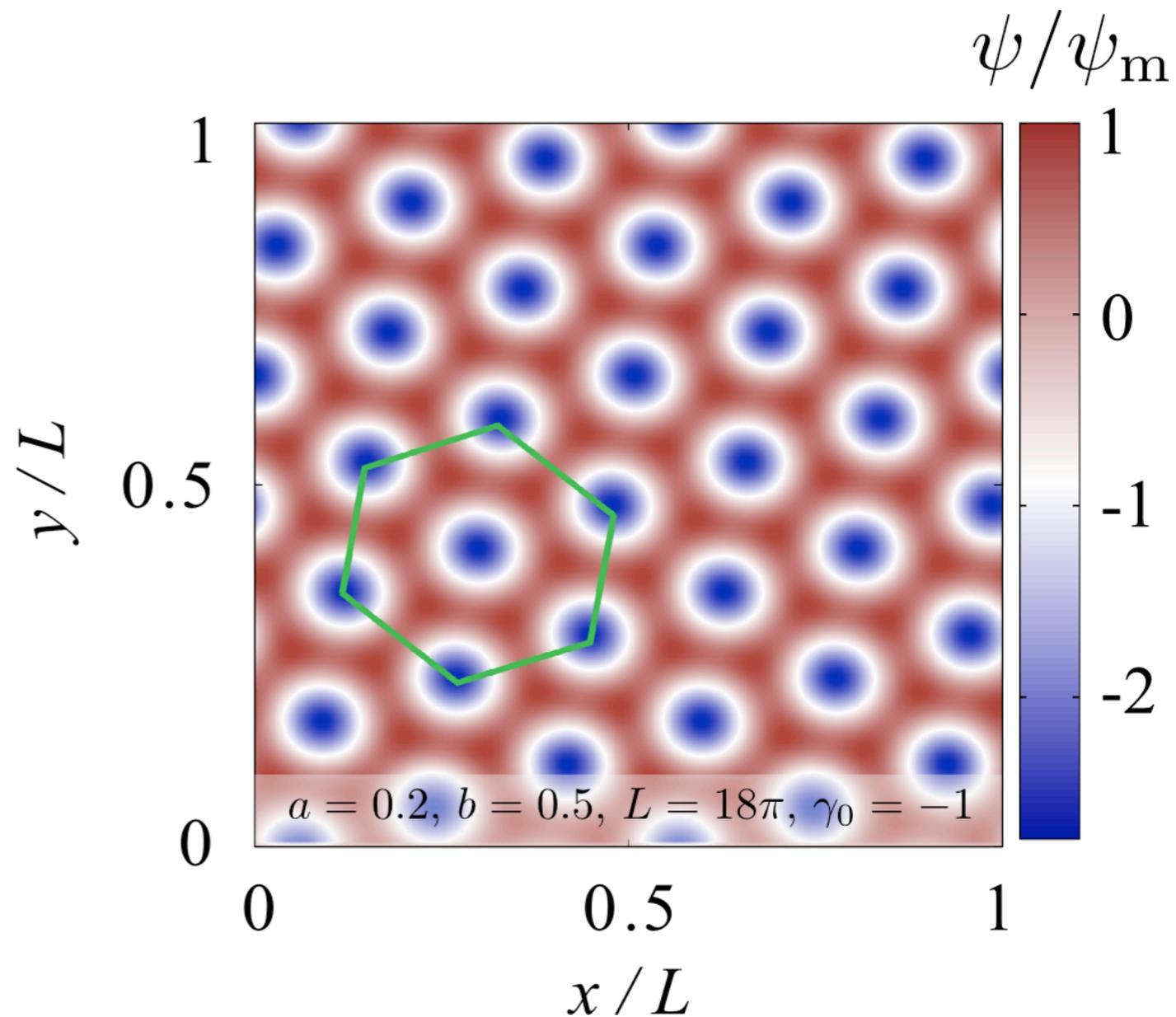
2d Swift-Hohenberg model

$$\partial_t \psi = -U'(\psi) + \gamma_0 \nabla^2 \psi - \gamma_2 (\nabla^2)^2 \psi$$

$$U(\psi) = \frac{a}{2} \psi^2 + \frac{b}{3} \psi^3 + \frac{c}{4} \psi^4$$

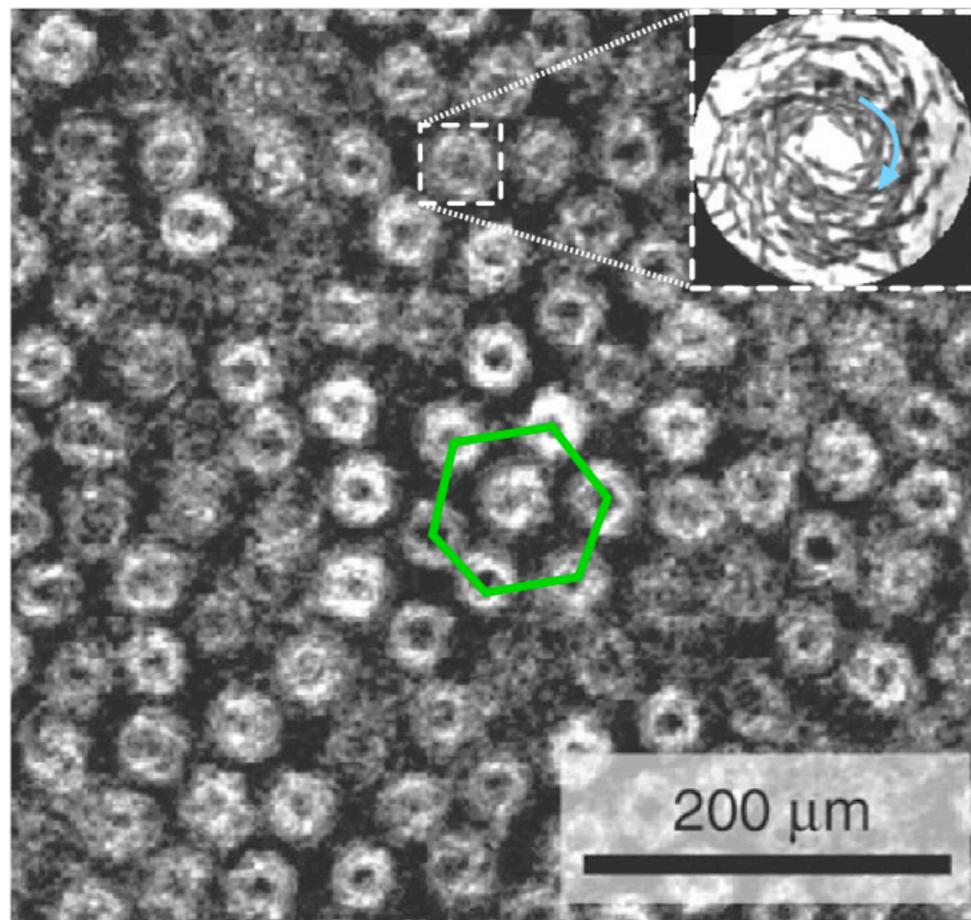
broken
reflection-symmetry

$$\psi \not\mapsto -\psi$$



2d Swift-Hohenberg model

Sea urchin sperm cells
near surface
(high concentration)



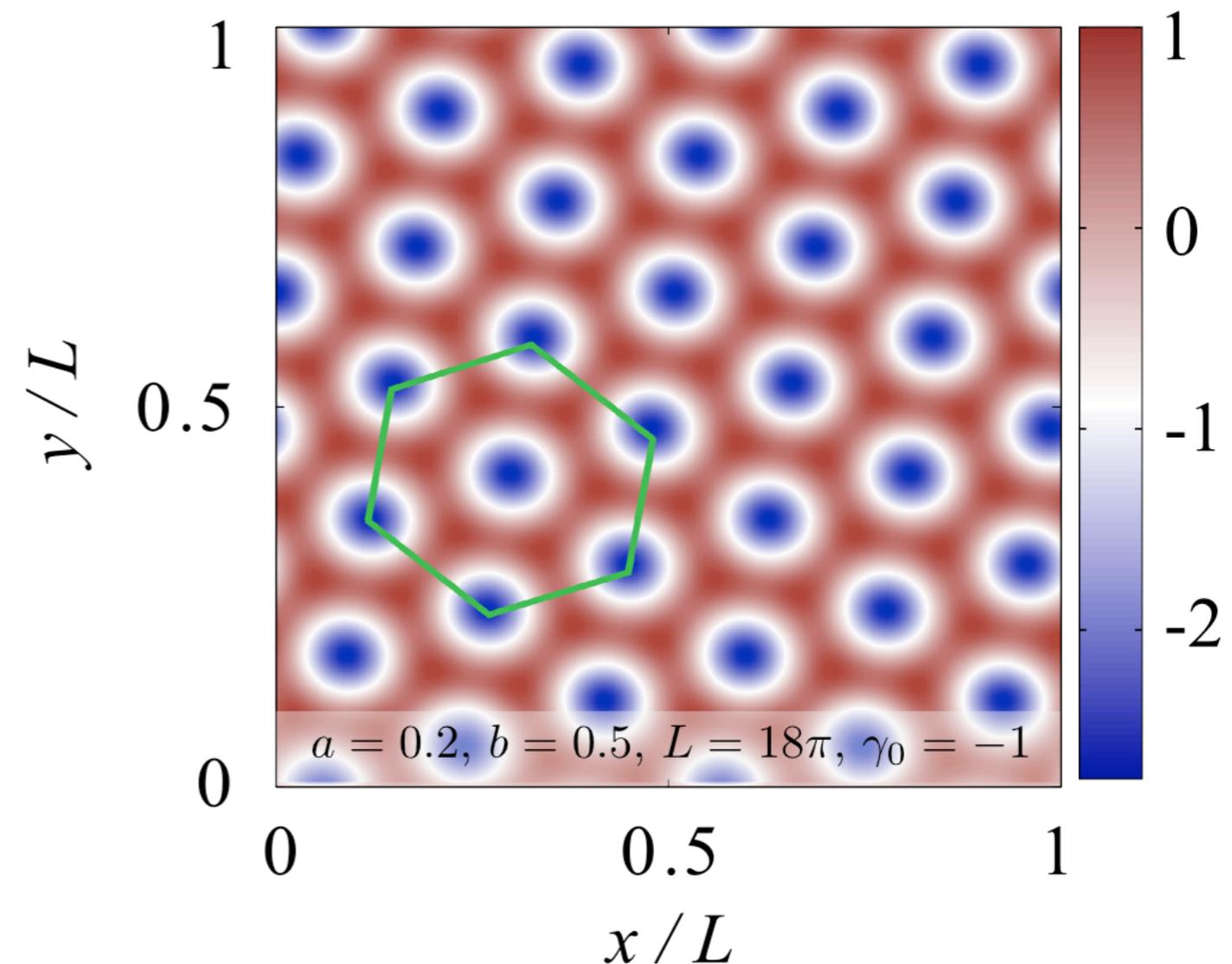
Riedel et al (2007) Science

broken
reflection-symmetry

$$b \neq 0$$

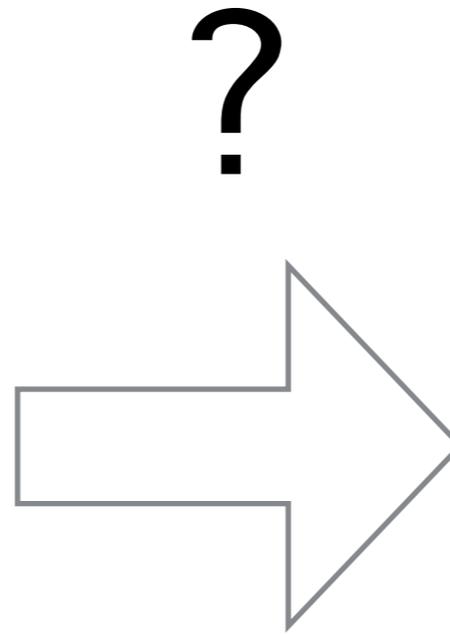
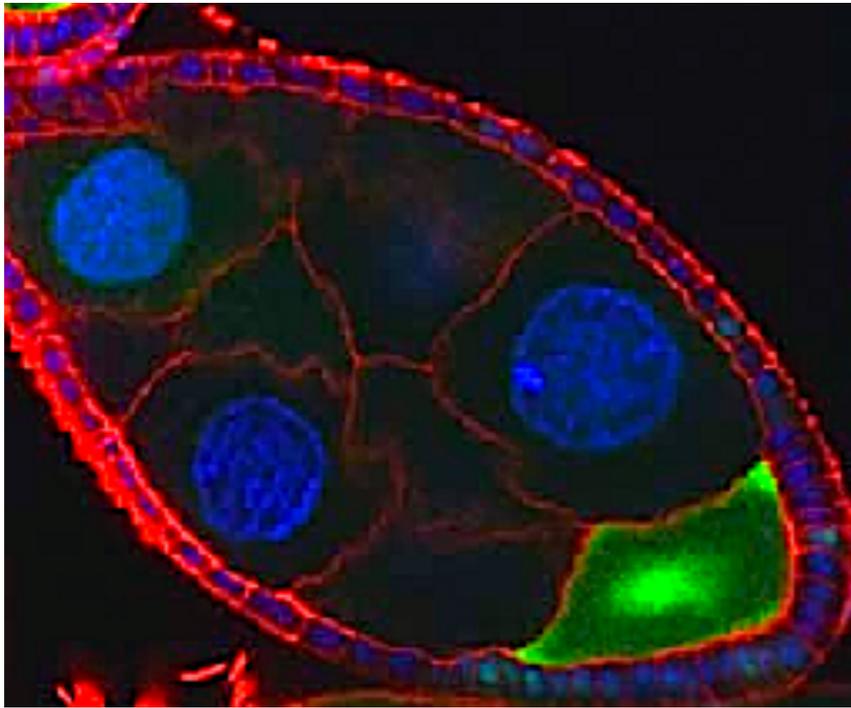
$$\psi \not\leftrightarrow -\psi$$

$$\psi / \psi_m$$



Mathematical description

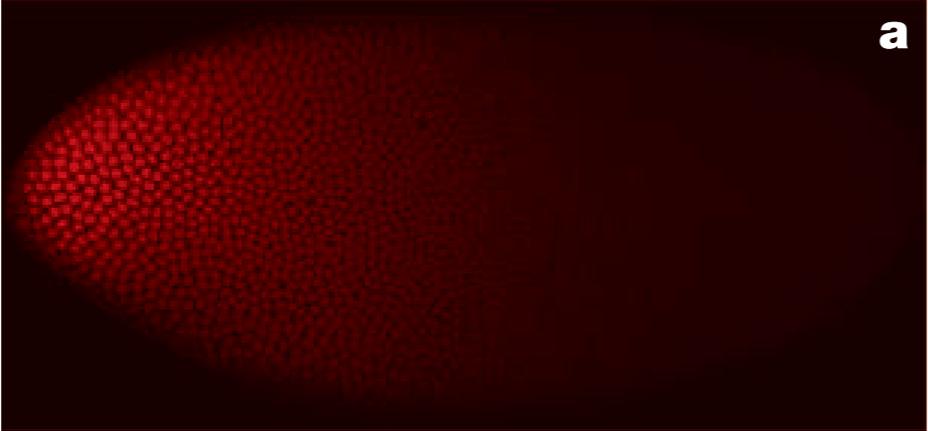
1. higher-than-second-order PDEs
2. many coupled second-order PDEs



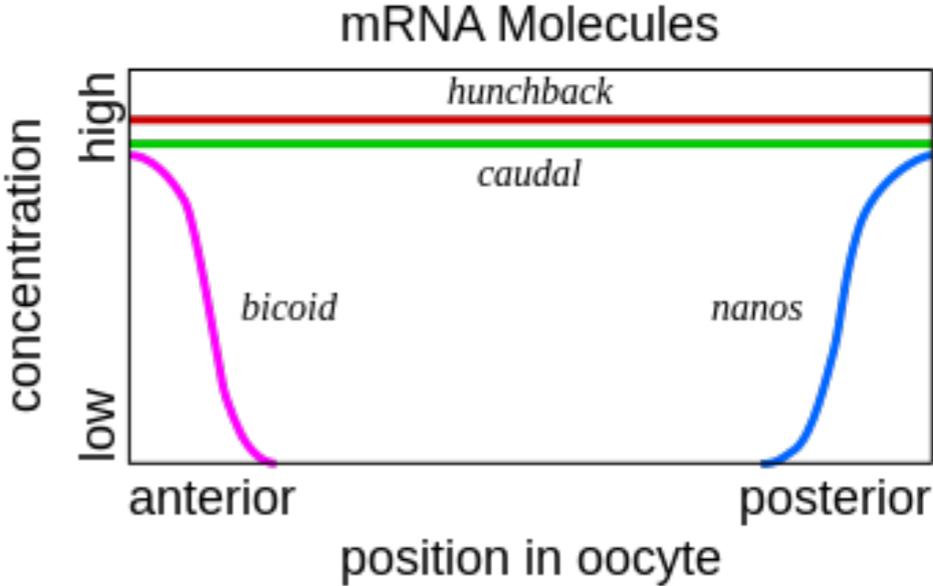
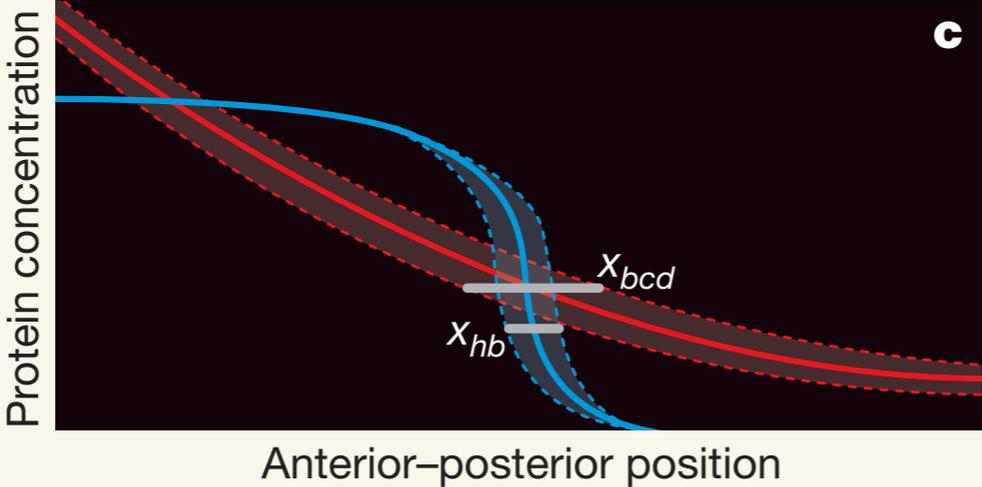
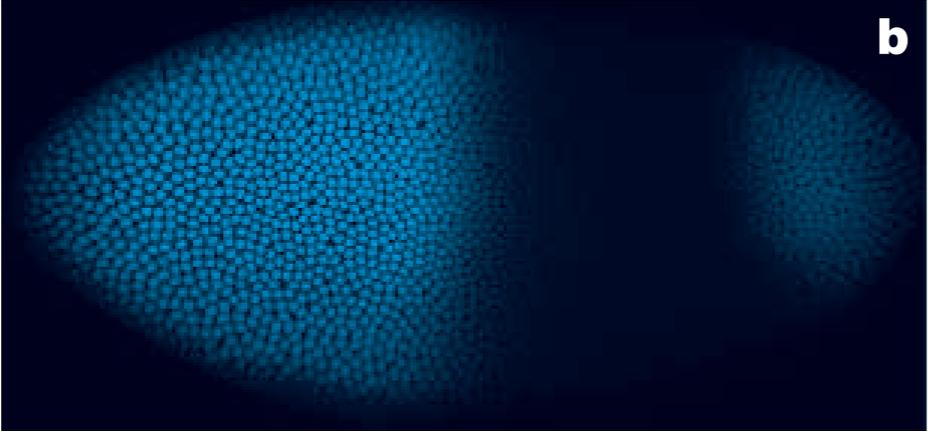
Maternal input

From: Reinitz

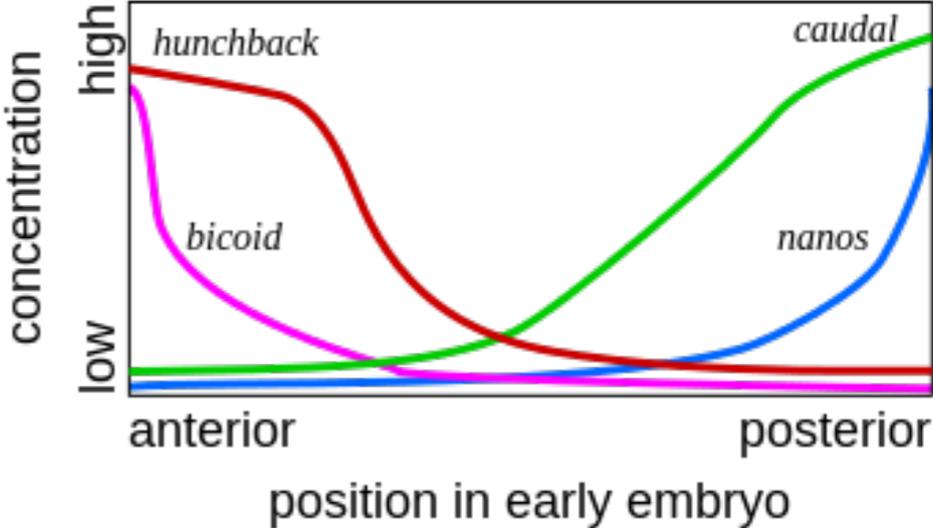
Bicoid



Hunchback

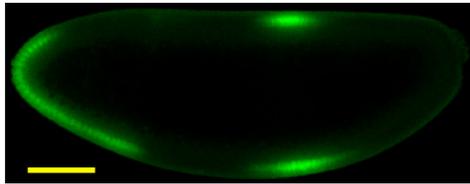


Proteins

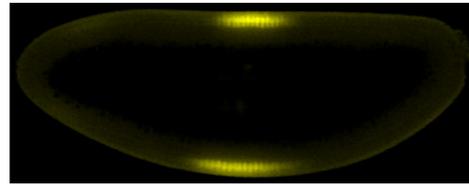


Intensity profiles

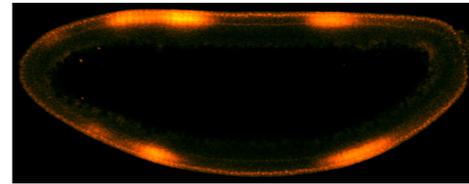
Kni (488nm)



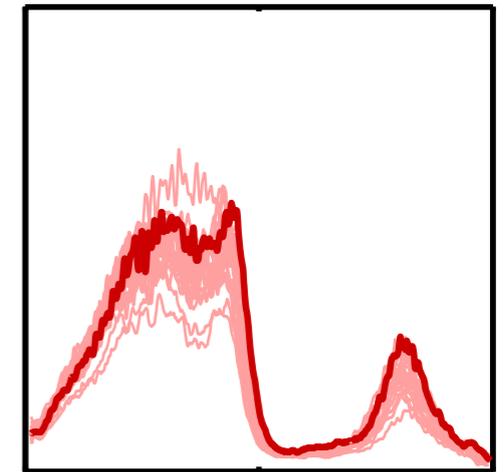
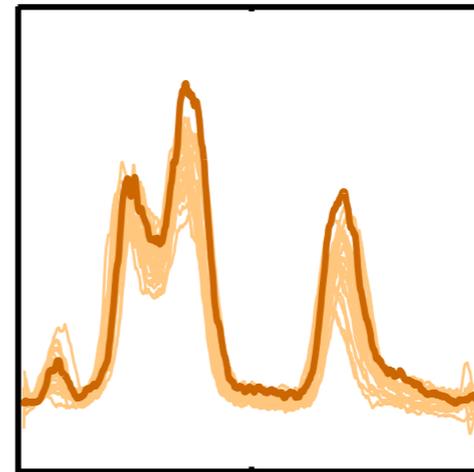
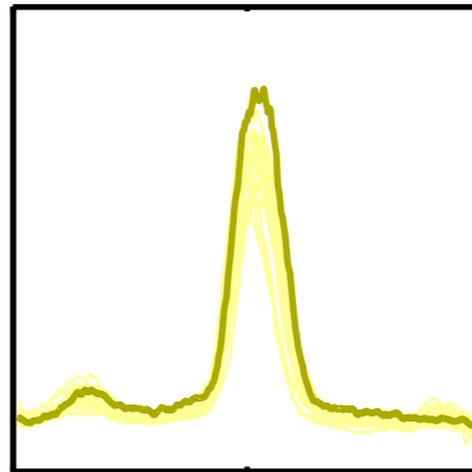
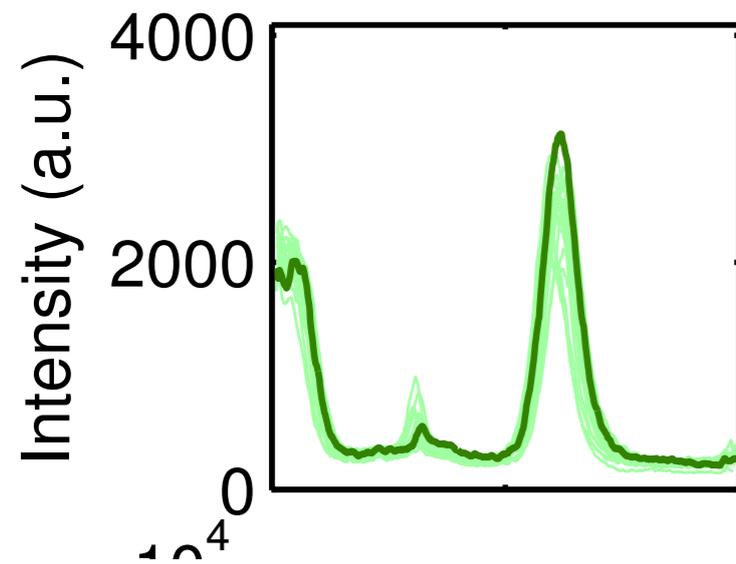
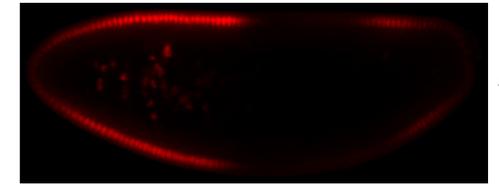
Kr (546nm)

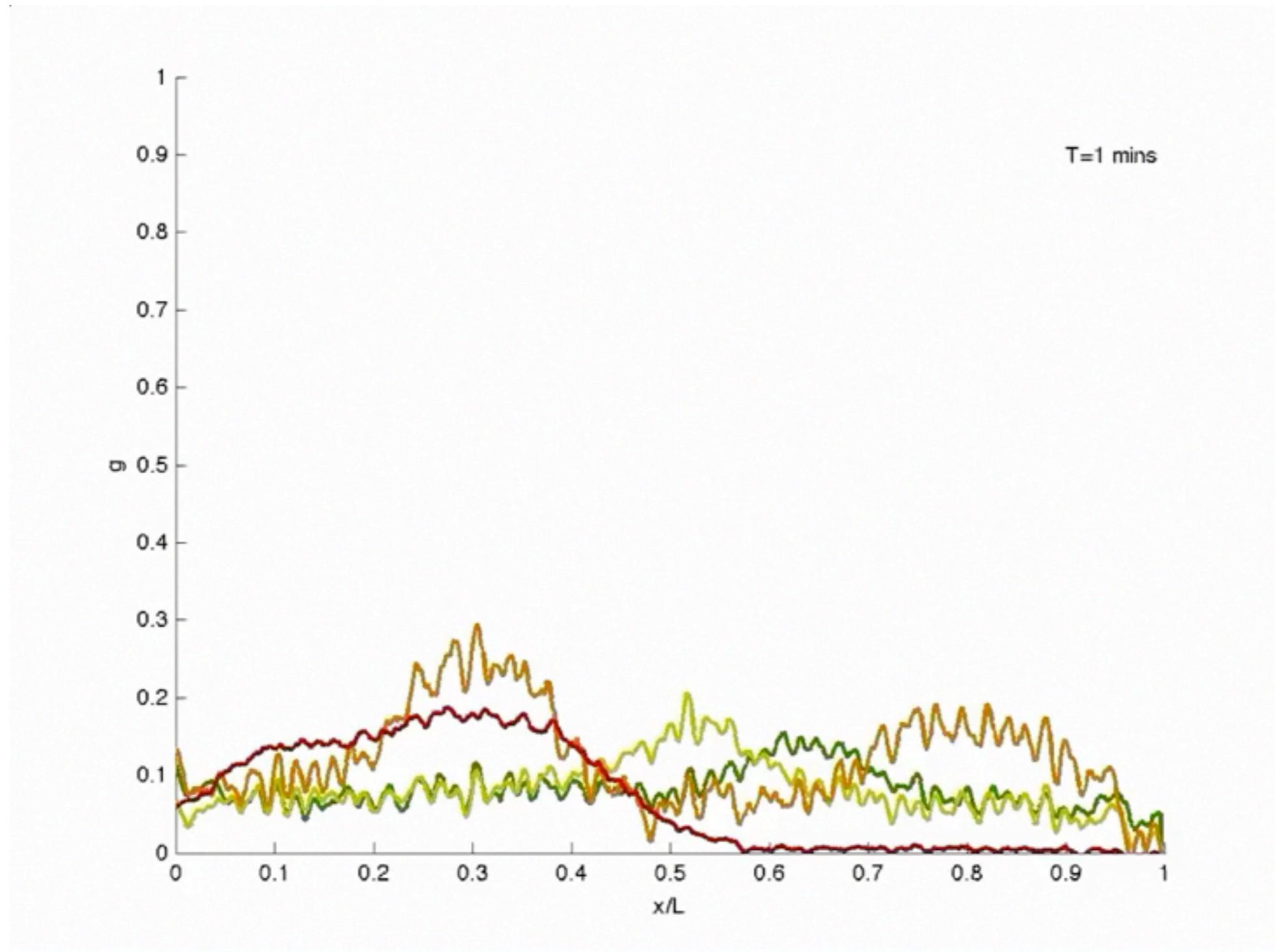


Gt (594nm)

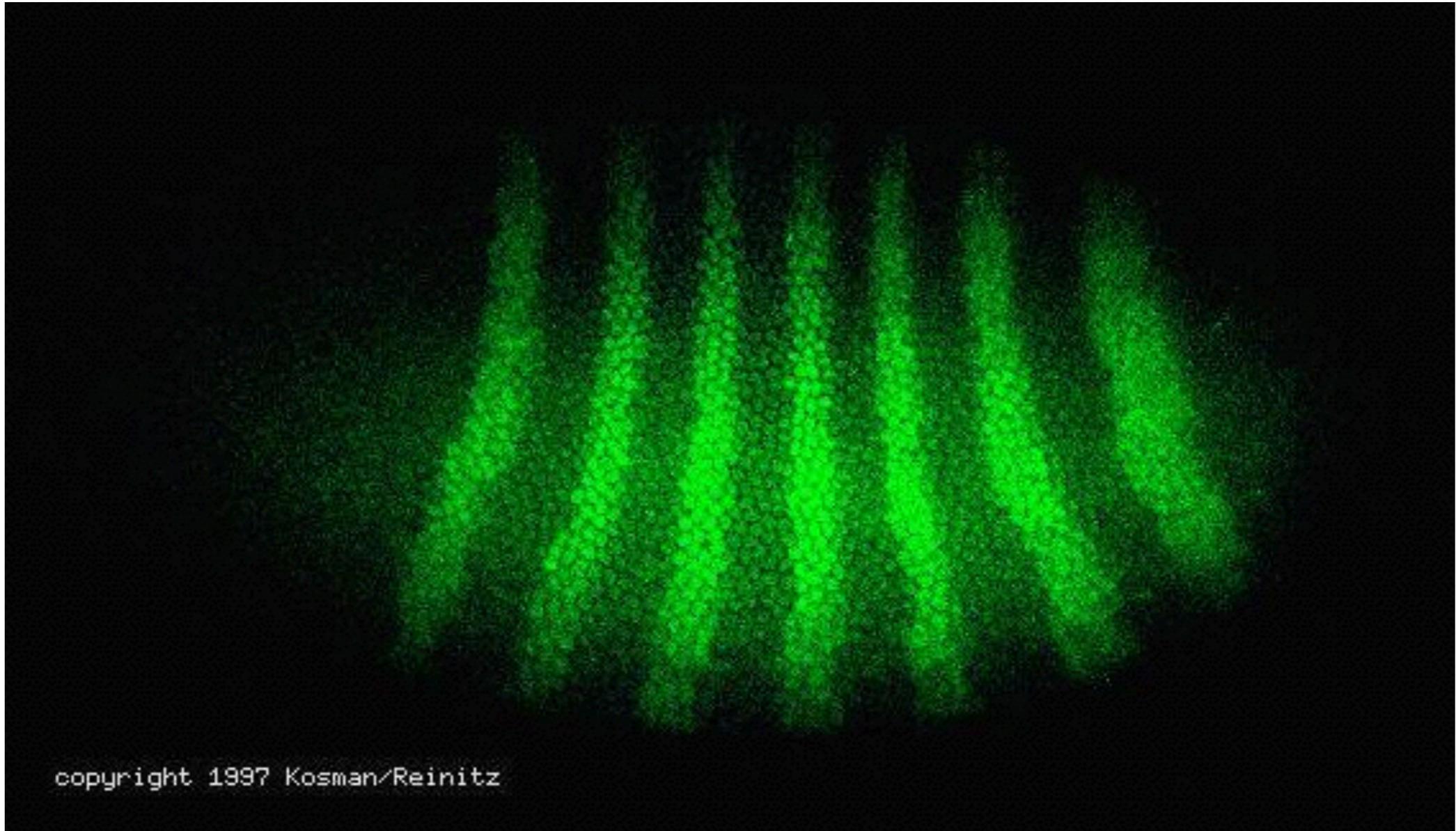


Hb (633nm)





Gregor Lab, Princeton



<http://www.iephb.nw.ru/hoxpro/ftz.html>

Philosophical Transactions of the Royal Society of London. Series B, Biological Sciences, Vol. 237, No. 641. (Aug. 14, 1952), pp. 37-72.

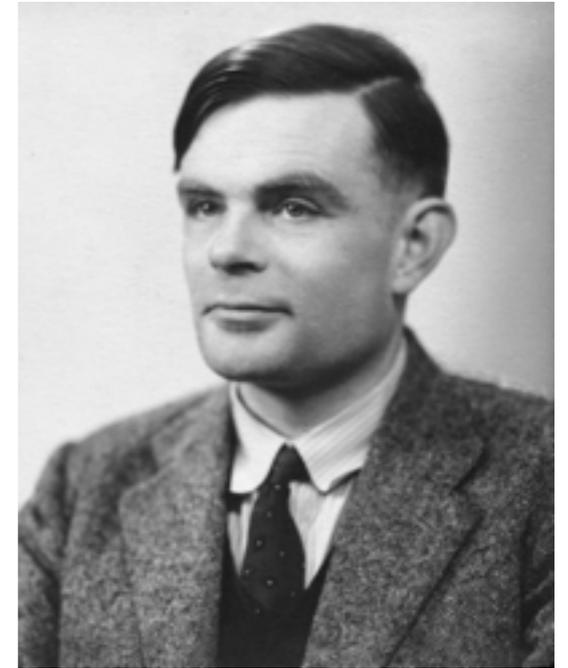
THE CHEMICAL BASIS OF MORPHOGENESIS

BY A. M. TURING, F.R.S. *University of Manchester*

(Received 9 November 1951—Revised 15 March 1952)

It is suggested that a system of chemical substances, called morphogens, reacting together and diffusing through a tissue, is adequate to account for the main phenomena of morphogenesis. Such a system, although it may originally be quite homogeneous, may later develop a pattern or structure due to an instability of the homogeneous equilibrium, which is triggered off by random disturbances. Such reaction-diffusion systems are considered in some detail in the case of an isolated ring of cells, a mathematically convenient, though biologically unusual system. The investigation is chiefly concerned with the onset of instability. It is found that there are six essentially different forms which this may take. In the most interesting form stationary waves appear on the ring. It is suggested that this might account, for instance, for the tentacle patterns on *Hydra* and for whorled leaves. A system of reactions and diffusion on a sphere is also considered. Such a system appears to account for gastrulation. Another reaction system in two dimensions gives rise to patterns reminiscent of dappling. It is also suggested that stationary waves in two dimensions could account for the phenomena of phyllotaxis.

The purpose of this paper is to discuss a possible mechanism by which the genes of a zygote may determine the anatomical structure of the resulting organism. The theory does not make any new hypotheses; it merely suggests that certain well-known physical laws are sufficient to account for many of the facts. The full understanding of the paper requires a good knowledge of mathematics, some biology, and some elementary chemistry. Since readers cannot be expected to be experts in all of these subjects, a number of elementary facts are explained, which can be found in text-books, but whose omission would make the paper difficult reading.

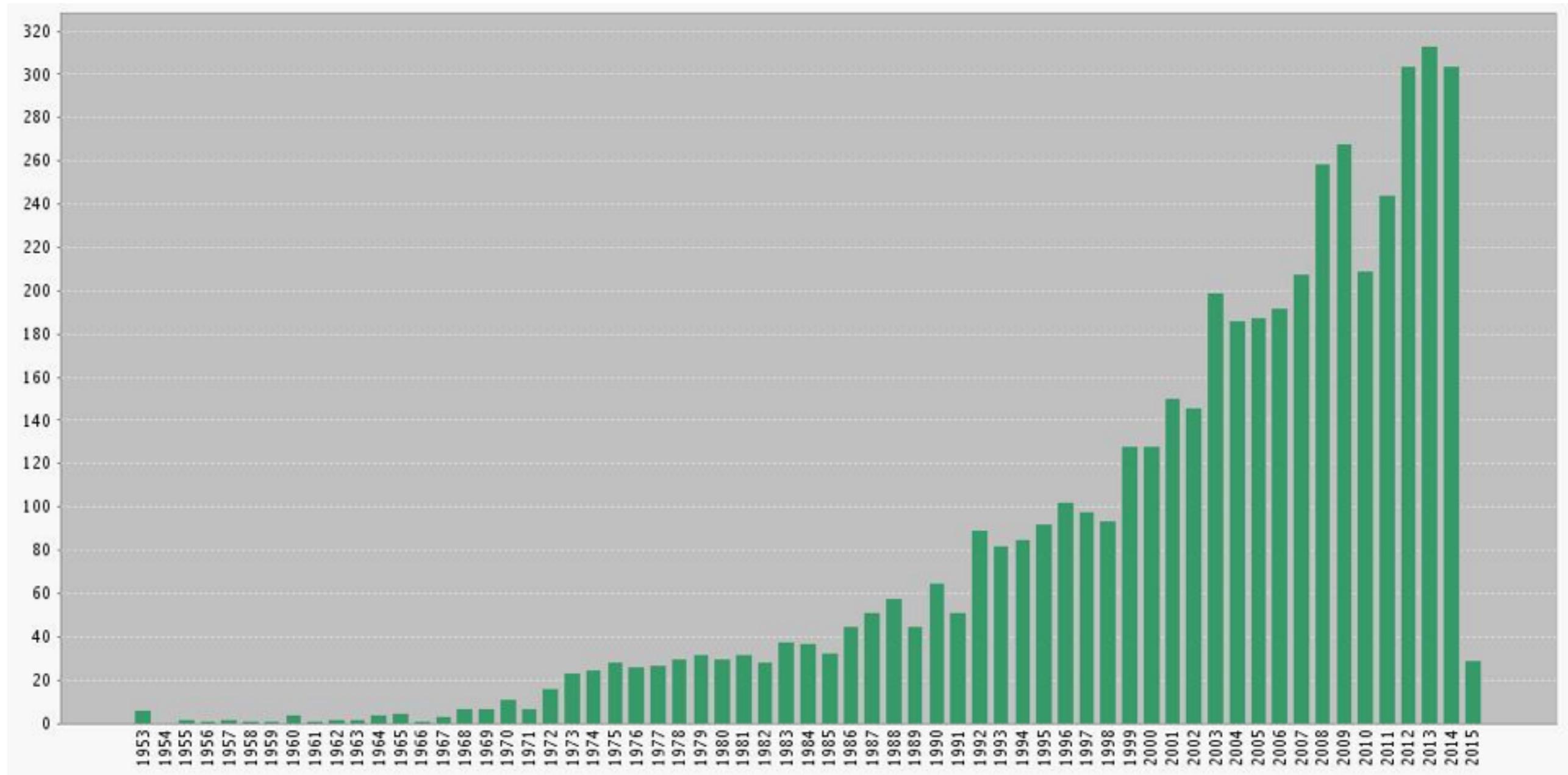


1912-1954

Google: 8721 citations

dunkel@math.mit.edu

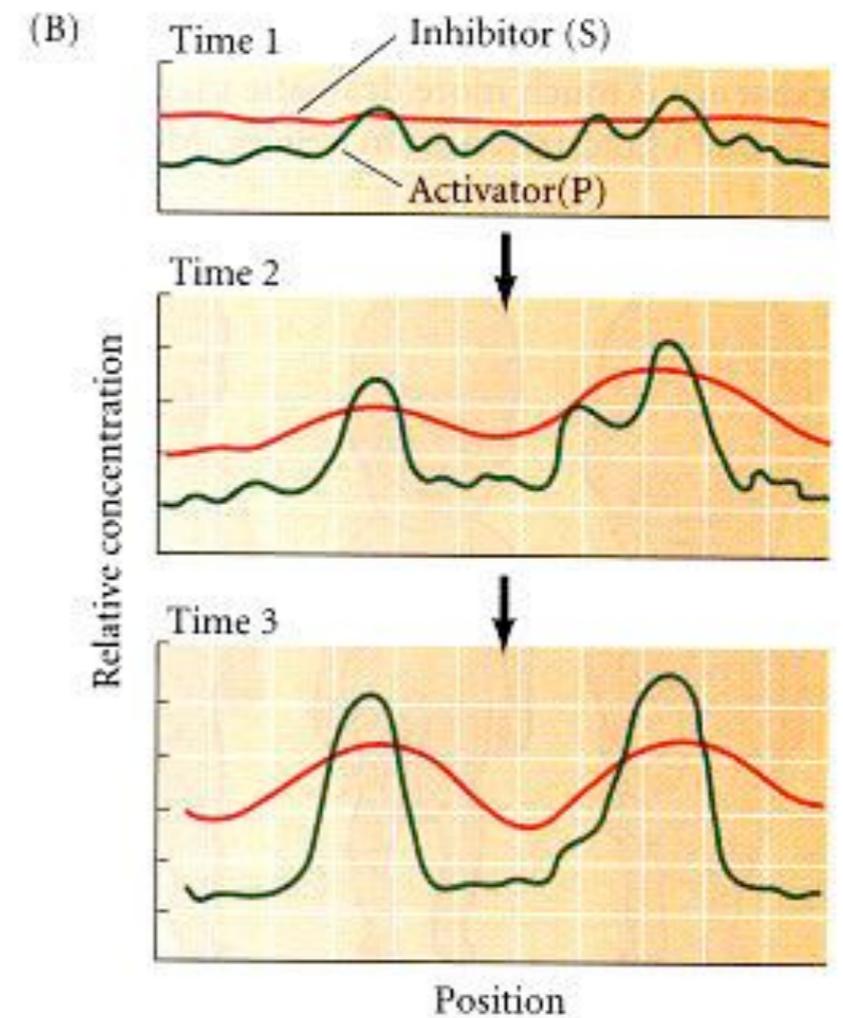
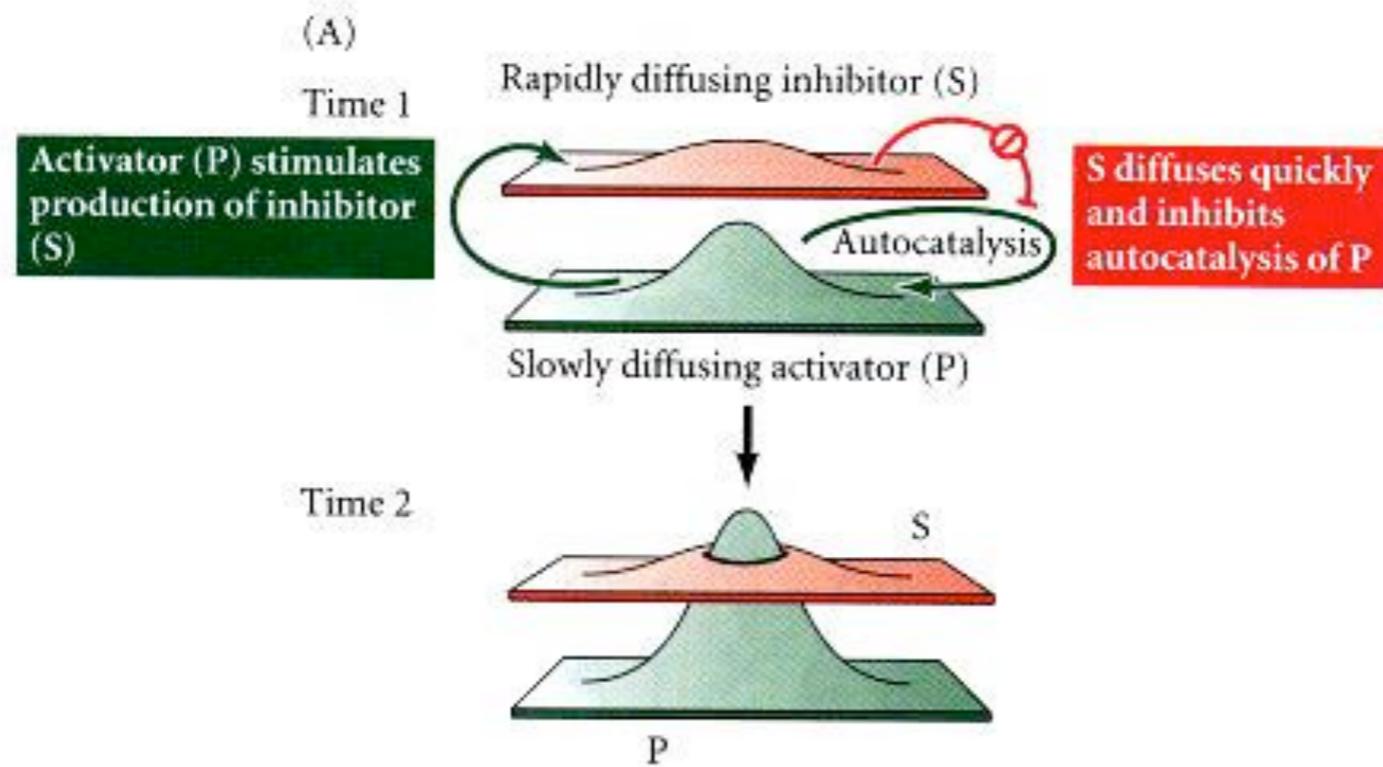
Philosophical Transactions of the Royal Society of London. Series B, Biological Sciences, Vol. 237, No. 641. (Aug. 14, 1952), pp. 37-72.



Web of Science >4800 citations

dunkel@math.mit.edu

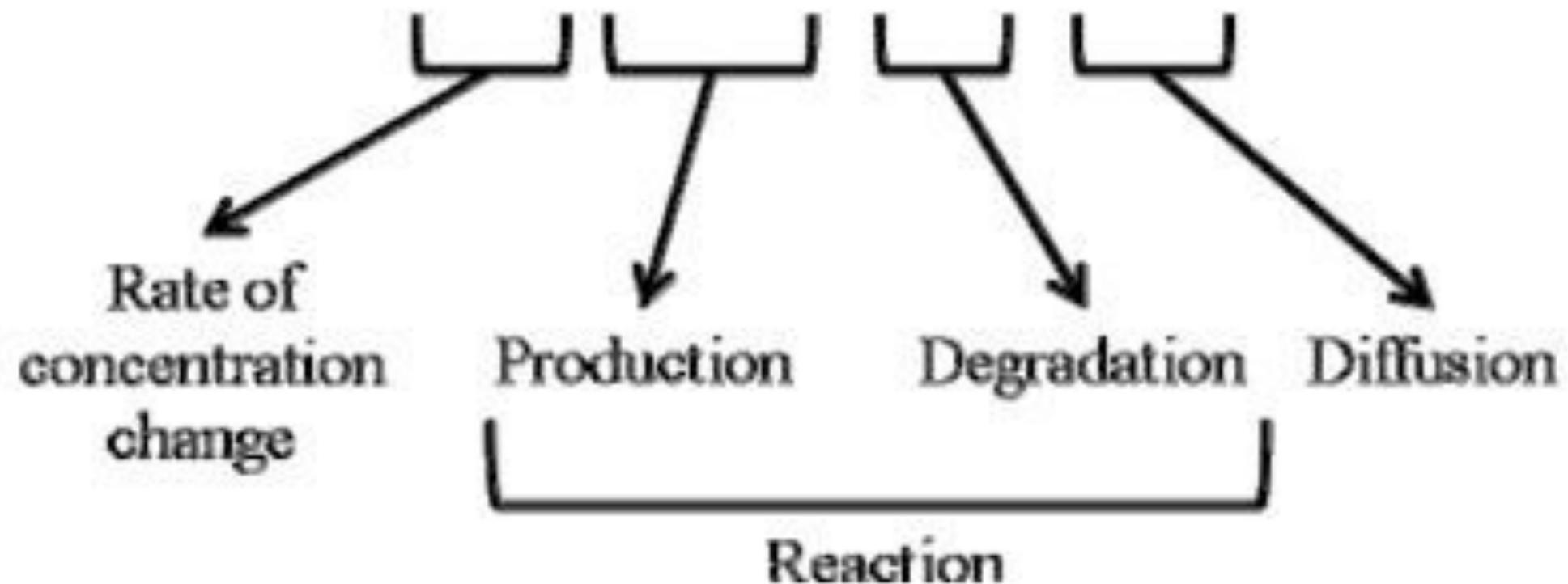
Turing model



Turing model

$$\frac{\partial u}{\partial t} = F(u, v) - d_u v + D_u \Delta u$$

$$\frac{\partial v}{\partial t} = G(u, v) - d_v v + D_v \Delta v$$



Turing pattern

Case VI (Turing pattern)



The matching of zebrafish stripe formation and a Turing model

