

(ii) chemotaxis

Deriving PDE models
continued

↳ movement in response to chemical signal

model

organism: *Dictyostelium discoideum*
(slime mold)

Single-cellular amoeba, eats bacteria

low food \rightarrow produce cAMP \rightarrow aggregate

goal: model the process of secreting cAMP

↳ aggregating in response to cAMP gradients

[again, could use cell-level agent-based model, likely to how many variables? \rightarrow v. complicated]

two densities $\left\{ \begin{array}{l} u(x,t) \text{ amoebae concentration} \\ c(x,t) \text{ concentration of cAMP} \end{array} \right.$

PDE model

note: run & tumble behavior

amoebae: diffusion + advection by external field

$v = \nabla c$ = gradient of cAMP

(why not birth & death? always consider relevant timescales)

cAMP: diffusion + creation + degradation

advection according

$u_t = D_u \Delta u - \chi \nabla \cdot (u \nabla c)$ to cAMP gradient

χ strength of response to gradient

$c_t = D_c \Delta c + \alpha u - \beta c$

prod. by amoebae degradation

Note : family of spatially homogeneous constant solns
 say we set $u(x,t) = u_0$ $c(x,t) = c_0$

$$u(x, t) = u_0$$

$$c(x, t) = c_0 \quad \forall x$$

$$\text{then } u_t = \cancel{D_u \vec{\nabla}^2 u} - \chi \vec{\nabla} \cdot (\vec{u} \vec{\nabla} c)$$

$$c_t = \cancel{D_c \vec{\nabla}^2 c} + \alpha u_0 - \beta c_0$$

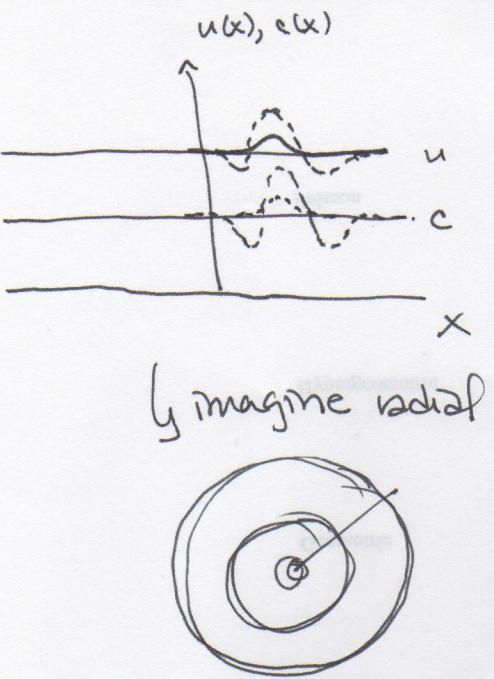
$$\Rightarrow \text{if } u_0 = \frac{\beta c_0}{\alpha} \text{ is steady soln for any } c$$

However, as w/ Turing bifurcation, spatially homogeneous steady state not nec. stable

↳ recall in activator-inhibitor system key was to have inhibitor diffuse much faster than activator to get pattern forming instability

↳ here more complicated, but similar idea, strength (rate of cAMP prod. / rate of response)
 roughly speaking want αX^* to dominate

βD_u (rate of degradation · diffusion of amoebae)



[only do if less than
very heuristic, γ_2 time gone]
like Bjorn's ex. of Turing)

(If response to cAMP & Incr.
cAMP prod. dominate
diffusion & cAMP degradation
can see this kind of pattern
forming behavior)

Spiral Fundamentally 2D - can't look at quite
this way

(iii) Calcium waves:

Nature review ~10 yr. ago:

"Almost everything we do is controlled by Ca^{2+} — how we move, how our hearts beat, and how our brains process information and store memories."

→ believe information in signals may have to do w/ wave frequency rather than amplitude

- intracellular oscillations

↳ Xenopus oocytes: spatial waves that propagate through cell

— full purpose not known, but involved in avoidance of polyspermy (multiple fertilization) & other key developmental processes



like 1D front

but also see spiral waves

targets (O)



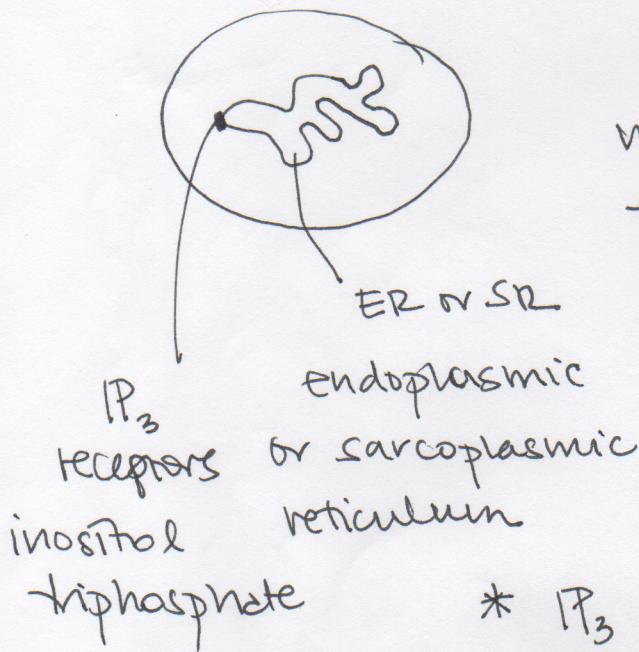
- intercellular oscillations

↳ airway epithelial cells: calcium waves generated by mechanical stimuli; purpose likely stimulation of cilia that move mucus: Ca^{2+} waves can coordinate movement of cilia (synchronized beat rhythm)

↳ contraction of heart muscle: [see visualizations on Scholarpedia]

- under normal conditions action potentials propagate along network of Purkinje fibers
(Ca^{2+} participates in creation of AP's)
- damaged tissue or changes in cellular structures can lead to abnormally long AP duration & shortened refractory times
 - ↳ can produce spiral waves that lead to tachycardia (heart rates 100-200 bpm)
 - or ventricular fibrillation (spiral breakup \rightarrow often fatal)

=



Ca^{2+} concentration in cytosol much lower than surrounding extracellular matrix \rightarrow actively pumped out (sequestered to ER)

- * $\text{IP}_3 \rightarrow \text{Ca}^{2+}$ released from ER to cytoplasm
- * Ca^{2+} exhibits positive feedback on short time scales (incr. activity of IP_3 receptors)
- * Ca^{2+} exhibits negative feedback on longer time scales

Model for Xenopus oocytes

(can use stochastic Markov-type models for individual receptors - we'll just use density spatio-temporal (PDE) model)

$$P_t = D_p \Delta P - \underbrace{k_p P}_{\text{degradation}}$$

could include production
↳ assume longer timescale/
not relevant here

$$C_t = D_c \Delta C + k_f \left(\mu_0 + \frac{\mu_1 P}{\mu_2 + P} \right) \left(b + \frac{(1-b)c}{k_1 + c} \right) h - \frac{\gamma c}{k_{xt} c} + \beta$$

↑
 rate const./
 scaling factor

IP₃ activation

calcium activation

pump out

calcium inactivation

leak M

[note what happens
when $P \rightarrow 0, \infty$; $C \rightarrow 0, \infty$]

$$\tau h_t = \frac{k_2}{k_2 + c} - h$$

say it starts at some const., $c=0$ what happens

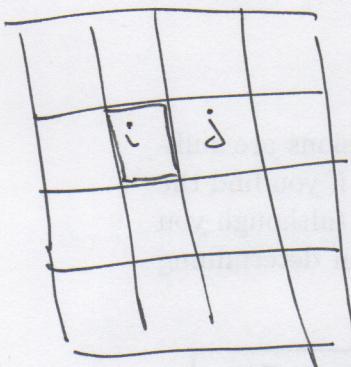
as $c \uparrow$, $\frac{k_2}{k_2 + c} \downarrow$ so ~ exponential decay

as h gets small, $\overset{(1)}{\downarrow}$ influx in c also gets small $\overset{(2)}{\downarrow}$

but note time scales not same

↳ this is how we capture initial pos. feedback
followed by later neg. feedback

To model intercellular waves, one approach is to use square lattice of cells w/ same internal dynamics as above, w/o Ca^{2+} across cells but P_3^* flux $\sqrt{2}$ gap junctions.



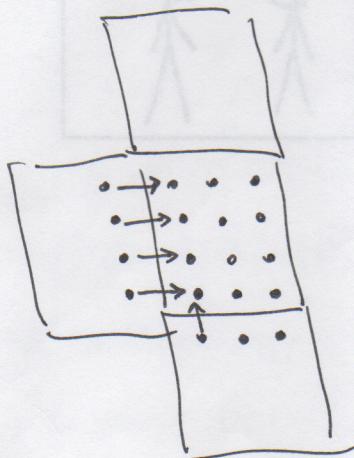
In particular use flux

$$\text{Flux} \propto k(p_i - p_j)$$

to couple cells

Solve numerically

using some # of grid points
inside cell and some # cells



Just scratching the surface - lots more in
papers posted on Canvas if interested
also Sneyd & Keener Math. Physiol.