Active Cells and Active Boundaries:

Analysis of a PDE/ODE model for Quorum and Diffusion Sensing in Biology

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Active Cells Coupled by Diffusion

Formulate and analyze a model of (ODE) dynamically active small "cells", with arbitrary intracellular kinetics, that are coupled spatially by a linear bulk-diffusion field (PDE) in a bounded 2-D domain.

Specific Questions:

- Can one trigger oscillations in the small cells (Hopf bifurcation), that would otherwise not occur without the coupling via bulk diffusion?
- Can we exhibit quorum sensing behavior by which cells oscillate and synchronize their dynamics when the population reaches a threshold?
 - In terms of the number m of cells per unit area, i.e. cell population density is $\rho = m/|\Omega|$.
 - What parameters regulate this threshold?
 - Usually studied from an ODE approach.
- Can we exhibit diffusion sensing behavior whereby cells oscillate and synchronize their dynamics based on:
 - cell spatial configuration (synchronization easier for clustered cells).
 - magnitude of diffusivity D of extracellular chemical (autoinducer).
 - Requires a PDE-based model.

Dynamical Quorum Sensing in Nature

Collective behavior in "cells" driven by chemical signalling between them.

- Collections of spatially segregated unicellular (eukaryotic) organisms such as starving yeast cells (glycolysis) coupled only through extracellular signalling molecules (autoinducer is Acetaldehyde). Ref: De Monte et al., PNAS 104(47), (2007).
- Amoeba colonies (Dicty) in low nutrient enviroments, with cAMP organizing the aggregation of starving colonies; Ref: Nanjundiah, Bio. Chem. 72, (1998), Gregor et al. Science, 328, (2010).
- Catalyst bead particles (BZ particles) interacting through a chemical diffusion field; Ref: Tinsley, Showalter, et al. "Dynamical Quorum Sensing... Collections of Excitable and Oscillatory Cataytic Particles", Physica D 239 (2010).

Key Ingredient: Need intracellular autocatalytic signal and an extracellular communication mechanism (bulk diffusion or autoinducer) that influences the autocatalytic growth. In the absence of coupling by bulk diffusion, the "cells" are in a quiescent state. Oscillations and ultimate sychronization occurs via a switchlike response to elevated levels of the autoinducer.

Amoeba Colony (Dictyostelium discoideum)

- About 180 cells are confined into an area of 420 μm in diameter (2-D).
- When resources are scarce, each cell secretes cAMP into the medium.
- Main Question: Is the oscillation an intrinsic property of the cells or does it only occur at the population level?



Caption: The cells secrete cAMP into the medium which first initiates a coordinated collective response.

On longer time-scale cells aggregate. **Ref:** The Onset of Collective Behavior in Social Amoebae, T. Gregor et al. Science 2010

Modeling Approaches

Large ODE system of weakly coupled system of oscillators. Prototypical is the Kuramoto type-models for the coupled phases of the oscillators, of the form

$$\frac{d\mathbf{x}_i}{dt} = \mathbf{F}(\mathbf{x}_i) + \sigma \sum_j C_{ij} \mathbf{H}(\mathbf{x}_j) \,,$$

Synchrony occurs between individual oscillators as the coupling strength σ increases. (Vast literature, but not the mechanism here).

- Homogenization approach of deriving RD systems through cell densities: Yields target and spiral wave patterns of cAMP in Dicty modeling (but phemenological).
- More Recent: PDE-ODE models coupling individual "cells" through a bulk diffusion field. Our framework related to:
 - Ref: J. Muller, C. Kuttler, et al. "Cell-Cell Communication by Quorum Sensing and...", J. Math. Bio. 53 (2006),
 - J. Muller, H. Uecker, J. Math. Bio. 67 (2013). (steady-state analysis in 3-D, dynamics).

Ref [GW]: J. Gou, M.J. Ward, J. Nonlinear Sci., 26(4), (2016), pp. 979–1029.

Formulation of the 2-D Model: I



- The *m* cells are circular and each contains *n* chemicals $\mu_j = (\mu_{1j}, \dots, \mu_{nj})^T$. When isolated they interact via ODE's $d\mu_j/dt = \mathbf{F}_j(\mu_j)$.
 - A scalar bulk diffusion field (autoinducer) diffuses in the space between the cells via

 $\mathcal{U}_T = D_B \Delta_X \mathcal{U} - k_B \mathcal{U} \,.$

There is an exchange across the cell membrane, regulated by permeability parameters, between the autoinducer and one intracellular species (Robin condition).

Scaling Limit: $\epsilon \equiv \sigma/L \ll 1$, where *L* is lengthscale for Ω . We assume that the permeability parameters are $\mathcal{O}(\epsilon^{-1})$. Parameters: Bulk diffusivity D_B , bulk decay k_B , permeabilities, ϵ , and time-scale of intracellular reactions.

Formulation of the 2-D Model: II

Our PDE-ODE coupled cell-bulk model in 2-D with m cells is

$$\mathcal{U}_T = \mathbf{D}_B \Delta_{\mathbf{X}} \mathcal{U} - \mathbf{k}_B \mathcal{U}, \quad \mathbf{X} \in \Omega \setminus \bigcup_{j=1}^m \Omega_j; \quad \partial_{n_{\mathbf{X}}} \mathcal{U} = 0, \quad \mathbf{X} \in \partial \Omega,$$
$$D_B \partial_{n_{\mathbf{X}}} \mathcal{U} = \beta_{1j} \mathcal{U} - \beta_{2j} \mu_j^1, \quad \mathbf{X} \in \partial \Omega_j, \quad j = 1, \dots, m.$$

Each cell $\Omega_j \in \Omega$ is a disk of radius σ centered at some $X_j \in \Omega$.

Inside each cell there are *n* interacting species with mass vector $\mu_j \equiv (\mu_j^1, \dots, \mu_j^n)^T$ whose dynamics are governed by *n*-ODEs, with (rank-one) coupling via integration over the *j*-th "cell"-membrane $\partial \Omega_j$:

$$\frac{d\boldsymbol{\mu}_j}{dT} = \boldsymbol{k_R} \mu_c \boldsymbol{F}_j \left(\boldsymbol{\mu}_j / \mu_c \right) + \boldsymbol{e}_1 \int_{\partial \Omega_j} \left(\boldsymbol{\beta}_{1j} \mathcal{U} - \boldsymbol{\beta}_{2j} \mu_j^1 \right) \, dS_j \,, \quad j = 1, \dots, m \,,$$

where $e_1 \equiv (1, 0, \dots, 0)^T$, and μ_c is typical mass.

- Only one species μ_i^1 can cross the *j*-th cell membrane into the bulk.
- $k_R > 0$ is intracellular reaction rate; β_{1j} , β_{2j} are permeabilities.
- The dimensionless function $F_j(u_j)$ models the intracellular dynamics.

Formulation of the 2-D Model: III

<u>Dimensionless Formulation</u>: The concentration of signalling molecule U(x, t) in the bulk satisfies the PDE:

$$\tau U_t = \mathbf{D}\Delta U - U, \qquad \mathbf{x} \in \Omega \setminus \bigcup_{j=1}^m \Omega_{\epsilon_j}; \quad \partial_n U = 0, \quad \mathbf{x} \in \partial \Omega,$$

$$\epsilon \mathbf{D}\partial_{n_j} U = \mathbf{d_{1j}} U - \mathbf{d_{2j}} u_j^1, \qquad \mathbf{x} \in \partial \Omega_{\epsilon_j}, \quad j = 1, \dots, m.$$

The cells are disks of radius $\epsilon \ll 1$ so that $\Omega_{\epsilon_j} \equiv \{x \mid |x - x_j| \le \epsilon\}$.

Inside each cell there are *n* interacting species $u_j = (u_j^1, \ldots, u_j^n)^T$, with intracellular dynamics for each $j = 1, \ldots, m$,

$$\frac{d\boldsymbol{u}_j}{dt} = \boldsymbol{F}_j(\boldsymbol{u}_j) + \frac{\boldsymbol{e}_1}{\epsilon\tau} \int_{\partial\Omega_{\epsilon_j}} (\boldsymbol{d}_{1j}U - \boldsymbol{d}_{2j}u_j^1) \, ds \,, \qquad \boldsymbol{e}_1 \equiv (1, 0, \dots, 0)^T$$

<u>**Remark:</u>** The time-scale is measured wrt intracellular reactions. The dimensionless bifurcation parameters are: d_{1j} , d_{2j} (permeabilities); τ (reaction-time ratio); D (effective diffusivity);</u>

$$\tau \equiv \frac{k_R}{k_B}, \quad D \equiv \left(\frac{\sqrt{D_B/k_B}}{L}\right)^2, \quad \beta_{1j} \equiv (k_B L) \frac{d_{1j}}{\epsilon}, \quad \beta_{2j} \equiv \left(\frac{k_B}{L}\right) \frac{d_{2j}}{\epsilon}.$$

Theoretical Framework

- Can one trigger oscillations in the small cells, via a Hopf bifurcation, that would otherwise not be present without the coupling via bulk diffusion? (i.e. each cell is a conditional oscillator). Intuition: Need reaction-time ratio τ in some interval $0 < \tau_{-} < \tau < \tau_{+} < \infty$.
- Can we exhibit quorum sensing and diffusion sensing behavior?

Three key regimes for D with different behaviors:

- D = $\mathcal{O}(1)$; Effect of spatial distribution of cells is a key factor whether oscillations are triggered or not (diffusion sensing behavior).
- D = $O(\nu^{-1})$ with $\nu = -1/\log \epsilon$; HB thresholds can occur for both synchronous and asynchronous modes. Spatial location of cells not important to leading order.
- $D \gg \mathcal{O}(\nu^{-1})$; In this "well-mixed" regime, the PDE-ODE cell-bulk model reduces to a finite dimensional dynamical system with global coupling. Quorum sensing behavior observed.
- Mathematical Framework: Use strong localized perturbation theory (SLPT) to construct steady-states, to formulate the linear stability problem, and to derive the limiting well mixed ODE system.

Steady-States: Matched Asymptotics

Main Result (Steady-State): In the outer region, the ss bulk diffusion field is

$$U(\boldsymbol{x}) = -2\pi \sum_{i=1}^{m} S_i G(\boldsymbol{x}, \boldsymbol{x}_i), \text{ where } \boldsymbol{S} \equiv (S_1, \dots, S_m)^T.$$

In terms of $\nu = -1/\log \epsilon$ and a Green's matrix \mathcal{G} , we obtain a nonlinear algebraic system for \mathbf{S} and $\mathbf{u}^1 \equiv (u_1^1, \dots, u_m^1)^T$, where $e_1 = (1, 0, \dots, 0)^T$:

$$\boldsymbol{F}_{j}(\boldsymbol{u}_{j}) + \frac{2\pi D}{\tau} \boldsymbol{S}_{j} \boldsymbol{e}_{1} = 0, \quad (\mathcal{H} + 2\pi \nu \mathcal{G}) \boldsymbol{S} = -\nu \mathcal{W} \boldsymbol{u}^{1}, \quad j = 1, \dots, m.$$

Here
$$\mathcal{W} \equiv \operatorname{diag}\left(\frac{d_{21}}{d_{11}}, \dots, \frac{d_{2m}}{d_{1m}}\right)$$
 and $\mathcal{H} \equiv \operatorname{diag}\left(\left(1 + \frac{\nu D}{d_{11}}\right), \dots, \left(1 + \frac{\nu D}{d_{1m}}\right)\right)$.

In this ss formulation, the entries of the $m \times m$ Green's matrix \mathcal{G} are

$$(\mathcal{G})_{ii} = R_i, \qquad (\mathcal{G})_{ij} = G(\boldsymbol{x}_i; \boldsymbol{x}_j), \quad i \neq j,$$

where, with $\varphi_0 \equiv 1/\sqrt{D}$, $G(\boldsymbol{x}; \boldsymbol{x}_j)$ is the reduced-wave G-function:

$$\begin{split} \Delta G - \varphi_0^2 G &= -\delta(\boldsymbol{x} - \boldsymbol{x}_j) \,, \quad \boldsymbol{x} \in \Omega \,; \qquad \partial_n G = 0 \,, \quad \boldsymbol{x} \in \partial \Omega \,. \\ G(\boldsymbol{x}; \boldsymbol{x}_j) &\sim -\frac{1}{2\pi} \log |\boldsymbol{x} - \boldsymbol{x}_j| + R_j + o(1) \,, \qquad \text{as} \quad \boldsymbol{x} \to \boldsymbol{x}_j \,. \end{split}$$

Globally Coupled Eigenvalue Problem (GCEP)

Linear Stability: For $\epsilon \to 0$, the perturbed bulk diffusion field satisfies

$$u(\boldsymbol{x},t) = U(\boldsymbol{x}) + e^{\lambda t} \eta(\boldsymbol{x}), \qquad \eta(\boldsymbol{x}) = -2\pi \sum_{i=1}^{m} c_i G_{\lambda}(\boldsymbol{x},\boldsymbol{x}_i).$$

Inside the *j*-th cell we have $u_j = u_{ej} + 2\pi D\tau^{-1}c_j e^{\lambda t}(\lambda I - J_j)^{-1}e_1$. Here $c = (c_1, \dots, c_m)^T$ is a nullvector of the GCEP:

$$\mathcal{M}\mathbf{c} = \mathbf{0}, \qquad \mathcal{M}(\lambda) \equiv 2\pi\nu\mathcal{G}_{\lambda} + \mathcal{H} + \nu\frac{2\pi D}{\tau}\mathcal{W}\mathcal{K}(\lambda).$$

In this GCEP, \mathcal{G}_{λ} is the Green's matrix formed from

$$egin{aligned} &\Delta G_\lambda - arphi_\lambda^2 G_\lambda = -\delta(oldsymbol{x} - oldsymbol{x}_j), \quad oldsymbol{x} \in \Omega\,; &\partial_n G_\lambda = 0\,, \quad oldsymbol{x} \in \partial \Omega\,, \ &G_\lambda(oldsymbol{x};oldsymbol{x}_j) &\sim -rac{1}{2\pi} \log |oldsymbol{x} - oldsymbol{x}_j| + R_{\lambda,j} + o(1)\,, & ext{as} \quad oldsymbol{x} o oldsymbol{x}_j\,, \end{aligned}$$

with $\varphi_{\lambda} \equiv D^{-1/2}\sqrt{1+\tau\lambda}$. Here \mathcal{K} is the diagonal matrix defined in terms of the Jacobian $J_j \equiv \mathbf{F}_{j,\mathbf{u}}(\mathbf{u}_{ej})$ of the intracellular kinetics \mathbf{F}_j :

$$\mathcal{K}_{j} = e_{1}^{T} (\lambda I - J_{j})^{-1} e_{1} = \frac{M_{j,11}(\lambda)}{\det(\lambda I - J_{j})}, \text{ where } e_{1} = (1, 0, \dots, 0)^{T}.$$

Shanghai – p.11

Numerics for the GCEP

Linear stability analysis: Nonlinear matrix eigenvalue problem of the form

 $\mathcal{M}(\lambda;\tau,D)\boldsymbol{c}=\boldsymbol{0}\,,$

<u>Definition</u>: An unstable "mode" is a root λ of $\mathcal{F}(\lambda) = \det(\mathcal{M}(\lambda)) = 0$ in $\operatorname{Re}(\lambda) > 0$. The number N of unstable modes is the total number of such roots. The eigenvector c determines the amplitude and phase at each cell.

- Determine N numerically from winding number computation of $\mathcal{F}(\lambda)$ over a large semi-circle in $\text{Re}(\lambda) > 0$. Gives a "stability map" in (τ, D) plane with N = 0 (white), N = 2 (grey), N = 4 (blue), etc..
- Hopf bifurcation boundaries, $\lambda = i\lambda_I(D)$ and $\tau = \tau(D)$ can have folds in *D*. Compute with $\text{Re}\mathcal{F} = 0$ and $\text{Im}\mathcal{F} = 0$ using psuedo-arclength.



Tractable: Ring and Ring + Center Hole Pattern:

- Small identical cells inside unit disk, evenly spaced on a concentric ring of radius r_0 .
- The center-cell can have different kinetics, or different permeabilities d_1 and d_2 .
- Matrix spectrum $\mathcal{M} oldsymbol{c} = \sigma oldsymbol{c}$ available analytically.

Intracellular Selkov Reaction-Kinetics

<u>Selkov Kinetics</u>: Let $u = (u_1, u_2)^T$ be intracellular dynamics given by Selkov model (used for modeling glycolysis oscillations):

$$F_1(u_1, u_2) = \alpha u_2 + u_2 u_1^2 - u_1, \quad F_2(u_1, u_2) = \epsilon_0 \left(\mu - (\alpha u_2 + u_2 u_1^2) \right)$$

For an *isolated cell* \exists a unique steady-state at $u_{1e} = \mu$, $u_{2e} = \mu/(\alpha + \mu^2)$. The determinant and trace of the Jacobian J_e is

trace
$$(J_e) = \frac{\left[\mu^2 - \alpha^2 - \epsilon_0(\alpha + \mu^2)^2\right]}{\alpha + \mu^2}$$
, det $(J_e) = \varepsilon_0(\alpha + \mu^2) > 0$.



- Fix Selkov parameters as $\alpha = 0.9$, and $\epsilon_0 = 0.15$ and plot versus μ .
- For $\mu = 2$ an isolated cell has a stable fixed point with no oscillations, but is near to stability threshold.

<u>Remark:</u> When coupled to the other cells there is a new (but unique) steady-state and the PDE-ODE coupling can trigger oscillations via a HB.

$D = \mathcal{O}(1)$: Ring Patterns



Analytically Tractable Example:

- m small cells inside the unit disk, evenly spaced on a concentric ring of radius r_0 .
- Assume identical kinetics and permeabilities, so that $F_j = F$, $d_{1j} = d_1$, and $d_{2j} = d_2$.

Spectral Problem (from GCEP): Must find the roots λ to $\mathcal{B}_j(\lambda) = 0$, where

$$\mathcal{B}_{j}(\lambda) \equiv \omega_{\lambda,j} + \frac{1}{2\pi\nu} \left(1 + \frac{D\nu}{d_{1}}\right) + \left(\frac{d_{2}D}{d_{1}\tau}\right) \frac{M_{11}}{\det(\lambda I - J)}, \qquad j = 1, \dots, m.$$

Here $\omega_{\lambda,j}$ are the eigenvalues of the λ -dependent Green's matrix \mathcal{G}_{λ} :

$$\mathcal{G}_{\lambda} \boldsymbol{v}_j = \boldsymbol{\omega}_{\boldsymbol{\lambda}, \boldsymbol{j}} \boldsymbol{v}_j, \qquad \boldsymbol{j} = 1, \dots, m,$$

- \exists a steady-state with $S_j = S_c$ for all j = 1, ..., m.
- **For the unit disk**, the Green's matrix \mathcal{G}_{λ} is given analytically in terms of an infinite series of modified Bessel functions of complex argument.

$D = \mathcal{O}(1)$: Ring Patterns: II

Linear Stability Computations (Theory):

- Phase Diagram: Compute Hopf Bifurcation (HB) boundaries in the τ versus D plane for each j = 1, ..., m by setting $\lambda = i\lambda_I$. Fix r_0 , $\epsilon = 0.05$, $d_1 = 0.8$, and $d_2 = 0.2$.
- Winding Number computations used to check where $\text{Re}(\lambda) > 0$ in open regions of the τ versus D plane.
- Cyclic Symmetric Matrices:: Matrix spectrum of \mathcal{G}_{λ} readily calculated. Note: $\mathbf{v}_1 = \mathbf{e} \equiv (1, \dots, 1)^T$ (synchronous mode), while $\mathbf{e}^T \mathbf{v}_j = 0$ for $j = 2, \dots, m$ are the asynchronous modes. However, mode degeneracy occurs due to cyclicity and symmetry of \mathcal{G}_{λ} . In particular, if m = 5, there are exactly two asynchronous branches.

Qualitative Questions: What is the effect of:

- \checkmark cell clustering (i.e. smaller r_0 ?)
- \checkmark the cell permeabilities d_1 and d_2 ?
- \checkmark the number m of cells?
- small changes in the intracellular kinetics?

$D = \mathcal{O}(1)$: HB Boundaries: m = 2 Cells



- HB boundaries τ vs. D for m = 2and $r_0 = 0.75$.
- Synchronous and asynchronous HB boundaries (heavy dashed).
- N = 2 (grey) and N = 4 (blue). (winding-number results)
- Asynchronous lobe exists only for D small.
- Predicts no oscillations for $D \gg 1$.

Shanghai - p.16

<u>Numerical Validation</u>: FlexPDE for a similar map with $r_0 = 0.25$





Caption: Let m = 2 and vary r_0 : HB boundaries in τ versus D for the synchronous mode (larger lobes) and the asynchronous mode (small lobes for D small).

- Asynchronous lobe is smallest when $r_0 = 0.25$ (i.e. for closely-spaced cells). Implies that *D* has to be only increased a bit before asynchronous oscillations are impossible.
- If $r_0 = 0.75$ the two cells are rather close to their images across the boundary of the disk (Neumann BC).
- Diffusion sensing: If D = 5 and $\tau = 0.6$, we are outside instability lobe for $r_0 = 0.5$ but within the lobes for $r_0 = 0.25$ and $r_0 = 0.75$. Thus a more clustered configuration will trigger oscillations for the same D.

$D = \mathcal{O}(1)$: HB Boundaries m = 5

HB boundaries: m = 5 cells and $r_0 = 0.5$.(Right is zoom of left)



•
$$N = 2$$
 (grey), $N = 6$ (red), $N = 10$ (cyan).

• Asynchronous lobes: only for D small. Two such lobes when m = 5.

Instability lobe for synchronous mode is now unbounded (left figure).

Implication: The unbounded lobe for the synchronous mode indicates that for the well-mixed limit $D \to \infty$ a Hopf bifurcation for the steady-state will occur when $\tau = \tau_{\pm}$ (horizontal asymptotes), and that an oscillatory instability occurs for $\tau_{-} < \tau < \tau_{+}$.

Ring + Center Pattern: Role of Permeability

Consider m = 5 with a defective cell at the center of the disk with different permeabilities than four identical cells on a ring of radius $r_0 = 0.75$.



- **Ring Cells:** $d_1 = 0.8$, $d_2 = 0.2$ (identical kinetics)
- Center Cell: Case I: $d_1 = 0.8$, $d_2 = 0.2$. Case II (Defective): $d_1 = 0.4$, $d_2 = 0.2$.
- \mathcal{M} is a 5×5 symmetric matrix with a 4×4 cyclic block with the fifth row being (b, b, b, b, r).



Caption: Left: Case I: all identical. Middle: Case II: center defective. Right: Zoom for small D with N = 0 (white), N = 2 (grey), N = 4 (blue), N = 6 (red), N = 8 (green), N = 10 (cyan).

Ring + Center Pattern: A Triggering Center Cell

Consider m = 5 with a defective cell at the center of the disk that has a different intracellular kinetic (Selkov) parameter closer to stability threshold of an isolated cell than the four identical cells on the ring.



Caption: Lobes of instability for the synchronous mode $c = (1, 1, 1, \xi)$: Left: all identical cells $d_1 = 0.3, d_2 = 0.2, \alpha = 0.9$. Right: center-cell has $\alpha = 0.86$.

- With more clustering ($r_0 = 0.25$), one can have a larger bulk diffusivity D before autoinducer wanders too far from cells to trigger collective behavior.

The Distinguished Limit $D = D_0/\nu$

Simplification: Assume identical intracellular dynamics: so $F_j = F$, $\forall j$:

•
$$G \sim D/|\Omega| + \mathcal{O}(1)$$
 and $G_{\lambda} \sim D/[(1 + \tau\lambda)|\Omega|] + \mathcal{O}(1)$ for $D \gg 1$.

- To leading order, the source strengths are independent of the locations of cells. No spatial information to leading order in $\nu = -1/\log \epsilon$.
- The GCEP becomes: $(\sigma_1 I + \sigma_2 E)c = 0$ for some $\sigma_j(\lambda)$, where $E = ee^T$ and $e = (1, ..., 1)^T$. Thus, $\exists m - 1$ asynchronous modes $c = q_j$, with $q_j^T e = 0$ for j = 2, ..., m. The synchronous mode is c = e.

Lemma: Steady-state is linearly stable to synchronous perturbations iff

$$\frac{M_{11}(\lambda)}{\det(\lambda I - J)} = -\frac{\tau}{2\pi d_2} \left(\frac{\kappa_1 \tau \lambda + \kappa_2}{\tau \lambda + 1}\right); \quad \kappa_1 \equiv \frac{d_1}{D_0} + 1, \ \kappa_2 \equiv \kappa_1 + 2\pi d_1 \frac{m}{|\Omega|},$$

has no eigenvalue in $\text{Re}(\lambda) > 0$. Here *J* is the Jacobian of F(u) at the leading-order steady-state for $D = O(\nu^{-1})$. $M_{11}(\lambda)$ is the (1, 1) cofactor.

<u>Lemma</u>: For $m \ge 2$, the steady-state is linearly stable to the asynchronous or competition modes iff no eigenvalue in $\text{Re}(\lambda) > 0$ for

$$\frac{M_{11}}{\det(\lambda I - J)} = -\frac{\tau}{2\pi d_2} \left(\frac{d_1}{D_0} + 1\right)$$

The Distinguished Limit $D = D_0/\nu$: II

Lemma: For n = 1 then no HB is possible for any intracellular dynamics F.

Next, let n = 2, so that there are two intracellular species $(u_1, u_2)^T$:

Synchronous Mode: Then, λ satisfies the cubic

$$\mathcal{H}(\lambda) \equiv \lambda^3 + \lambda^2 p_1 + \lambda p_2 + p_3 = 0; \qquad p_1 \equiv \tau^{-1} (\gamma + \zeta) - \operatorname{tr}(J),$$
$$p_2 \equiv \det(J) - \frac{\gamma}{\tau} G_{u_2}^e + \frac{1}{\tau} \left(\frac{\gamma}{\tau} - \zeta \operatorname{tr}(J) \right), \qquad p_3 \equiv \frac{1}{\tau} \left(\zeta \det(J) - \frac{\gamma}{\tau} G_{u_2}^e \right).$$

Here γ and ζ are defined in terms of the effective "cell density" $m/|\Omega|$ by

$$\gamma \equiv \frac{2\pi d_2 D_0}{d_1 + D_0} > 0, \qquad \zeta \equiv 1 + \frac{2\pi d_1 D_0}{(d_1 + D_0)} \frac{m}{|\Omega|} > 1.$$

HB criterion: By Routh-Hurwitz we must have $p_1 > 0$, $p_3 > 0$, $p_1p_2 = p_3$.

Asynchronous Mode: When n = 2, λ satisfies the quadratic

$$\lambda^2 - \lambda q_1 + q_2 = 0$$
; where $q_1 \equiv \operatorname{tr}(J) - \frac{\gamma}{\tau}$, $q_2 \equiv \det(J) - \frac{\gamma}{\tau} G_{u_2}^e$

HB criterion: we must have $q_1 = 0$ and $q_2 > 0$.

The Well-Mixed Regime $D \gg \mathcal{O}(\nu^{-1})$: I

Goal: Derive and analyze a reduced finite-dimensional dynamical system characterizing the cell-bulk interations from PDE-ODE system for $D \rightarrow \infty$.

An asymptotic analysis yields that in the bulk that $u(x,t) \sim U_0(t)$, where

$$egin{aligned} U_0'&=-rac{1}{ au}U_0-rac{oldsymbol{
ho}}{ au}\left(rac{1}{m}\sum_{j=1}^m\left[\kappa_{1,j}U_0-\kappa_{2,j}u_j^1
ight]
ight)\,,\ u_j'&=oldsymbol{F}_j(oldsymbol{u}_j)+rac{1}{ au}\left[\kappa_{1,j}U_0-\kappa_{2,j}u_j^1
ight]oldsymbol{e}_1\,,\qquad j=1\,,\ldots,m\,, \end{aligned}$$

where $e_1 = (1, 0, ..., 0)^T$. Here ρ is the effective cell density and

$$\rho \equiv \frac{m}{|\Omega|}, \qquad \kappa_{1,j} \equiv 2\pi d_{1,j}, \qquad \kappa_{2,j} \equiv 2\pi d_{2,j}.$$

Large system of ODEs with weak but global coupling when $0 < d_{1j} << 1$ and $0 < d_{2j} \ll 1$, or when $\tau \gg 1$.

<u>Identical Cells</u>: Look for $u_j = u$, $\forall j$. We get

$$U_0' = -\frac{1}{\tau} (1 + \kappa_1 \rho) U_0 + \rho \frac{\kappa_2}{\tau} u_1, \qquad u' = F(u) + \frac{1}{\tau} [\kappa_1 U_0 - \kappa_2 u_1] e_1.$$

The Well-Mixed Regime $D \gg \mathcal{O}(\nu^{-1})$: II Selkov with $d_1 = 0.8$, $d_2 = 0.2$ and $|\Omega| = \pi$. Global Bifurcation Study.



Caption: Global solution branches u_{1e} versus τ for m = 5 cells: Heavy (thin) solid is stable (unstable) steady-steady. Dots indicate stable periodic solution branch. HB points at $\tau_{H-} =$ 0.2187 and $\tau_{H+} = 0.6238$.

Key: Stable synchronous oscillations occur in some τ interval. Limiting well-mixed ODE dynamics is independent of cell locations and D.

Quorum sensing (Qualitative): Collective behavior of "cells" in response to changes in their population size. There is a threshold number m_c of cells or a critical cell density ρ that is needed to initiate a collective behavior.

Quorum sensing (Math): For what range of m, do the well-mixed ODEs have a stable periodic solution on $\tau_{H-} < \tau < \tau_{H+}$ with HB points at $\tau_{H\pm}$?

Quorum Sensing Behavior

What parameters control control QS behavior? We will study QS behavior as the permeability d_1 is varied and $d_2 = 0.2$: Recall:

 $\partial_{n_j} U = \mathbf{d_1} U - \mathbf{d_2} u_j^1$, on $\partial \Omega_{\mathcal{E}_j}$, $j = 1, \dots, m$.

<u>**Remark:**</u> Equivalent to finding the range of m for which the instability lobe for the synchronous mode is unbounded in the τ versus D plane.

Left: Quorum threshold m_c vs. d_1 from ODEs. Right: τ vs. D for $d_1 = 0.3$, $r_0 = 0.5$.



Key: m_c sensitive to small changes in d_1 $d_1 = 0.8$, $m_c = 3$; $d_1 = 0.3$, $m_c = 7$; $d_1 = 0.2$, $m_c = 12$; $d_1 = 0.1$, $m_c = 19$.

Large Cell Populations: Synchronization I

In the well-mixed limit $D \rightarrow \infty$, the PDE-ODE system reduces to

$$U'_{0} = -\frac{1}{\tau}U_{0} - \frac{\rho}{m\tau} \sum_{j=1}^{m} \left[\kappa_{1,j}U_{0} - \kappa_{2,j}u_{j}^{1}\right],$$
$$u'_{j} = F_{j}(u_{j}) + \frac{1}{\tau} \left[\kappa_{1,j}U_{0} - \kappa_{2,j}u_{j}^{1}\right] e_{1}, \qquad j = 1, \dots, m,$$

where $\rho = m/|\Omega|$ is the "cell density" $\kappa_{1,j} \equiv 2\pi d_{1,j}$ and $\kappa_{2,j} \equiv 2\pi d_{2,j}$.

<u>Non-Identical Cells</u>: We take $\tau = 0.5$, and fix common permeability parameters $d_{1j} = 0.8$ and $d_{2j} = 0.2 \quad \forall j$. The intracellular kinetics F_j are not identical. Selkov parameters $\varepsilon_0 = 0.15$ and $\mu = 2$ are fixed for each cell, but α can vary from cell to cell. Isolated cells are not oscillatory.

Kuramoto order parameter: (measures the degree of oscillator phase synchrony):

$$R = \left\langle \left| N^{-1} \sum_{j=1}^{N} \exp[i\theta_j(t)] - \left\langle N^{-1} \sum_{j=1}^{N} \exp[i\theta_j(t)] \right\rangle \right| \right\rangle, \quad 0 \le R \le 1.$$

R = 1 (Perfect phase synchrony); R = 0 (No phase coherence);

Large Cell Populations: Synchronization II

Computations of order parameter R with respect to ρ . Iyaniwura (UBC)



Identical cells: $\alpha = 0.9$. "Defective" cells: α is random in $0.921 \le \alpha \le 0.952$.

Population density ρ plays a dual role of triggering and quenching oscillations

Interval of ρ where synchrony occurs decreases as the number of defective cells increases.

Cell-Bulk Model: Further Directions

Let D = O(1). Consider *m* "randomly" placed cells in a disk. Can we observe clusters of oscillating and non-oscillating cells? (i.e. "chimera"-type states.)



- How do we solve the spectral problem in arbitrary domains? (fast multipole methods for G and G_{λ})
- Numerics for the GCEP for large numbers of cells.
- What if the steady-state solution is not unique (hysteresis) or if intracellular dynamics has a time-delay?
- Intracellular dynamics to model a specific biological system (LuxIR circuit in Vibrio fischeri).
- Derive a RD system in the homogenized limit of $m \gg 1$ but $m\epsilon^2 \ll 1$.
- Two bulk-diffusing (autoinducer) species.
- PDE-ODE Model in 3-D. (interactions are, in general, much weaker owing to 1/r decay of Green's function).

PDE-ODE Cell-Bulk Model in 3-D

The dimensionless bulk concentration $U(\mathbf{x}, t)$ satisfies

$$\frac{\partial U}{\partial t} = \mathbf{D} \,\Delta U - \kappa \,U, \quad \mathbf{x} \in \Omega \setminus \bigcup_{j=1}^{m} \Omega_{\varepsilon_j} \,; \quad \partial_n \,U = 0, \quad \mathbf{x} \in \partial\Omega \,,$$
$$\varepsilon \mathbf{D} \,\partial_n U = \mathbf{d}_{1,j} \,U - \frac{\mathbf{d}_{2,j}}{\varepsilon} u_j^1 \,, \quad \mathbf{x} \in \partial\Omega_{\varepsilon_j} \,, \quad j = 1, \dots, m,$$

which is coupled to the dimensionless intracellular dynamics for the j^{th} cell

$$\frac{d\boldsymbol{u}_{j}}{dt} = \boldsymbol{F}_{j}\left(\boldsymbol{u}_{j}\right) + \boldsymbol{e}_{1} \int_{\partial\Omega_{\varepsilon_{j}}} \left(\frac{\boldsymbol{d}_{1,j}}{\varepsilon}U - \frac{\boldsymbol{d}_{2,j}}{\varepsilon^{2}}u_{j}^{1}\right) \, dS \,, \quad j = 1, 2, \dots, m,$$

where $u_j = (u_j^1, \dots, u_j^n)^T$, $e_1 \equiv (1, 0, \dots, 0)^T$, and $d_{2,j} = \mathcal{O}(1)$.

<u>Near Well-Mixed Limit</u>: An interesting limit where there is $\mathcal{O}(1)$ interaction between the cells is when

$$D = \mathcal{O}(\varepsilon^{-1}), \ \kappa = \mathcal{O}(1), \ d_{1,j} = \frac{\widetilde{d}_{1,j}}{\varepsilon}, \ \text{where} \ \widetilde{d}_{1,j} = \mathcal{O}(1)$$

In this regime, Quorum and Diffusing sensing can be studied through a common limiting system.

ODE System in Near Well-Mixed Limit

In this limit, the PDE-ODE system reduces to

$$U_0' = -\kappa U_0 + \frac{4\pi}{|\Omega|} \sum_{j=1}^m (p_{2,j} v_j^1 - p_{1,j} U_0) - \frac{16\pi^2 \varepsilon}{|\Omega|} \sum_{j=1}^m p_{1,j} (\mathcal{G} c)_j + \dots,$$

 $\frac{d\boldsymbol{v}_j}{dt} = \boldsymbol{F}_j(\boldsymbol{v}_j) + 4\pi \boldsymbol{e}_1(\boldsymbol{p}_{1,j}U_0 - \boldsymbol{p}_{2,j}v_j^1) + 16\varepsilon\pi^2 \boldsymbol{e}_1 \boldsymbol{p}_{1,j}(\mathcal{G}\boldsymbol{c})_j + \dots, \ j = 1,\dots,m,$

where $\boldsymbol{c} = (c_1, \ldots, c_m)^T$, $\boldsymbol{\mathcal{G}}$ is Neumann Green's matrix in 3-D and

$$p_{1,j} \equiv \frac{D_0 \,\widetilde{d}_{1,j}}{\widetilde{d}_{1,j} + D_0}, \quad p_{2,j} \equiv \frac{D_0 \,d_{2,j}}{\widetilde{d}_{1,j} + D_0}, \quad c_j \equiv \frac{d_{2,j} v_j^1 - \widetilde{d}_{1,j} U_0}{\widetilde{d}_{1,j} + D_0}, \quad j = 1, \dots, m.$$

■ For $D_0 \rightarrow 0$, then $p_{1,j} \rightarrow 0$ and $p_{2,j} \rightarrow 0$ (no cell-cell communication).

- For $D_0 \to \infty$ (well-mixed), then $p_{1,j} \to \tilde{d}_{i,j}$, $p_{2,j} \to d_{2,j}$, and $c_j \to 0$ (maximal cell-cell communication, but cell configuration insignificant).
- For $D_0 = O(1)$ dependence on cell configuration and shape of confining domain Ω is at O(ε) term through Neumann G-matrix G.
- ODE system: reveals both quorum sensing and diffusion sensing behavior.

Thanks For Your Attention!

Topic II: 1-D PDE-ODE Bulk-Cell Model

Mathematical Model:

- One compartment (cell) at each endpoint of the domain [0, 2L].
- N dynamically interacting substances within each cell, but only one substance can be secreted into the bulk 0 < x < 2L.
- The signaling substance diffuses and is degraded in the bulk.
- Distinct from "quasi-static" models where compartments yield nonlinear flux-type boundary conditions. (Glass et al, Othmer, Riecke).



Ref [GLNW]: J. Gou, Y. X. Li, W. Nagata, M. J. Ward, *Synchronized Oscillatory Dynamics for a 1-D Model of Membrane Kinetics Coupled by Linear Bulk Diffusion*, SIADS, **14**(4), (2015), pp. 2096–2137.

1-D Theory: General Model

The bulk diffusion field C(x, t) for the signalling molecule satisfies

$$\tau C_t = DC_{xx} - C, \qquad t > 0, \quad 0 < x < 2L,$$

$$DC_x(0,t) = G(C(0,t), u_1(t)), \qquad -DC_x(2L,t) = G(C(2L,t), v_1(t)).$$

Inside each compartment, there are N species that can interact, and that their dynamics are described by N-ODE's of

$$\frac{d\boldsymbol{u}}{dt} = \boldsymbol{\mathcal{F}}(\boldsymbol{u}) + \boldsymbol{\beta}\boldsymbol{\mathcal{P}}(C(0,t),u_1)\boldsymbol{e_1}, \qquad \frac{d\boldsymbol{v}}{dt} = \boldsymbol{\mathcal{F}}(\boldsymbol{v}) + \boldsymbol{\beta}\boldsymbol{\mathcal{P}}(C(2L,t),v_1)\boldsymbol{e_1}.$$

where $\boldsymbol{u} = (u_1, u_2, ..., u_N)^T$ and $\boldsymbol{e}_1 = (1, 0, ..., 0)^T$. Thus, only one component can diffuse into the bulk.

Special Case: Linear coupling is a special case

$$G(a,b) = \kappa_1(a-b), \qquad \mathcal{P}(a,b) = a-b.$$

<u>Conditional Oscillator</u>: When $\beta = 0$, we assume that the isolated ODE system has a linearly stable steady state. With coupling to the bulk the steady-state is modified, and can trigger oscillations through a HB.

Steady State and Linear Stability

Assuming identical compartments, the symmetric steady-state solution satisfies a nonlinear algebraic system

 $-C_e^0 \tanh(\omega_0 L) = \omega_0 G(C_e^0, u_{1e}), \quad \mathcal{F}(\boldsymbol{u}) + \beta \mathcal{P}(C(0, t), u_1) \boldsymbol{e_1} = 0.$

To study its linear stability, we introduce

$$C(x,t) = C_e(x) + e^{\lambda t} \eta(x), \quad \boldsymbol{u} = \boldsymbol{u}_e + e^{\lambda t} \boldsymbol{\phi}.$$

Upon linearizing, we obtain a Steklov-type spectral problem for ϕ and $\eta(x)$ on 0 < x < L:

$$D\eta_{xx} - (1 + \tau \lambda)\eta = 0, \qquad 0 < x < L; \qquad D\eta_x(0) = G_c^e \eta_0 + G_{u_1}^e \phi_1,$$
$$J_e \phi + \beta (\mathcal{P}_c^e \eta_0 + \mathcal{P}_{u_1}^e \phi_1) e_1 = \lambda \phi.$$

For the boundary condition of $\eta(x)$ at the midline x = L, we have two possibilities:

$\phi_x(L) = 0 ,$	Even: In-Phase Synchronization
$\phi(L) = 0,$	Odd: Anti-Phase Synchronization

Linear Stability Analysis

For both cases, the eigenvalue λ are roots of $\mathcal{G}(\lambda) = 0$, where

$$\mathcal{G}(\lambda) = 1 - p_{\pm}(\lambda) \frac{M_{11}(\lambda)}{\det \left(J_e - \lambda I\right)} \,,$$

- J_e = $(\frac{\partial F_i}{\partial u_j})_{ij}$ is the Jacobian matrix of the uncoupled ODE system evaluated at the new steady-state
- \blacksquare M_{11} is the cofactor of the element $a_{1,1}$ of the matrix $J_e \lambda I$.
- $\mathbf{P}_{\pm}(\lambda)$, determined by the bulk diffusion field, is

$$p_{+}(\lambda) \equiv \beta \left(\frac{G_{u_{1}}^{e} \mathcal{P}_{c}^{e} - \mathcal{P}_{u_{1}}^{e} G_{c}^{e} - \mathcal{P}_{u_{1}}^{e} D\Omega_{\lambda} \tanh(\Omega_{\lambda}L)}{G_{c}^{e} + D\Omega_{\lambda} \tanh(\Omega_{\lambda}L)} \right) , \quad \text{(In-phase)}$$
$$p_{-}(\lambda) \equiv \beta \left(\frac{G_{u_{1}}^{e} \mathcal{P}_{c}^{e} - \mathcal{P}_{u_{1}}^{e} G_{c}^{e} - \mathcal{P}_{u_{1}}^{e} D\Omega_{\lambda} \coth(\Omega_{\lambda}L)}{G_{c}^{e} + D\Omega_{\lambda} \coth(\Omega_{\lambda}L)} \right) , \quad \text{(Anti-phase)},$$

where we take the principal value of $\varphi_{\lambda} = \sqrt{\frac{1+\tau\lambda}{D}}$.

Theoretical Framework for Analysis

Linearized Analysis:

- Find HB points for in-phase and anti-phase modes.
- Use winding number criterion of complex analysis for information on linear stability, to get phase diagrams.
- Image Rigorous spectral results for one-ODE and $L \to \infty$.

Global Bifurcation Analysis: Track global branches of in-phase and anti-phase periodic solutions branches emanating from HB points. Method of lines for Bulk Diffusion and XPPAUT. Identify secondary bifurcations such as Hopf-Hopf points, Torus bifurcations, etc.

Full Numerical Simulations of the PDE-ODE to verify bifurcation studies.

Weakly Nonlinear Analysis:

- Determine whether HB points are sub or supercritical.
- Sey Challenge: Derive amplitude equations with Steklov structure.

Selkov Compartmental Dynamics: I

Suppose u(t) = (V(t), W(t)), and that there is linear coupling $G = \kappa(V(t) - C(0, t))$. Choose Selkov membrane dynamics

$$\frac{dV}{dt} = f(V,W) + \beta(C(0,t) - V), \qquad f(V,W) \equiv \alpha W + WV^2 - V,$$
$$\frac{dW}{dt} = g(V,W) = \epsilon_0(\mu - (\alpha W + WV^2)).$$

We fix the Selkov parameters $\mu = 2$, $\alpha = 0.9$ and $\epsilon_0 = 0.15$.

Linear stability phase diagram in *D* vs β plane for $\kappa = k = L = 1$.



Caption: In-phase and anti-phase oscillations occur within the open loops bounded by the blue solid and red dashed curves. Above the faint-hashed curve, the in-phase periodic solution is stable; below the magenta dot-dashed curve the anti-phase periodic solution is stable.

Selkov Compartmental Dynamics: II



Full numerics for C(x,t) confirming the theory at the blue and red dots in the phase diagram. Left: (blue dot) is in-phase. Right: (red dot) anti-phase.





Full numerics for D = 0.4 and $\beta = 0.8$, showing different longtime results, either in-phase or anti-phase, depending on the initial conditions. Parameter values are within both in-phase and antiphase loops.

Selkov Compartmental Dynamics: III

Global Bifurcation Study: slices through linear stability phase diagram



Caption: Bifurcation diagram of *V* for slices through linear stability phase diagram. Left: *V* versus *D* for $\beta = 0.8$ (vertical slice). Right: *V* versus β for D = 0.4 (horizontal slice). The solid (dashed) lines are linearly stable (unstable) branches of steady-states. Closed loops are branches of in-phase and anti-phase periodic solutions, with solid (open) circles indicating stable (unstable) periodic solutions. Left: bifurcating branch near D = 1 is the in-phase synchronous branch. Right: the outer loop is the anti-phase branch. Torus bifurcations occur where the periodic solution branches lose stability.

Modeling and References

Biological "Realistic" Models:

- Simplified version of the GnRH neuron hormone model from (Li-Khadra, 2008) where C(x,t) is the GnRH concentration in the bulk medium.
- Cell-cell signaling in Dictyostelium (Goldbeter 1990), where C(x,t) is the concentration of the cAMP in the bulk region, and u is the total fraction of cAMP receptor in the active state on the two membranes.

Ref [GLNW]: J. Gou, Y. X. Li, W. Nagata, M. J. Ward, *Synchronized Oscillatory Dynamics for a 1-D Model of Membrane Kinetics Coupled by Linear Bulk Diffusion*, SIADS, **14**(4), (2015), pp. 2096–2137.

Ref [G]: J. Gou et al. A Theory of Synchrony by Coupling Through a Diffusive Chemical Signal, Physica D, **339**, (2017), pp. 1–17.

Ref [GW]: J. Gou, M. J. Ward, Oscillatory Dynamics for a Coupled Membrane-Bulk Diffusion Model with Fitzhugh-Nagumo Kinetics, SIAP, **76**(2), (2016), pp. 776-804.

Topic III: Bulk-Surface RD Systems

Coupling passive diffusion in a bulk domain with a reaction-diffusion process on the domain boundary through a Robin boundary condition.



- H. Levine Membrane Bound Turing Patterns, Phys. Rev. E. (2005). Turing patterns occur even with equal diffusivities.
- A. Madzvamuse et al, Proc. Roy. Soc. A, (2015). General Turing stability analysis of spatially uniform state.
- Modeling: A. Madzvamuse et al, A coupled bulk-surface model for cell polarization, J. Theo. Bio. (2019). Giese, Frey et al: various models of protein pattern formation.

<u>Our Focus</u>: For a class of coupled bulk-surface RD model in a disk, develop a weakly nonlinear theory for pattern formation near bifurcation points. Derive and analyze amplitude equations for Hopf, Turing, Turing-Hopf instabilities. Thesis work of Paquin-Lefebvre (UBC)

Coupled Membrane-Bulk System I

Dimensionless Formulation: Let $\Omega = {\mathbf{x} \in \mathbb{R}^2 \mid ||\mathbf{x}| < R}$. In the bulk, we assume passive diffusion

$$\frac{\partial U}{\partial t} = D_u \Delta U - \sigma_u U \,, \quad \frac{\partial V}{\partial t} = D_v \Delta V - \sigma_v V \,, \quad \mathbf{x} \in \Omega \,, \quad t > 0 \,,$$

coupled to the surface with the Robin boundary condition

$$D_u \left. \frac{\partial U}{\partial r} \right|_{r=R} = K_u \left(u - U |_{r=R} \right) , \qquad D_v \left. \frac{\partial V}{\partial r} \right|_{r=R} = K_v \left(v - V |_{r=R} \right) .$$

This 2-D bulk problem is coupled to a nonlinear 1-D RD system on the boundary (membrane) of the circular disk

$$\frac{\partial u}{\partial t} = \frac{d_u}{R^2} \frac{\partial^2 u}{\partial \theta^2} - K_u \left(u - U |_{r=R} \right) + f(u, v),$$

$$\frac{\partial v}{\partial t} = \frac{d_v}{R^2} \frac{\partial^2 v}{\partial \theta^2} - K_v \left(v - V |_{r=R} \right) + g(u, v).$$

Ref: [PNW] F. Paquin-Lefebvre, W. Nagata, M. J. Ward, *Pattern Formation and Oscillatory Dynamics in a 2-D Coupled Bulk-Surface Reaction Diffusion System*, to appear, SIADS, (2019), (48 pages).

Coupled Membrane-Bulk System II

Outline of Analysis:

- Construct radially symmetric steady-state solution $\mathbf{u}_e(r)$, and linearize introducing $\mathbf{u} = \mathbf{u}_e + e^{\lambda t + in\theta} \Phi$. Derive the eigenvalue relation. Linearization is not around a spatially uniform state.
- Plot marginal stability curves for Hopf n = 0 and Turing n = 1 branches. The bifurcation parameters are taken as D_v and K_v , and we consider any such two-parameter path crossing a marginal stability boundary.
- Key step: formulate appropriate adjoint of linearized operator, inner product, orthogonality relation, and solvability condition.
- Multi-scale expansion in order to derive normal form amplitude equations for Hopf, Turing, and Turing-Hopf instabilities. Central is to derive explicit "computable" formulae for the coefficients for arbitrary *f* and *g*.

Membrane-Bulk: WNA (Technical I)

$$\dot{W} = \mathbf{F}(W) = \begin{pmatrix} D_u \Delta U - \sigma_u U \\ D_v \Delta V - \sigma_v V \\ \frac{d_u}{R^2} u_{\theta\theta} - K_u (u - U) + f(u, v) \\ \frac{d_v}{R^2} v_{\theta\theta} - K_v (v - V) + g(u, v) \end{pmatrix}$$

for functions satisfying

$$W \in \mathcal{W} \equiv \left\{ \begin{pmatrix} U(r,\theta) \\ V(r,\theta) \\ u(\theta) \\ v(\theta) \end{pmatrix} \middle| \begin{array}{l} D_u \partial_r U|_{r=R} = K_u \left(u - U|_{r=R} \right) \\ D_v \partial_r V|_{r=R} = K_v \left(v - V|_{r=R} \right) \end{array} \right\}$$

For a radially symm. base-state $W_e(r) \in \mathcal{W}$, let $\tilde{W} = W - W_e$. Expand

$$\tilde{\tilde{W}} = \mathcal{L}\tilde{W} + \mathcal{B}(\tilde{W}, \tilde{W}) + \mathcal{C}(\tilde{W}, \tilde{W}, \tilde{W}) + \dots,$$

where $\mathcal{L}\tilde{W}$ is the linearized operator with eigenfunctions $\mathcal{L}\Phi_n = \lambda \Phi_n$,

$$\mathcal{L}(\tilde{W}) = \begin{pmatrix} D_u \Delta \tilde{U} - \sigma_u \tilde{U} \\ D_v \Delta \tilde{V} - \sigma_v \tilde{V} \\ \frac{d_u}{R^2} \tilde{u}_{\theta\theta} - K_u \left(\tilde{u} - \tilde{U}\right) + f_u^e \tilde{u} + f_v^e \tilde{v} \\ \frac{d_v}{R^2} \tilde{v}_{\theta\theta} - K_v \left(\tilde{v} - \tilde{V}\right) + g_u^e \tilde{u} + g_v^e \tilde{v} \end{pmatrix}, \quad \text{where} \quad \Phi_n = \begin{pmatrix} \tilde{W}_1(r) e_1^T \phi_n \\ \tilde{W}_2(r) e_2^T \phi_n \\ \phi_n \end{pmatrix} e^{in\theta}.$$

Membrane-Bulk: WNA (Technical II)

Stability Threshold: Re($\lambda_{max}(n, \mu_0)$) = 0 for n = 0, 1, 2, ... and $\mu_0 \equiv (K_v, D_v)^T$. Derive the adjoint $\mathcal{L}^*(W^*)$ and introduce inner product

$$\langle W^{\star}, W \rangle = \int_{0}^{2\pi} \int_{0}^{R} \left[\overline{U^{\star}}U + \overline{V^{\star}}V \right] r dr d\theta + \int_{\partial\Omega} \left[\overline{u^{\star}}u + \overline{v^{\star}}v \right] d\sigma \,,$$

where $W \equiv (U, V, u, v)^T$ and $W^* \equiv (U^*, V^*, u^*, v^*)^T$.

Fredholm Alternative Lemma: Let λ_c denote the critical eigenvalue at a given bifurcation point $\mu_0 = (K_v^c, D_v^c)$. Then,

$$\mathcal{L}(\mu_0; \mathbf{\Phi}_n) = \lambda_c \mathbf{\Phi}_n, \quad \mathcal{L}^{\star}(\mu_0; \mathbf{\Phi}_n^{\star}) = \overline{\lambda_c} \mathbf{\Phi}_n^{\star}, \quad \lambda_c \equiv \begin{cases} i\lambda_I & n = 0\\ 0 & n \neq 0 \end{cases}$$

Consider the inhomogeneous problem

$$\lambda_c X - \mathcal{L}(\mu_0; X) = \mathcal{F} \quad \text{with} \quad \left[\left. \partial_r \begin{pmatrix} D_u x_1 \\ D_v x_2 \end{pmatrix} \right|_{r=R} - \begin{pmatrix} K_u (x_3 - x_1|_{r=R}) \\ K_v (x_4 - x_2|_{r=R}) \end{pmatrix} \right] = \begin{pmatrix} \xi(\theta) \\ \eta(\theta) \end{pmatrix},$$

where $X \equiv (x_1(r, \theta), x_2(r, \theta), x_3(\theta), x_4(\theta))^T$. A necessary condition for a solution is

$$\langle \mathbf{\Phi}_n^{\star}, \mathcal{F} \rangle + \int_{\partial \Omega} \overline{U_n^{\star}} \xi \, d\sigma + \int_{\partial \Omega} \overline{V_n^{\star}} \eta \, d\sigma = 0 \, .$$

Shanghai – p.45

Membrane-Bulk: Normal Forms

Two-parameter sweep across linear stability boundary $(K_v^c, D_v^c)^T$:

$$\mu = (K_v^c, D_v^c)^T + \varepsilon^2 \mu_1 \,.$$

The weakly nonlinear theory (WNA) yields Normal Forms:

$$\begin{aligned} \frac{dA_0}{d\tau} &= g_{1000}^T \mu_1 A_0 + g_{2100} |A_0|^2 A_0 \,, \quad \text{(Hopf)} \,, \\ \frac{dA_n}{d\tau} &= g_{0010}^T \mu_1 A_n + g_{0021} |A_n|^2 A_n \,, \quad \text{(Pitchfork)} \,. \end{aligned}$$

and the codimension-two Turing-Hopf:

$$\frac{dA_0}{d\tau} = g_{1000}^T \mu_1 A_0 + g_{2100} |A_0|^2 A_0 + g_{1011} |A_n|^2 A_0 ,$$

$$\frac{dA_n}{d\tau} = g_{0010}^T \mu_1 A_n + g_{0021} |A_n|^2 A_n + g_{1110} |A_0|^2 A_n .$$

WNA provides explicit formulae for all the coefficients in the normal form. Example: Consider Brusselator membrane kinetics

$$f(u,v) = a - (b+1)u + u^2v \,, \quad g(u,v) = bu - u^2v \,; \quad a > 0 \,, \quad 0 < b < a^2 + 1 \,.$$

Membrane-Bulk: Linear Stability, WNA



Caption: Linear stability phase diagram in (K_v, D_v) plane with R = 1, $D_u = 1$ $\sigma_u = \sigma_v = 0.01$, $K_u = 0.1$, $d_u = d_v = 0.5$, a = 3 and b = 7.5 (left) and b = 8.7 (right). Right: "o" indicates supercritical and "+" indicates subcritical.



Caption: Transition from a super to sub-critical Hopf bifurcation. Plot of the normal form coefficient $\text{Re}(g_{2100})$ along Hopf stability curve for b = 8.7.

Membrane-Bulk: Supercritical Hopf



Caption: Supercritical Hopf: b = 7.5 and $K_v = 5$. Left: global periodic solution branches (AUTO). Right: local branching: weakly nonlinear vs. AUTO



Caption: Left: parameter path $\mu = (K_v, D_v) = (5, 2.32)^T + \varepsilon^2 (0, 1)^T$, $\varepsilon = 0.1$. Right: weakly nonlinear vs. full numerics for membrane oscillations.

Membrane-Bulk: Subcritical Hopf



Caption: Subcritical Hopf: b = 8.7 and $D_v = 9$. Left: global periodic solution branches (AUTO). Right: local branching: weakly nonlinear vs. AUTO



Caption: Left: parameter path with $D_v = 9$ and $\varepsilon = 0.1$. Middle: U(r, t) for bulk. Right: relaxation oscillations for membrane oscillations.

Membrane-Bulk: Subcritical Pitchfork



Left: parameter path with $D_v = 5$, b = 7.5, and $\varepsilon = 0.075$. Right: WNA approx (solid), stable branches by time-stepping to steady state in full numerics.



Stable membrane pattern at t = 1000 (red curve) as evolved from the unstable branch near a subcritical pitchfork.

Membrane-Bulk: Supercritical Pitchfork



Same parameters except now b = 5 and $d_u = d_v = 1.0$. Pitchfork is now supercritical



Left: bifurcation diagram (AUTO) versus WNA. Right: stable Turing pattern.

Further Directions

- I-D Periodic Chains: "active units" on a ring coupled by bulk diffusion. The synchronous mode has the best stability properties.
- I-D Non-Identical Compartments: Eigenfunctions no longer of in-phase and anti-phase type. New behavior: oscillations can exist in only one compartment, with the other being essentially quiescent.
- I-D Theory: Numerical global bifurcation study. Follow Torus (secondary) bifurcation branches.
- Membrane-bulk RD systems: coupling 3-D passive bulk diffusion to an RD process on the surface (modeling, linear and weakly nonlinear stability theory).
- Localized membrane-bound RD patterns obtained from coupling to bulk-diffusion. Thesis work of Daniel Gomez (UBC). (Movie)

Thanks For Your Attention!