

# Reduction and Diffusion-Driven Instability Analysis In a Receptor-Based Model for Hydra Morphogenesis



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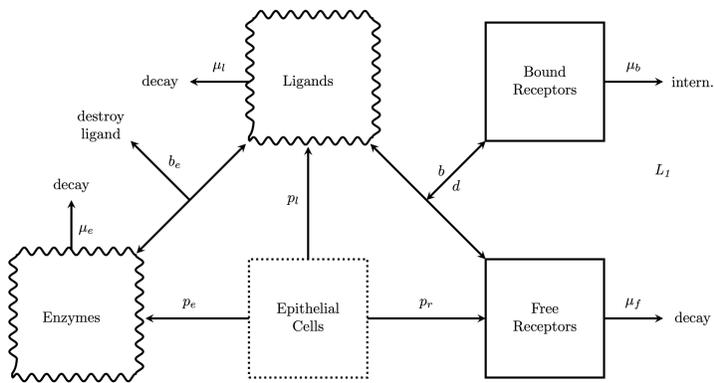
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## Abstract

How do complex structures emerge from an initially symmetric state? One mathematical approach, proposed by Alan Turing, involves studying reaction-diffusion equations. When given the right conditions, almost symmetrical initial datum to these equations can evolve into stable, spatially heterogeneous solutions known as Turing patterns. Such reaction-diffusion equations can be coupled to standard ODEs to model localized phenomena, opening the door to various new observable patterns - which are not necessarily Turing patterns - and a necessary condition for them to appear is diffusion-driven instability (DDI). We begin with a receptor-based model for Hydra morphogenesis [3], which we reduce by a quasi-static approximation before investigating the phenomenon of DDI in this new toy model.

## Original model

**Introduction:** This model describes the time evolution of four chemical components (two diffusive, two non-diffusive) according to a classical reaction scheme. It originates from [3] and suggests a simple dynamics for limb regeneration in Hydra and admits a visual representation like follows:



## Model Equations:

$$\begin{aligned} \partial_t r_f &= -\mu_f r_f + m_1 \frac{r_b}{1+r_b} - br_f \ell + dr_b & \text{in } \Omega \times (0, T), & r_f(\cdot, 0) = r_f^0 \text{ in } \Omega, \\ \partial_t r_b &= -\mu_b r_b + br_f \ell - dr_b & \text{in } \Omega \times (0, T), & r_b(\cdot, 0) = r_b^0 \text{ in } \Omega, \\ \partial_t \ell &= d_1 \Delta \ell - \mu_l \ell + m_2 \frac{r_b}{1+r_b} - br_f \ell + dr_b - b_e \ell e & \text{in } \Omega \times (0, T), & \ell(\cdot, 0) = \ell^0 \text{ in } \Omega, \\ \partial_t e &= d_2 \Delta e - \mu_e e + m_3 \frac{r_b}{1+r_b} & \text{in } \Omega \times (0, T), & e(\cdot, 0) = e^0 \text{ in } \Omega, \\ \partial_\nu \ell &= 0, \quad \partial_\nu e = 0 & \text{on } \partial\Omega \times (0, T). \end{aligned}$$

**Domain:**  $\Omega = (0, L)$ , where  $L \in \mathbb{R}_{>0}$ . We assume  $L = 1$  except stated otherwise.

## Model reduction: quasi-static approximation and rescaling

### Idea of the approximation:

- Goal is to simplify the analysis of chemical systems kinetics by assuming that some species are reaching their steady-state concentrations (much) faster than others.
- In practice, introduce a second time scale in the system  $\tau = \varepsilon t$  and let  $\varepsilon \rightarrow 0$
- Results in a differential-algebraic equation where, in most cases, one can explicitly solve the algebraic equation (otherwise, restriction of the differential equation to an algebraic manifold)

### Singular limit:

Introduce  $0 < \varepsilon \ll 1$ , small parameter in the equation on  $\partial_t r_b$ . Let  $\varepsilon \rightarrow 0$  and solve for  $r_b$ :

$$r_b(t, x) = \frac{b}{d + \mu_b} r_f(t, x) \ell(t, x), \quad (t, x) \in \Omega \times (0, T).$$

### Change of variable and parameter rescaling:

$$\text{new functions: } u = \sqrt{\frac{b}{d + \mu_b}} r_f, \quad v = \sqrt{\frac{b}{d + \mu_b}} \ell, \quad w = \sqrt{\frac{b}{d + \mu_b}} e$$

$$\text{new parameters: } \tilde{m}_1 = \sqrt{\frac{b}{d + \mu_b}} m_1, \quad \tilde{m}_2 = \sqrt{\frac{b}{d + \mu_b}} m_2, \quad \tilde{m}_3 = b_e m_3, \quad \tilde{\mu}_b = \sqrt{\frac{b}{d + \mu_b}} \mu_b$$

### Reduced model equations:

$$\begin{aligned} \partial_t u &= -\mu_f u + \tilde{m}_1 \frac{uv}{1+uv} - \tilde{\mu}_b uv & \text{in } \Omega \times (0, T), & u(\cdot, 0) = u^0 \text{ in } \Omega, \\ \partial_t v &= d_1 \Delta v - \mu_l v + \tilde{m}_2 \frac{uv}{1+uv} - \tilde{\mu}_b uv - v w & \text{in } \Omega \times (0, T), & v(\cdot, 0) = v^0 \text{ in } \Omega, \\ \partial_t w &= d_2 \Delta w - \mu_e w + \tilde{m}_3 \frac{uv}{1+uv} & \text{in } \Omega \times (0, T), & w(\cdot, 0) = w^0 \text{ in } \Omega, \\ \partial_\nu v &= 0, \quad \partial_\nu w = 0 & \text{on } \partial\Omega \times (0, T). \end{aligned}$$

**Notation:** We drop the tilde  $\tilde{\square}$  notation on parameters in the future. Moreover, we denote respectively by  $f, g, h$ , each reaction term for  $u, v, w$  and define the vector field  $\phi = (f, g, h)$ . The quantity  $\mathbf{X}$  denotes the vector  $(u, v, w)$ .

**Theorem:** Let  $C_u, C_v, C_w$  be positive constants satisfying

$$C_u \geq \frac{m_1}{\min(\mu_f, \mu_b)}, \quad C_v \geq \frac{m_2}{\min(\mu_l, \mu_b, 1)}, \quad C_w \geq \frac{m_3}{\mu_e},$$

then the region  $\Sigma := [0, C_u] \times [0, C_v] \times [0, C_w]$  is an invariant rectangle of the system, providing uniform global estimates on  $u(t, x), v(t, x), w(t, x)$  for any  $\mathbf{X}^0 \in \Sigma$  and all  $t > 0, x \in \Omega$ .

## Diffusion-driven instability and Turing patterns

We are interested in pattern formation linked to destabilization of a spatially homogeneous state.

### Diffusion-driven instability (DDI):

- DDI is a bifurcation leading to a loss of stability of a constant steady state.
- The steady state is asymptotically stable as a solution of the kinetic system, i.e.  $d_1 = d_2 = 0$ .
- The steady state is unstable in the system with diffusion.

### Turing Patterns:

- Turing patterns are continuous patterns which arise due to DDI of a constant steady state.
- They are formed around the constant steady state (close-to-equilibrium patterns).

### Far-from-equilibrium patterns:

- This is another type of patterns which exhibit singularities.
  - Patterns with jump-discontinuity (arise in systems with multiple constant stationary solutions).
  - Spike patterns (arise e.g. due to diffusion-driven unbounded growth). [1]

## Steady states & a necessary and sufficient condition for DDI

### Steady states:

- There exists (up to) four steady states given by the relation

$$\bar{\mathbf{X}}_0 = (0, 0, 0), \quad \bar{\mathbf{X}} = \left( \frac{m_1}{\mu_f + \mu_b V} - \frac{1}{V}, \quad V, \quad \frac{m_3}{\mu_e} \left( 1 - \frac{\mu_f + \mu_b V}{m_1 V} \right) \right),$$

where  $V$  is one of the (up to) three roots of a polynomial of degree three such that  $V > \mu_f / (m_1 - \mu_b)$ .

- Multiple cases are identified for the sign and amount of roots ( $V$ ) of this polynomial:

$$(-), \quad (+), \quad (-, -), \quad (-, +), \quad (+, +), \quad (-, -, -), \quad (-, -, +), \quad (-, +, +), \quad (+, +, +)$$

- We show there is only one scenario occurring:  $(-, +, +)$ . This yields three nonnegative steady states:  $\bar{\mathbf{X}}_0, \bar{\mathbf{X}}_u, \bar{\mathbf{X}}$ , of type (stable, unstable, stable), which is ideal for observing jump-discontinuity patterns.

### DDI in the model:

- $\bar{\mathbf{X}}_0$ , cannot exhibit DDI since it is unconditionally asymptotically stable.
- $\bar{\mathbf{X}}_u$  cannot exhibit DDI since it is unstable.
- $\bar{\mathbf{X}}$  can exhibit DDI (see theorem below), and we identify two cases:
  - $d_1 \geq 0$  sufficiently small and  $d_2 > 0$ , fixed.
  - $d_1 > 0$ , fixed and  $d_2 \geq 0$ , sufficiently large (either with conditions or unconditionally if  $L$  large)

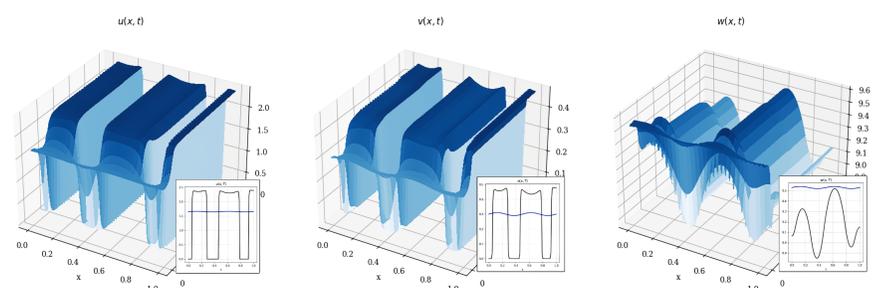
**Theorem:** Let  $\bar{\mathbf{X}}$  be a constant, non-zero, positive steady state of the system. Then,  $\bar{\mathbf{X}}$  can exhibit DDI if and only if

$$\det \left[ \nabla_{u,v} \begin{pmatrix} f \\ g \end{pmatrix} (\bar{\mathbf{X}}) \right] < 0$$

- For a fixed domain  $\Omega = (0, L)$ , the smallness condition on  $d_1$  is given by

$$d_1 < \frac{L^2}{\pi^2 \nabla_{u,v} f(\bar{\mathbf{X}})} \det \left[ \nabla_{u,v} \begin{pmatrix} f \\ g \end{pmatrix} (\bar{\mathbf{X}}) \right].$$

### Simulations:



### Parameters:

$$\mu_f = 0.87, \quad \mu_b = 0.68, \quad \mu_l = 0.05, \quad \mu_e = 0.60, \quad m_1 = 5.36, \quad m_2 = 9.68, \quad m_3 = 17.27, \\ d_1 = 0.0001, \quad d_2 = 0.1.$$

**For culture:** A well-known condition for DDI is  $\nabla_{u,v} f(\bar{\mathbf{X}}) > 0$ , also called *autocatalysis condition*. Unfortunately, all the patterns induced by that condition are unstable and thus impossible to observe in practice. Such case does not occur in our model.

## Conclusion and open problems

- We have precise conditions for DDI but DDI is not sufficient for Turing patterns.
- So far we only proved existence of Turing patterns numerically with simulations
  - How do we recognize Turing patterns and distinguish them from other patterns in simulations?
  - How can we prove the existence of Turing patterns analytically?

## References

- [1] Steffen Harting and Anna Marciniak-Czochra. Spike patterns in a reaction-diffusion-ode model with Turing instability. *arXiv*, March 2013.
- [2] Chris Kowall, Anna Marciniak-Czochra, and Finn Munnich. Stability results for bounded stationary solutions of reaction-diffusion-ODE systems, 2022.
- [3] Anna Marciniak-Czochra. *Developmental models with cell surface receptor densities defining morphological position*. PhD thesis, Universitat Heidelberg, 2004.

**Acknowledgements** This project has been funded by the Ministry of Science, Research and Arts Baden-Wurttemberg, grant no: BW6-07.