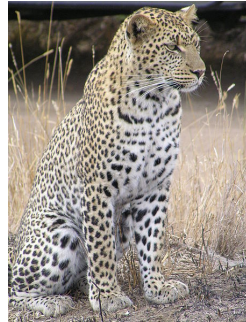


Lecture 4 – Spatially extended systems

February 18, 2020

- 1 Turing instability revisited.
- 2 Turing patterns on symmetric networks.
- 3 The theory of Turing patterns on directed networks.
- 4 Stochastic patterns for reaction-diffusion systems.

Self-organized patterns are ubiquitous in nature



The Belousov-Zhabotinsky reaction.



Highlighting the peculiarities:

- First system to display self-organization
- Regular oscillations between homogeneous states.

Experimental evidence

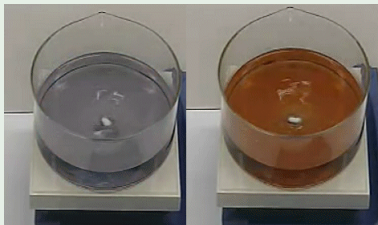
The Belousov-Zhabotinsky reaction.



Highlighting the peculiarities:

- First system to display self-organization
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Experimental evidence



Belousov e Zhabotinsky
Med. Publ. Moscow (1959) - *Biofizika* (1964)

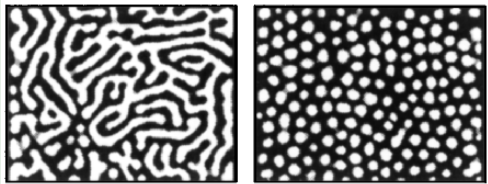
The Belousov-Zhabotinsky reaction.



Highlighting the peculiarities:

- First system to display self-organization
- Regular oscillations between homogeneous states.

Experimental evidence



Spatially organized patterns develop (**Turing instability**) - Vanag e Epstein
Phys. Rev. Lett. (2001)

Alan Turing (1912-1954)



- 1 **Turing machine**, a general purpose computer: concepts of algorithm and computation with the Turing machine.
- 2 Pivotal role in **cracking** intercepted coded messages during II world war.
- 3 Mathematical biology: chemical basis of **morphogenesis**.

Morphogenesis

Morphogenesis is the biological process that causes an organism to develop its **shape**.

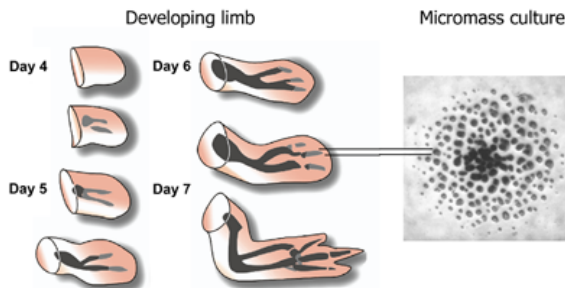
Morphogenesis addresses the problem of biological form at many levels, from the structure of **individual cells**, through the formation of **multicellular arrays and tissues**, to the higher order assembly of tissues into **organs** and whole **organisms**.



Morphogenesis

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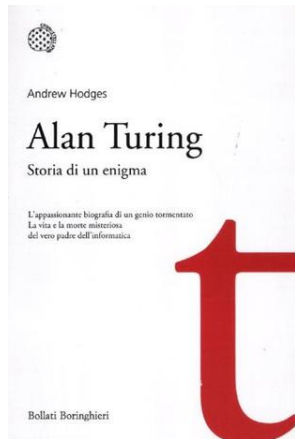
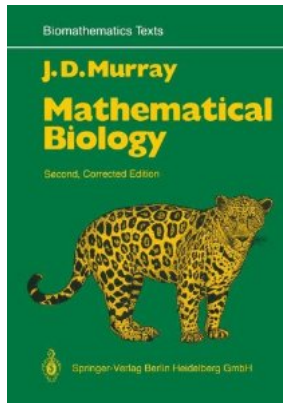


The complete and fine detailed understanding of the mechanisms involved in actual organisms required the **discovery of DNA** and the development of **molecular biology** and **biochemistry**.

Although the mechanism must be **genetically controlled**, the genes themselves cannot create the patterns. They only provide a blue print or recipe, for the pattern generation.

Turing suggested that under certain conditions, chemicals can **react** and **diffuse** in such a way as to produce **steady state heterogeneous spatial patterns** of chemical or morphogen concentration.

Two books



Deterministic reaction-diffusion systems

Alan Turing



The Turing instability (1952)

$$\begin{aligned}\partial_t \phi &= f(\phi, \psi) + D_\phi \nabla^2 \phi \\ \partial_t \psi &= g(\phi, \psi) + D_\psi \nabla^2 \psi\end{aligned}$$

where:

- 1 $\phi(r, t)$ and $\psi(r, t)$ are the species concentrations.
- 2 D_ϕ and D_ψ denote the diffusion coefficients

An informative albeit unrealistic image (Murray) (1)

- 1 Consider a **field of dry grass** with a large number of **grasshoppers** which can generate moisture by sweating if they get warm.
- 2 Suppose the grass is **set alight** at some point and the front starts to **propagate**.
- 3 When the grasshoppers get warm enough they can generate enough moisture to dampen the fire: when the flames will reach the pre-moistened area the **grass will not burn**.
- 4 The fire starts to spread. When the grasshoppers ahead of the flame front feel it coming they **move well ahead** of it ($D_G > D_F$).
- 5 The burned area is hence **restricted** to a given domain which depends on the parameters of the game.

An informative albeit unrealistic image (Murray) (2)

- 1 If instead of a initial single fire there was a **random scattering** of them, the process would result in a final **spatially inhomogeneous distribution** of burnt and preserved patches.
- 2 Notice that the **inhibitors** (grasshoppers) diffuse faster than the **activator** (fire).

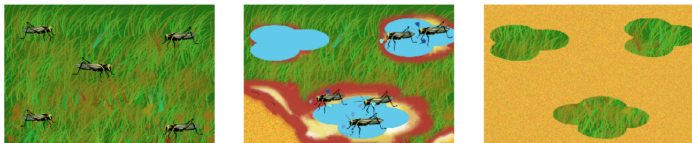


Figure 1.2: In a field with sweating grasshoppers, their combined efforts to prevent the fire from spreading would cause patches of grass in a charred area once the process has finished.

I. Stable fixed point of the aspatial model

Assume a **stable homogeneous fixed point** of the dynamics to exist and label it (ϕ^*, ψ^*) :

$$\begin{aligned}f(\phi^*, \psi^*) &= 0 \\g(\phi^*, \psi^*) &= 0\end{aligned}$$

The Jacobian matrix

$$J = \begin{pmatrix} f_\phi & f_\psi \\ g_\phi & g_\psi \end{pmatrix}$$

The **stability** of the fixed point implies $\text{Tr } J < 0$ and $\det J > 0$.

II. The perturbation.

Introduce a **small non homogeneous perturbation** of the fixed point:

$$w = \begin{pmatrix} \phi - \phi^* \\ \psi - \psi^* \end{pmatrix}.$$

and linearize the reaction-diffusion equations to get:

$$\dot{w} = Jw + D\nabla^2 w,$$

where

$$D = \begin{pmatrix} D_\phi & 0 \\ 0 & D_\psi \end{pmatrix}.$$

III. Laplacian's eigenfunctions

To solve the linearized system one introduces $W_k(x)$ such that:

$$\nabla^2 W_k(x) = -k^2 W_k(x),$$

Expand the perturbation w as

$$w(x, t) = \sum_{k \in \sigma} c_k e^{\lambda(k)t} W_k(x),$$

- 1 c_k refer to the initial condition.
- 2 Equivalent to **Fourier transforming** the original equation.
- 3 $\lambda(k)$ defines the dispersion relation

Substituting the ansatz in the linear system yields:

$$\lambda W_k = JW_k - k^2 DW_k$$

or equivalently:

$$\begin{pmatrix} f_\phi - D_\phi k^2 - \lambda & f_\psi \\ g_\phi & g_\psi - D_\psi k^2 - \lambda \end{pmatrix} W_k = 0$$

We require non trivial solutions for W_k which implies that λ is determined by the roots of the **characteristic polynomial**:

$$\det(\lambda(k)I - J - Dk^2) = 0$$

The **Turing instability** occurs if one can isolate a finite domain in k for which $\text{Re}(\lambda(k)) > 0$.

A simple calculation (done on the blackboard) yields the following general condition for the **Turing instability** to sets in:

$$\begin{aligned}(D_\phi g_\psi + D_\psi f_\phi)^2 &> 4D_\phi D_\psi (f_\phi g_\psi - f_\psi g_\phi) \\ (D_\phi g_\psi + D_\psi f_\phi) &> 0\end{aligned}$$

which sum up to the aforementioned conditions:

$$f_\phi + g_\psi < 0$$

$$f_\phi g_\psi - f_\psi g_\phi > 0$$

Important remarks

- 1 f_ϕ and g_ψ must be of **opposite** sign.
- 2 Assume $f_\phi > 0$ (**activator**) and $g_\psi < 0$ (**inhibitor**). Then, $f_\phi + g_\psi < 0$ implies:

$$f_\phi < |g_\psi|$$

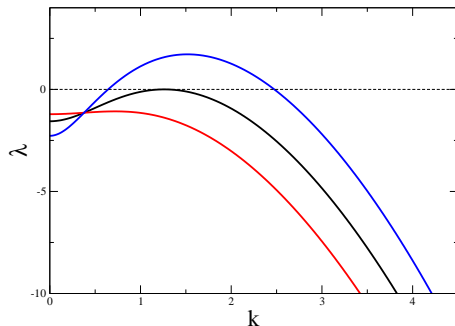
and thus:

$$\frac{D_\psi}{D_\phi} > \frac{|g_\psi|}{f_\phi} > 1$$

the inhibitor must diffuse **faster** than the activator.

- 3 Boundary conditions do matter.

The Brusselator model

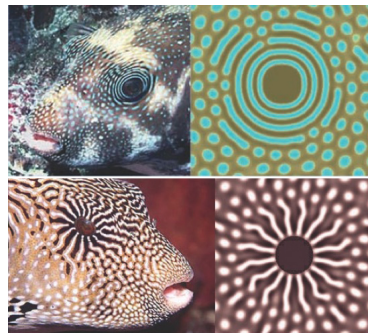


- 1 Species ϕ is the **activator**,
- 2 ψ play the role of the **inhibitor**.

$$\begin{aligned}f(\phi, \psi) &= a - (b + d)\phi + c\phi^2\psi \\g(\phi, \psi) &= b\phi - c\phi^2\psi\end{aligned}$$

From a **random perturbation** of the homogeneous fixed point to a **stationary pattern**.

Turing patterns are widespread



Patterns on a (symmetric) network.

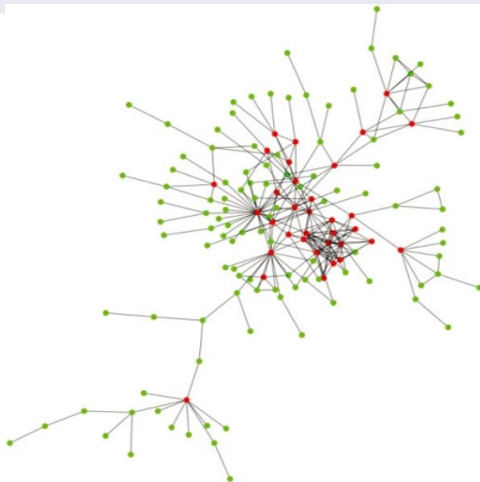
Adjacency matrix

$$\mathbf{W} = \begin{bmatrix} 1 & 0 & 0 & \dots \\ \dots & 0 & 1 & \dots \\ \vdots & \vdots & \dots & \vdots \\ \dots & 1 & 0 & \dots \end{bmatrix}$$

$W_{ij} = 1$, if nodes i and j
are connected ($i \neq j$), and
 $W_{ij} = 0$ otherwise

$$k_i = \sum_{j=1}^{\Omega} W_{ij} \text{ (node degree)}$$

Scale-free network



Patterns on a (symmetric) network.

Adjacency matrix

$$\mathbf{W} = \begin{bmatrix} 1 & 0 & 0 & \dots \\ \dots & 0 & 1 & \dots \\ \vdots & \vdots & \dots & \vdots \\ \dots & 1 & 0 & \dots \end{bmatrix}$$

$W_{ij} = 1$, if nodes i and j are connected ($i \neq j$), and $W_{ij} = 0$ otherwise

$$k_i = \sum_{j=1}^{\Omega} W_{ij} \text{ (node degree)}$$

Reaction-diffusion equations

$$\partial_t \phi_i = f(\phi_i, \psi_i) + D_{\phi} \sum_{j=1}^{\Omega} \Delta_{ij} \phi_j$$

$$\partial_t \psi_i = g(\phi_i, \psi_i) + D_{\psi} \sum_{j=1}^{\Omega} \Delta_{ij} \psi_j$$

where $i = 1, \dots, \Omega$ and $\Delta_{ij} = W_{ij} - k_i \delta_{ij}$ is the **discrete Laplacian operator**.

Linear stability analysis on networks

Perturbation near the homogeneous fixed point

$$\phi_i = \phi^* + \delta\phi_i \quad \psi_i = \psi^* + \delta\psi_i$$

- **Linearize** the equations for ϕ_i and ψ_i
- Introduce the **eigenvectors** of the Laplacian

$$\sum_j \Delta_{ij} \Phi_j^{(\alpha)} = \Lambda^{(\alpha)} \Phi_i^{(\alpha)}$$

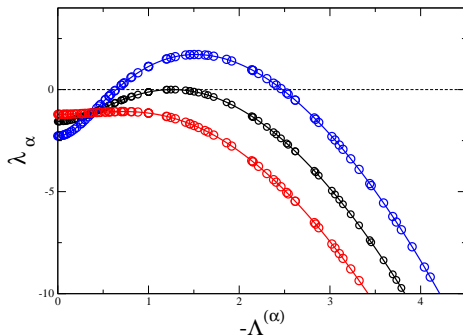
The **eigenvalues** $\Lambda^{(\alpha)}$ are **real** and **negative**.

The set of eigenvectors defines a **basis** on which we can expand the perturbation.

$$\delta\phi_i = \sum_{\alpha=1}^{\Omega} c_{\alpha} e^{\lambda_{\alpha}\tau} \Phi_i^{(\alpha)}$$
$$\delta\psi_i = \sum_{\alpha=1}^{\Omega} c_{\alpha} \beta_{\alpha} e^{\lambda_{\alpha}\tau} \Phi_i^{(\alpha)}$$

Inserting in the linearized equation one gets a **dispersion relation** for λ_{α} versus $\Lambda^{(\alpha)}$, which controls the **instability**.

Dispersion relation for the Brusselator model



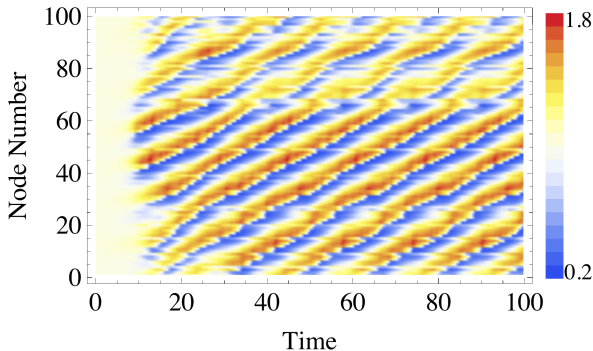
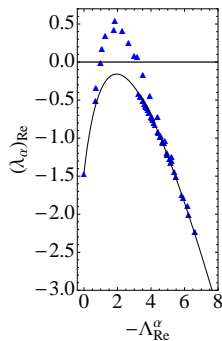
The dispersion relation is defined on the discrete support of Ω eigenvalues $\Lambda^{(\alpha)}$.

The curve relative to the continuum case is recovered by replacing $\Lambda^{(\alpha)}$ with $-k^2$.

A **segregation** in activator/inhibitors rich/poor nodes is found as follows the linear instability.

The active role of topology: **directed** networks

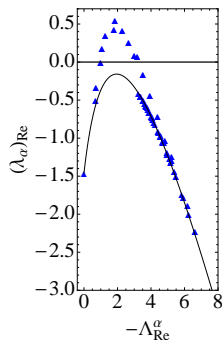
Self-organized **waves** can develop instigated by the network topology.



M. Asllani et al, Nature
Communications
(2014)

The active role of topology: **directed** networks

Self-organized **waves** can develop instigated by the network topology.



M. Asllani et al, Nature
Communications
(2014)

From the linear stability analysis...

$$\Lambda^{(\alpha)} = \Lambda_{\text{Re}}^{(\alpha)} + i\Lambda_{\text{Im}}^{(\alpha)}$$

The modified Jacobian matrix:

$$J_{\alpha} = J + D\Lambda^{(\alpha)}$$

yields the **dispersion relation**:

$$(\lambda_{\alpha})_{\text{Re}} = \frac{1}{2} [(\text{tr} J_{\alpha})_{\text{Re}} + \gamma]$$

where γ is a functions of J_{α}

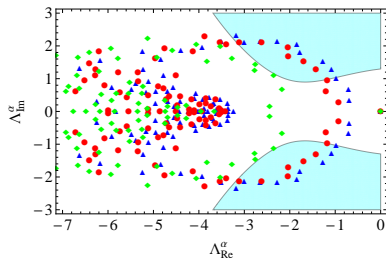
Region of instability

The instability develops when $(\lambda_{\alpha})_{\text{Re}}$ is **positive**, namely when:

$$S_2(\Lambda_{\text{Re}}^{(\alpha)}) [\Lambda_{\text{Im}}^{(\alpha)}]^2 \leq -S_1(\Lambda_{\text{Re}}^{(\alpha)}),$$

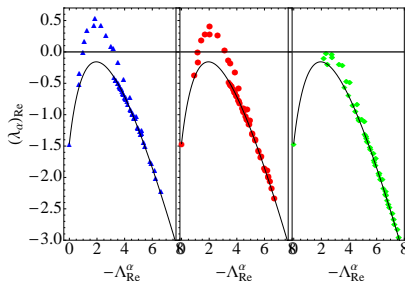
where S_1 and S_2 are polynomials of **fourth** and **second** degree in $\Lambda_{\text{Re}}^{(\alpha)}$

The active role of topology



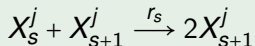
The instability region (shaded) in the $(\Lambda_{\text{Re}}^{(\alpha)}, \Lambda_{\text{Im}}^{(\alpha)})$ plane.

Dispersion relations for **Newman-Watts** networks with different p , the probability of long-range links

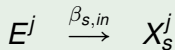
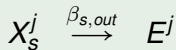


Example 1. Spatial autocatalytic model

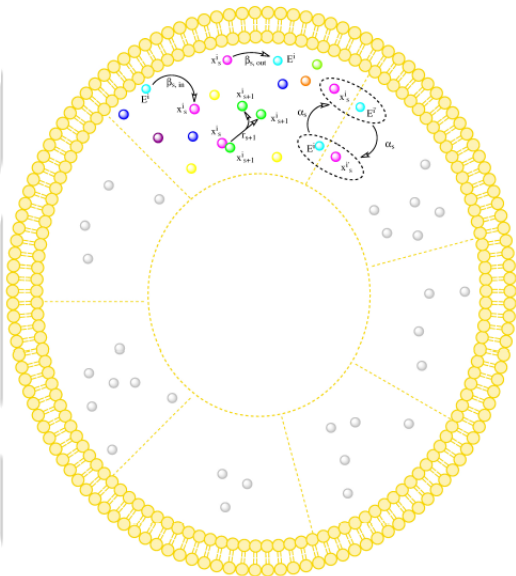
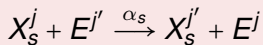
Autocatalytic reaction



Exchange with the bulk

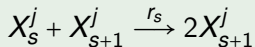


Migration between urns

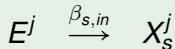
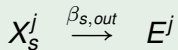


Example 2. Spatial autocatalytic model

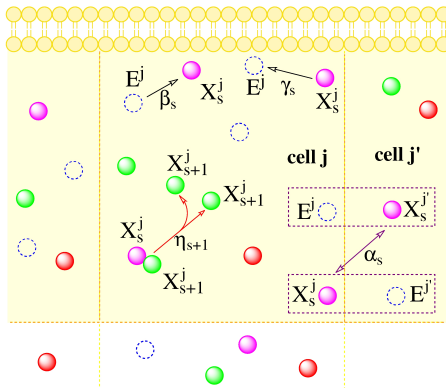
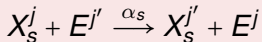
Autocatalytic reaction



Exchange with the bulk



Migration between urns



Transition associated to migration

$$\begin{aligned}T(n_s^j - 1, n_s^{j'} + 1 | n_s^j, n_s^{j'}) &= \frac{\alpha_s}{z\Omega} \frac{n_s^j}{N} \left(1 - \sum_{m=1}^k \frac{n_m^{j'}}{N}\right), \\T(n_s^j + 1, n_s^{j'} - 1 | n_s^j, n_s^{j'}) &= \frac{\alpha_s}{z\Omega} \frac{n_s^{j'}}{N} \left(1 - \sum_{m=1}^k \frac{n_m^j}{N}\right)\end{aligned}$$

where z is the number of nearest neighbors that each micro-cell has.
Use has been made of the condition:

$$\sum_{s=1}^k \frac{n_s^j}{N} + \frac{n_E^j}{N} = 1,$$

The master equation

Introduce $\mathbf{n} = (\mathbf{n}^1, \mathbf{n}^2, \dots, \mathbf{n}^\Omega)$ where $\mathbf{n}^j = (n_1^j, n_2^j, \dots, n_k^j)$. Then:

$$\begin{aligned} \frac{dP(\mathbf{n}, t)}{dt} = & \sum_{j=1}^{\Omega} T_{loc}^j P(\mathbf{n}, t) + \sum_{j=1}^{\Omega} \sum_{j' \in j} T_{mig}^{jj'} P(\mathbf{n}, t) \\ & + \sum_{j=1}^{\Omega} T_{env}^j P(\mathbf{n}, t), \end{aligned} \quad (1)$$

where the three terms on the right-hand side refer to the **local terms** for the chemical reactions, the **migration** of chemical species between the micro-cells, and the **interaction** with the environment, respectively. The notation $j' \in j$ means that the cell j' is a nearest-neighbor of the cell j .

The van Kampen expansion

In analogy with above we set:

$$\frac{n_s^j}{N} = \phi_s^j + \frac{1}{\sqrt{N}} \xi_s^j$$

At the leading order in $\frac{1}{\sqrt{N}}$ we get:

$$\begin{aligned} \frac{d}{d\tau} \phi_s^j &= \frac{r}{\Omega} (\phi_{s-1}^j \phi_s^j - \phi_s^j \phi_{s+1}^j) + \frac{\alpha}{\Omega} \left(\Delta \phi_s^j (1 - \sum_m \phi_m^j) + \phi_s^j \sum_m \Delta \phi_m^j \right) \\ &+ \frac{\beta_{in}}{\Omega} (1 - \sum_m \phi_m^j) - \frac{\beta_{out}}{\Omega} \phi_s^j \end{aligned}$$

where Δ is the discrete Laplacian operator $\Delta f_s^j = (2/z) \sum_{j' \in j} (f_s^{j'} - f_s^j)$.

The next-to-leading order: Fokker Planck equation

$$\frac{\partial \Pi}{\partial \tau} = - \sum_p \frac{\partial}{\partial \xi_p} \left[A_p(\xi) \Pi \right] + \frac{1}{2} \sum_{l,p} B_{lp} \frac{\partial^2 \Pi}{\partial \xi_l \partial \xi_p},$$

where the matrix A can be re-written as

$$A_p(\xi) = \sum_l M_{pl} \xi_l.$$

To specify the Fokker-Planck equation we need to give the form of the two $(k\Omega) \times (k\Omega)$ matrices M and B .

The equivalent Langevin formulation

$$\frac{d\xi_s^j}{d\tau} = \sum_{j',r} M_{sr}^{jj'} \xi_r^{j'} + \lambda_s^j(\tau),$$

where

$$\langle \lambda_s^j(\tau) \lambda_r^{j'}(\tau') \rangle = B_{sr}^{jj'} \delta(\tau - \tau').$$

Going to Fourier space

$$\frac{d\xi_s^{\mathbf{k}}}{d\tau} = \sum_r M_{sr}^{\mathbf{k}} \xi_r^{\mathbf{k}} + \lambda_s^{\mathbf{k}}(\tau),$$

where

$$\langle \lambda_s^{\mathbf{k}}(\tau) \lambda_r^{\mathbf{k}'}(\tau') \rangle = B_{sr}^{\mathbf{k}} \Omega a^d \delta_{\mathbf{k}+\mathbf{k}',0} \delta(\tau - \tau'),$$

and where \mathbf{k} is the wavevector. Both $M^{\mathbf{k}}$ and $B^{\mathbf{k}}$ are simply $k \times k$ matrices (recall that k is the number of chemical species) and the analysis from now on is as in the non-spatial case.

The matrix $M^{\mathbf{k}}$

$$M_{sr}^{\mathbf{k}} = M_{sr}^{(NS)} + M_{sr}^{(SP)} \Delta_{\mathbf{k}},$$

where the two matrices $M^{(NS)}$ and $M^{(SP)}$ are given by

$$M_{ss}^{(NS)} = -\beta - \gamma$$
$$M_{sr}^{(NS)} = \begin{cases} -\eta\phi^* - \beta, & \text{if } r = s + 1 \\ \eta\phi^* - \beta, & \text{if } r = s - 1 \\ -\beta, & \text{if } |s - r| > 1, \end{cases}$$

and

$$M_{ss}^{(SP)} = \alpha_s [1 + (1 - k)\phi^*]$$

$$M_{sr}^{(SP)} = \alpha_s \phi^* \text{ if } s \neq r.$$

Fourier transform of the discrete Laplacian

$$\Delta_{\mathbf{k}} = \frac{2}{d} \sum_{\gamma=1}^d [\cos(\mathbf{k}_{\gamma} \mathbf{a}) - 1].$$

The matrix B^k

$$B_{sr}^k = B_{sr}^{(NS)} + B_{sr}^{(SP)} \Delta_k,$$

where the two matrices $M^{(NS)}$ and $M^{(SP)}$ are given by

$$B_{ss}^{(NS)} = a^d \left[\beta(1 - k\phi^*) + \gamma\phi^* + 2\eta(\phi^*)^2 \right] \quad (2)$$

$$B_{sr}^{(NS)} = \begin{cases} -a^d \eta(\phi^*)^2, & \text{if } r = s + 1 \\ -a^d \eta(\phi^*)^2, & \text{if } r = s - 1 \\ 0 & \text{if } |s - r| > 1, \end{cases} \quad (3)$$

and

$$B_{ss}^{(SP)} = -2a^d \alpha_s \phi^* (1 - k\phi^*) \quad (4)$$

$$B_{sr}^{(SP)} = 0 \text{ if } s \neq r. \quad (5)$$

Calculating the power spectrum of fluctuations

$$\sum_{r=1}^k (-i\omega\delta_{sr} - M_{sr}^{\mathbf{k}}) \tilde{\xi}_r^{\mathbf{k}}(\omega) = \tilde{\lambda}_s^{\mathbf{k}}(\omega),$$

Defining the matrix $\Phi_{sr}^{\mathbf{k}}(\omega) = (-i\omega\delta_{sr} - M_{sr}^{\mathbf{k}})$:

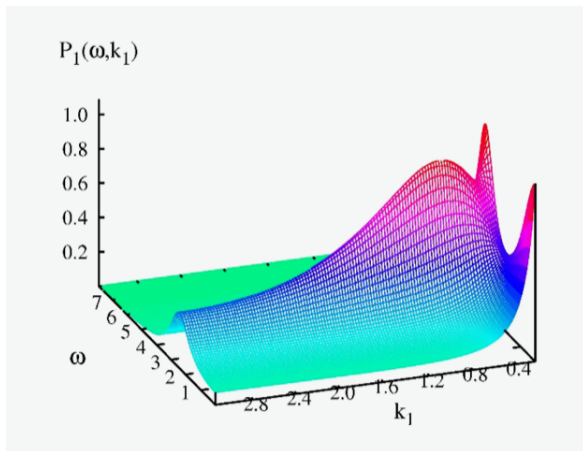
$$\tilde{\xi}_r^{\mathbf{k}}(\omega) = \sum_{s=1}^k [\Phi^{\mathbf{k}}(\omega)]_{rs}^{-1} \tilde{\lambda}_s^{\mathbf{k}}(\omega).$$

and:

$$P_s(\mathbf{k}, \omega) \equiv \langle |\xi_s^{\mathbf{k}}(\omega)|^2 \rangle = \Omega a^d \sum_{r=1}^k \sum_{u=1}^k [\Phi^{\mathbf{k}}(\omega)]_{sr}^{-1} B_{ru}^{\mathbf{k}} [\Phi^{\mathbf{k}\dagger}(\omega)]_{us}^{-1}.$$

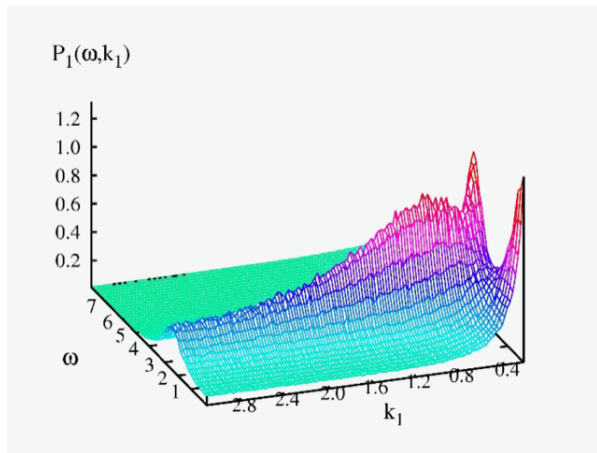
Analytical power spectrum species 1, $\alpha_i \neq 0$

- The **homogeneous** fixed point is stable: no deterministic patterns
- Stochastic patterns (**wave like**) do exist!



Numerical power spectrum species 1, $\alpha_j \neq 0$

- The **homogeneous** fixed point is stable: no deterministic patterns
- Stochastic patterns (**wave like**) do exist!



Summing up...

- **Autocatalytic** reactions are presumably important
- **Stochastic** (spatial/aspatial) model of autocatalytic **cycles**
- **Deterministic** models predict **homogeneous** fixed points
- **Patterns** (quasi cycles, waves) exist as seeded by **finite size** effects.

On the synchronization issue (highly speculative!)

- Imagine that the **vesicle** containing the chemical species **grows** because of the inclusion of successive membrane constituents

On the synchronization issue (highly speculative!)

- Imagine that the **vesicle** containing the chemical species **grows** because of the inclusion of successive membrane constituents
- Suppose that the vesicle is **filled by a discrete population** of chemical constituents, organized in a **autocatalytic cycle**.

On the synchronization issue (highly speculative!)

- Imagine that the **vesicle** containing the chemical species **grows** because of the inclusion of successive membrane constituents
- Suppose that the vesicle is **filled by a discrete population** of chemical constituents, organized in a **autocatalytic cycle**.
- The chemicals experience a first rapid evolution toward the stationary state, where **enhanced oscillations** appear due to the **intrinsic finiteness**.

On the synchronization issue (highly speculative!)

- Imagine that the **vesicle** containing the chemical species **grows** because of the inclusion of successive membrane constituents
- Suppose that the vesicle is **filled by a discrete population** of chemical constituents, organized in a **autocatalytic cycle**.
- The chemicals experience a first rapid evolution toward the stationary state, where **enhanced oscillations** appear due to the **intrinsic finiteness**.
- Such oscillations might **seed** an instability which could **resonate** with the innate ability of the container to divide, so initiating the **splitting process**.

In other words...

These oscillations could act as an **effective switch** by signaling to the membrane that the genetic evolution had been virtually taken to completion and that the **fission could now occur**, so ensuring that the genetic material is passed on to the **daughter protocells**.

Remark: discrete (network/lattice) support.

Linear Langevin equation

$$\frac{d\xi_{si}}{d\tau} = \sum_{r,j} M_{sr,ij} \xi_{r,j} + \eta_{si}(\tau)$$

where $\langle \eta_{si}(\tau) \eta_{rj}(\tau') \rangle = B_{sr,ij} \delta_{\tau\tau'}$

Generalized transform: expand along $\phi_i^{(\alpha)}$

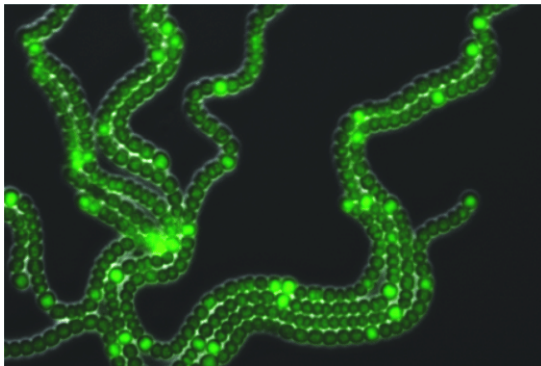
Transform

$$\tilde{f}_\alpha(\omega) = \int_0^\infty dt \sum_{i=1}^\Omega f_i(t) \phi_i^{(\alpha)} e^{-j\omega t}$$

Inverse Transform

$$f_i(t) = \frac{1}{2\pi} \int_{-\infty}^\infty d\omega \sum_{\alpha=1}^\Omega \tilde{f}_\alpha(\omega) \phi_i^{(\alpha)} e^{j\omega t}$$

Example 2. Pattern formation in Anabaena



- Under **nitrogen deprivation**, Anabaena develops **patterns** of nitrogen-fixing cells.
- **Heterocysts** are separated by nearly regular intervals of photosynthetic **vegetative** cells.

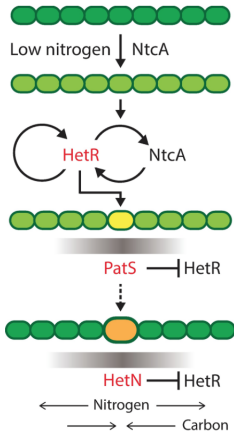


Fig 1. Basic regulatory network leading to heterocyst differentiation. In response to combined nitrogen deprivation, the NtcA protein is activated, leading to activation of expression of HetR in some cells. NtcA and HetR undergo mutual amplification, resulting in increased levels of the two regulators. In addition, HetR positively regulates its own production. During the early stages of differentiation, HetR induces expression of PatS in cells that can potentially form heterocysts (yellow). A PatS-derived peptide signal is thought to diffuse to neighbouring cells (grey gradient), where it interferes with the DNA-binding activity of HetR, causing its degradation and creating HetR gradients along filaments. At late stages (dashed arrow), HetN is produced in heterocysts (orange), and a HetN-derived signal is conveyed to neighbouring cells (grey gradient), where it inhibits HetR function and heterocyst formation. During the differentiation process, cells lose phycobilisomes and their autofluorescence declines (different shades of green). Phycobilisomes are restored when nitrogen compounds produced by the heterocysts reach the vegetative cells. For details, see [Introduction](#). The dynamical variables of our model, the activator HetR and its inhibitors PatS and HetN, have been emphasized in red. Adapted with modifications from [6].

- **Nitrogen** deprivation enhances expression **HetR**.
- HetR positively regulates its **own production**.
- HetR induces expression of **PatS** in cells that can potentially form **heterocysts** (yellow).

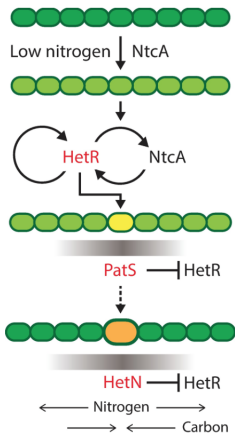


Fig 1. Basic regulatory network leading to heterocyst differentiation. In response to combined nitrogen deprivation, the NtcA protein is activated, leading to activation of expression of HetR in some cells. NtcA and HetR undergo mutual amplification, resulting in increased levels of the two regulators. In addition, HetR positively regulates its own production. During the early stages of differentiation, HetR induces expression of PatS in cells that can potentially form heterocysts (yellow). A PatS-derived peptide signal is thought to diffuse to neighbouring cells (grey gradient), where it interferes with the DNA-binding activity of HetR, causing its degradation and creating HetR gradients along filaments. At late stages (dashed arrow), HetN is produced in heterocysts (orange), and a HetN-derived signal is conveyed to neighbouring cells (grey gradient), where it inhibits HetR function and heterocyst formation. During the differentiation process, cells lose phycobilisomes and their autofluorescence declines (different shades of green). Phycobilisomes are restored when nitrogen compounds produced by the heterocysts reach the vegetative cells. For details, see [Introduction](#). The dynamical variables of our model, the activator HetR and its inhibitors PatS and HetN, have been emphasized in red. Adapted with modifications from [6].

- A PatS-derived peptide signal is thought to **diffuse** to neighbouring cells (grey gradient).
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- The **dynamical variables** of our model are the activator HetR and its inhibitors PatS and HetN.

Deterministic Turing patterns

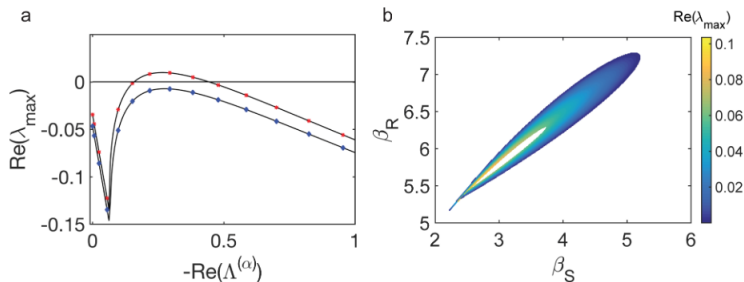
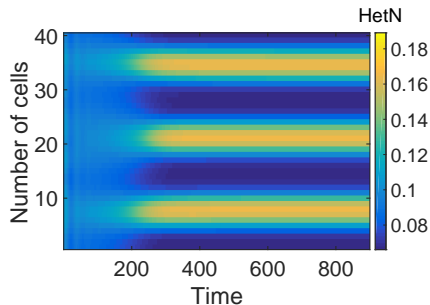


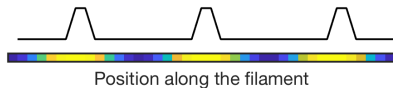
Fig 3. Conditions for a deterministic Turing instability. (a) Dispersion relations for $\beta_R = 6.5$ and $\beta_S = 3.65$ (blue diamonds) and $\beta_R = 6.5$ and $\beta_S = 3.7$ (red stars). The data used in this figure are included in S1 Data. (b) Region in the plane (β_S, β_R) where the maximum of $\lambda_{\text{Re}}(\Lambda^{(\alpha)})$ is positive and the equilibrium point is stable, for a ratio of diffusion coefficients $\frac{D_S}{D_N} = 3$. Parameters are set as $k_R = 0.2$, $\alpha_R = 0.2$, $K = 2$, $k_S = 0.1$, $\alpha_S = 0.1$, $\mu_S = 0.1$, $k_N = 0.7$, $\alpha_N = 0.3$, $\mu_N = 3$, $D_S = 3$, $D_N = 1$, and $\Omega = 40$.

<https://doi.org/10.1371/journal.pbio.2004877.g003>

Deterministic Turing patterns

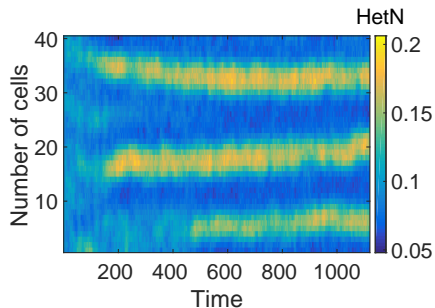


Project on the **1D** chain.



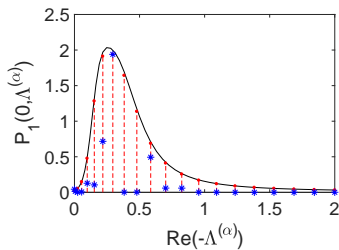
Pattern of **HetN**.

Stochastic Turing patterns

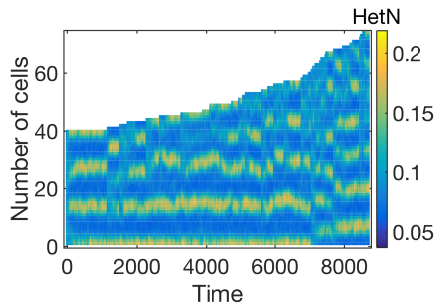
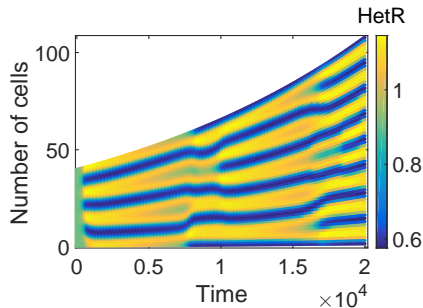


Pattern of **HetN**.

Power spectrum.



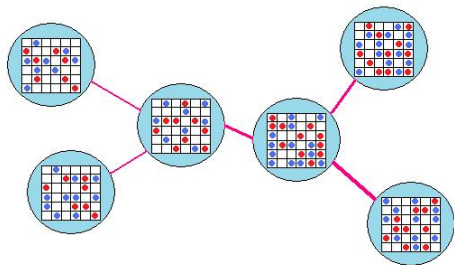
Accounting for cell duplication.



Example 3. Patterns seeded by stochastic finite size effects

Adjacency matrix

- X_i and Y_i identify individual elements of the two species.
- The index i stands for the node to which the elements belong.
- **Reactions** are local.
- **Migration** between adjacent nodes is allowed.

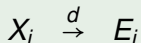
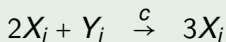
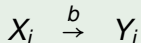
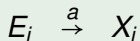


M. Asllani, Phys Rev E (2012)

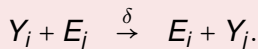
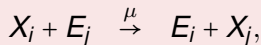
M. Asllani, Europ. Phys. J. B.(2013)

Stochastic moves: the Brusselator

Reaction rules



Diffusion between adjacent nodes



- $n_i \rightarrow$ number of molecule X in node i
- $m_i \rightarrow$ number of molecule Y in node i
- $N \rightarrow$ number of molecules which can be hosted in any node.

On the technical details

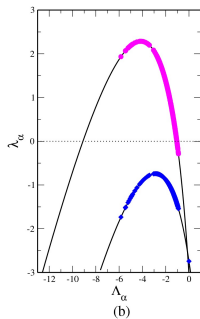
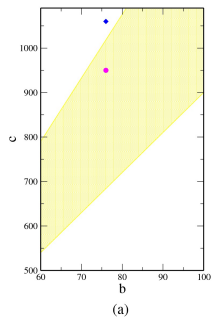
- Define the **discrete** concentrations $\mathbf{n} = (n_1, \dots, n_i, \dots, n_\Omega)$ and $\mathbf{m} = (m_1, \dots, m_i, \dots, m_\Omega)$
- Write down the **Master Equation** that governs the dynamics of the probability $P(\mathbf{n}, \mathbf{m}, t)$
- Working under the **linear noise approximation**:

$$\frac{n_i}{N} = \phi_i(t) + \frac{\xi_i}{\sqrt{N}}, \quad \frac{m_i}{N} = \psi_i(t) + \frac{\eta_i}{\sqrt{N}},$$

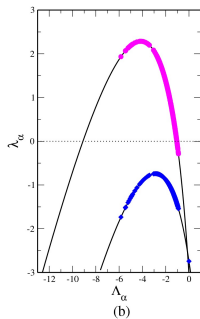
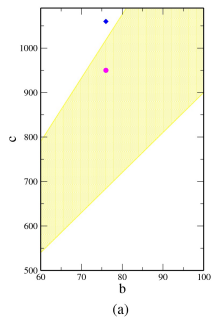
where $\phi_i(t)$ and $\psi_i(t)$ represent the **continuous** concentration.

- Perform a **perturbative** (system size) expansion in the small parameter $1/\sqrt{N}$.

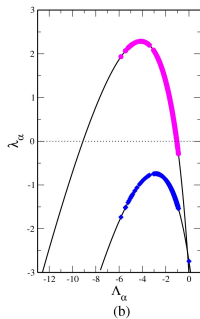
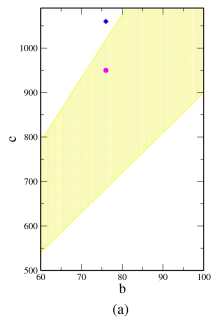
Leading order ($N^{-1/2}$) : mean-field equations



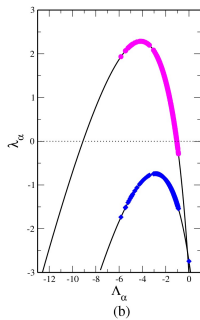
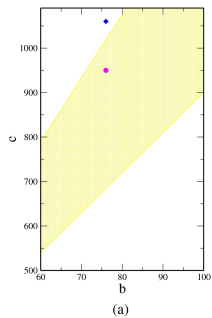
Leading order ($N^{-1/2}$) : mean-field equations



Leading order ($N^{-1/2}$) : mean-field equations



Stochastic patterns

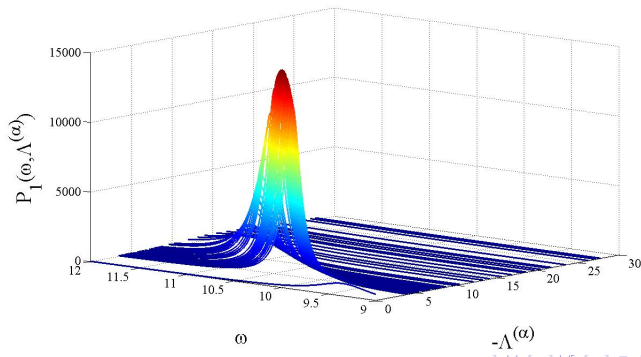


The next-to-leading order (N^{-1})

Linear Langevin equation

$$\frac{d\xi_{si}}{d\tau} = \sum_{r,j} M_{sr,ij} \xi_{r,j} + \eta_{si}(\tau)$$

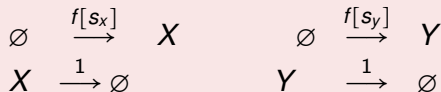
where $\langle \eta_{si}(\tau) \eta_{rj}(\tau') \rangle = B_{sr,ij} \delta_{\tau\tau'}$



Example 4. Excitatory-inhibitory neurons.

Label X and Y **individual** excitatory and inhibitory elements.

Birth-death scheme



where:

- $s_x = -r \left(\frac{n_Y}{V} - \frac{1}{2} \right)$.
- $s_y = r \left(\frac{n_X}{V} - \frac{1}{2} \right)$.
- $r > 0$ is the only free parameter.
- n_X and n_Y identify the number of elements of type X and Y .

Logic flow

- Introduce $P_n(t)$ to label the probability for the system to be in state $\mathbf{n} = (n_X, n_Y)$ at time t .
- The dynamics of the system is governed by a **master equation**.
- Perform a **Kramers-Moyal** expansion, $1/\sqrt{V}$ acting as small parameter.

Fluctuating hydrodynamic approximation

$$\begin{aligned}\dot{x} &= -x + f\left[-r\left(y - \frac{1}{2}\right)\right] + \frac{1}{\sqrt{V}} \left[x + f\left(-r\left(y - \frac{1}{2}\right)\right) \right]^{1/2} \eta_x \\ \dot{y} &= -y + f\left[r\left(x - \frac{1}{2}\right)\right] + \frac{1}{\sqrt{V}} \left[y + f\left(r\left(x - \frac{1}{2}\right)\right) \right]^{1/2} \eta_y\end{aligned}$$

- stochastic **non linear** equations.
- **multiplicative** noise.
- η_x and η_y are delta correlated Gaussian variables.

Deterministic limit, $V \rightarrow \infty$

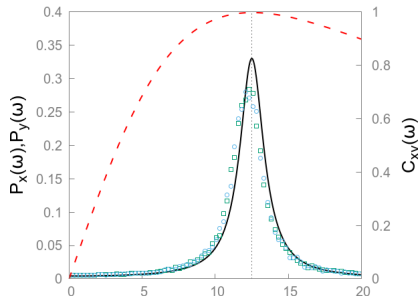
The deterministic model admits a **fixed point** $x^* = y^* = 1/2$. The linear stability analysis returns $\lambda = -1 \pm i\frac{r}{4}$

Fluctuations and quasi-cycles

Linear noise approximation

- $x(t) = x^* + V^{-1/2} \xi_1$
 $y(t) = y^* + V^{-1/2} \xi_2$
- $\dot{\xi}_i = \sum_j J_{ij} \xi_j + \eta_i$ with $i = 1, 2$

where $\eta_i(t)$ is a Gaussian noise with $\langle \eta_i(t) \eta_j(t') \rangle = \delta_{ij} \delta(t - t')$.



Power spectral density matrix

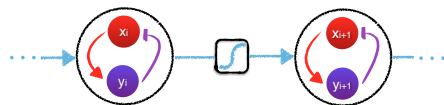
$$P_{ij}(\omega) = \langle \tilde{\xi}_i(\omega) \tilde{\xi}_j^*(\omega) \rangle = \sum_{l=1}^2 \sum_{m=1}^2 \Phi_{il}^{-1}(\omega) \delta_{lm} (\Phi^\dagger)^{-1}_{mj}(\omega)$$

where $\Phi_{ij} = -J_{ij} - i\omega \delta_{ij}$

Visualizing quasi-regular oscillations

Finite size corrections do matter: macroscopic **order** can emerge as mediated by the microscopic **disorder** (inherent granularity and stochasticity)

Directed chain of neuromorphic units



Birth-death scheme

$$\begin{array}{ccccc}
 \emptyset & \xrightarrow{f[s_{X_i}]} & X_i & \emptyset & \xrightarrow{f[s_{Y_i}]} & Y_i \\
 X_i & \xrightarrow{1} & \emptyset & Y_i & \xrightarrow{1} & \emptyset
 \end{array}$$

- $$s_{X_i} = -r \left(\frac{n_{Y_i}}{V} - \frac{1}{2} \right) + D \left(\frac{n_{X_{i-1}}}{V} - \frac{n_{X_i}}{V} \right) - D \left(\frac{n_{Y_{i-1}}}{V} - \frac{n_{Y_i}}{V} \right).$$
- $$s_{Y_i} = r \left(\frac{n_{X_i}}{V} - \frac{1}{2} \right) + D \left(\frac{n_{X_{i-1}}}{V} - \frac{n_{X_i}}{V} \right) - D \left(\frac{n_{Y_{i-1}}}{V} - \frac{n_{Y_i}}{V} \right).$$

- $r > 0$ and $D > 0$ are free parameters.
- n_{X_k} and n_{Y_k} identify the number of elements of type X and Y in cell k .

Deterministic limit

$$\begin{aligned}\dot{x}_i &= -x_i + f \left[-r \left(\frac{n_{Y_i}}{V} - \frac{1}{2} \right) + D \left(\frac{n_{X_{i-1}}}{V} - \frac{n_{X_i}}{V} \right) - D \left(\frac{n_{Y_{i-1}}}{V} - \frac{n_{Y_i}}{V} \right) \right] \\ \dot{y}_i &= -y_i + f \left[r \left(\frac{n_{X_i}}{V} - \frac{1}{2} \right) + D \left(\frac{n_{X_{i-1}}}{V} - \frac{n_{X_i}}{V} \right) - D \left(\frac{n_{Y_{i-1}}}{V} - \frac{n_{Y_i}}{V} \right) \right]\end{aligned}$$

Homogeneous fixed point:

$$x^* = y^* = 1/2 \quad \forall i.$$

Jacobian and stability.

$$J = \begin{pmatrix} E_1 & 0 & 0 & 0 & 0 \\ S_2 & E_2 & 0 & 0 & 0 \\ 0 & S_3 & E_3 & 0 & 0 \\ 0 & 0 & \ddots & \ddots & 0 \\ 0 & 0 & 0 & S_\Omega & E_\Omega \end{pmatrix}$$

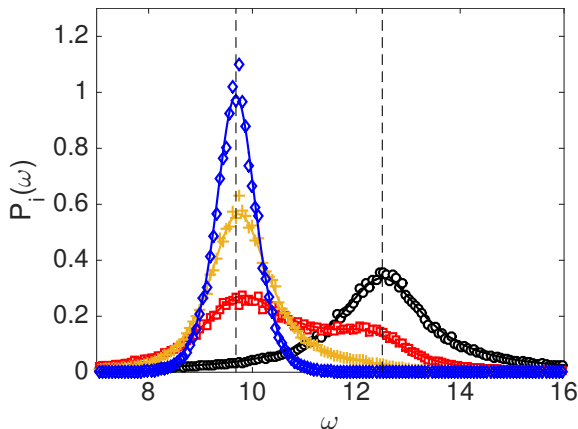
E_i are identical for $i \geq 2$

- $\lambda_{1,2} = -1 \pm i \frac{r}{4}$
- $(\lambda_i)_{3,4} = -1 \pm \sqrt{-\frac{r}{8} \left(\frac{r}{2} - D \right)}$
 $i = 2, \dots, \Omega$

Stable for $D < r/2$.

$$\omega_0 = r/4, \quad \omega_1 = \sqrt{\frac{r}{8} \left(\frac{r}{2} - D \right)}.$$

Linear noise approximation: theory vs. simulations.

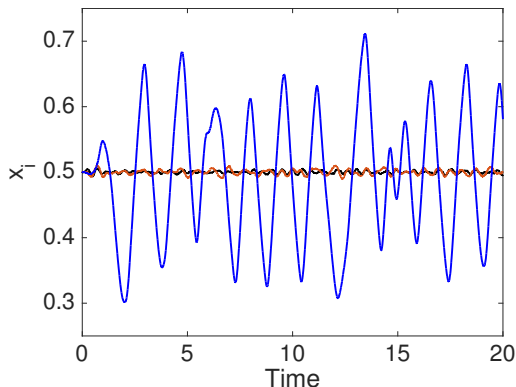


Act on **D** to **select** a frequency and **amplify** the signal along the chain.

On the amplification process

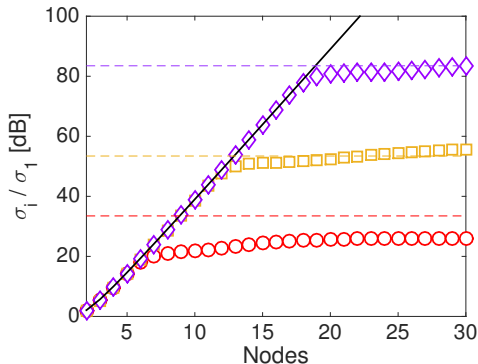
Stochastic trajectories as seen on different nodes of the chain.

Superposing the outcome



The amplification proceeds steadily along the chain. **Saturation** is attained when the fluctuations hit the boundaries.

The amplification is **exponential**.

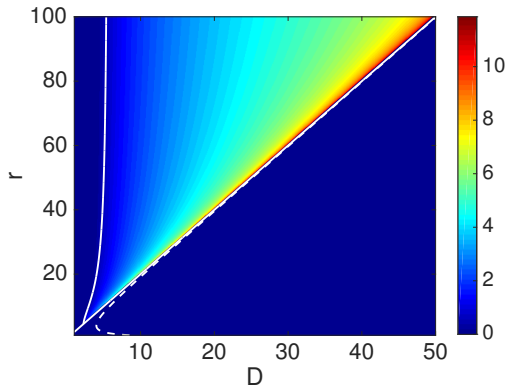


- Π is the distribution of fluctuations
- σ_i is the associated standard deviation on node i

Fokker-Planck equation

$$\frac{\partial}{\partial \tau} \Pi = - \sum_{i=1}^{2\Omega} \frac{\partial}{\partial \zeta_i} [(J\zeta)_i \Pi] + \frac{1}{2} \sum_{i=1}^{2\Omega} \frac{\partial^2}{\partial \zeta_i^2} \Pi$$

The rate of exponential amplification in the (D,r) plane.



The region of parameters that yields the sought amplification is delimited by the white solid curves (**exact** [right] and **approximate** [left]).

I. Tuning the frequency

II. Detecting a small amplitude noisy signal

Deterministic + noisy external input on node 1.