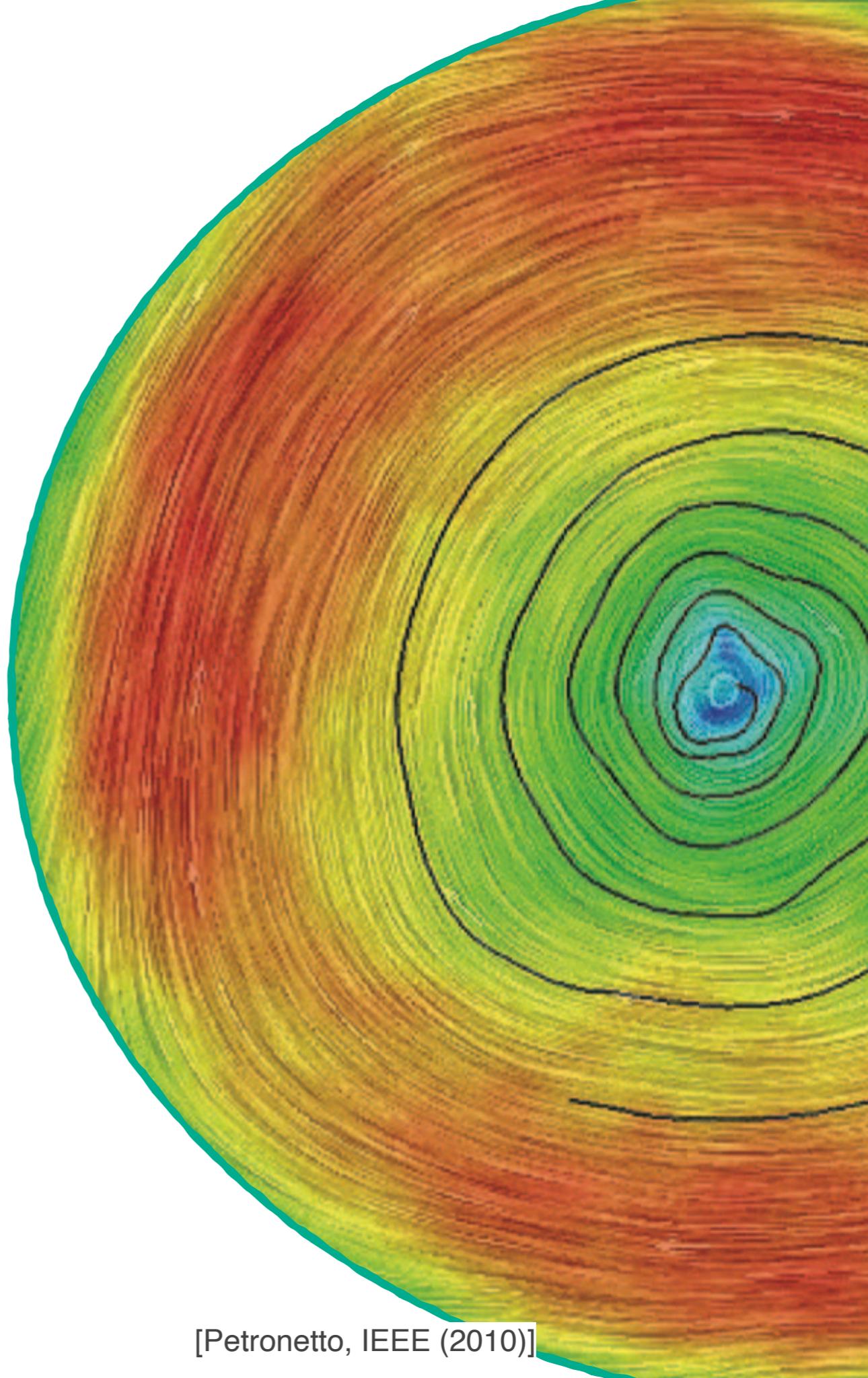


Schnakenberg without Schnakenberg

Sara Dal Cengio
LIPhy Grenoble

WOST III
31/05/22



[Petronetto, IEEE (2010)]

Identifying nonequilibrium forces

Identifying nonequilibrium forces

Helmholtz decomposition ($\mathbb{R}^2, \mathbb{R}^3$)

$$\mathbf{f} = -\nabla V + \nabla \times \mathbf{A}$$

The diagram illustrates the Helmholtz decomposition of a vector field \mathbf{f} into a conservative part $-\nabla V$ and a non-conservative part $\nabla \times \mathbf{A}$. On the left, a circular arrangement of black arrows pointing radially inward represents a conservative field. In the center, a radial arrangement of blue arrows pointing toward a central point represents another conservative field. On the right, a circular arrangement of red arrows forming a closed loop represents a non-conservative field, specifically a circulation or vorticity.

Conservative Non-conservative

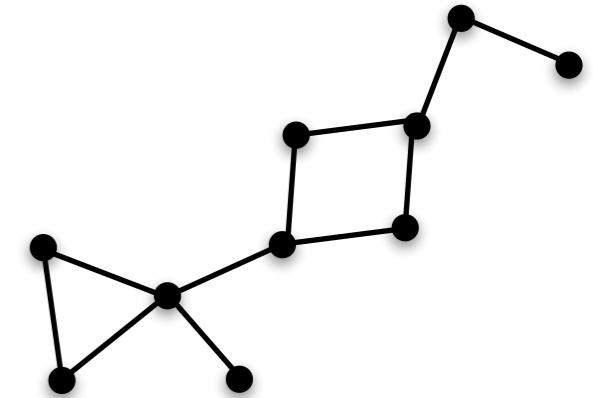
In physics: $\mathbf{f}_{nc} = 0 \iff$ Equilibrium

Identifying nonequilibrium forces

Helmholtz decomposition ($\mathbb{R}^2, \mathbb{R}^3$)

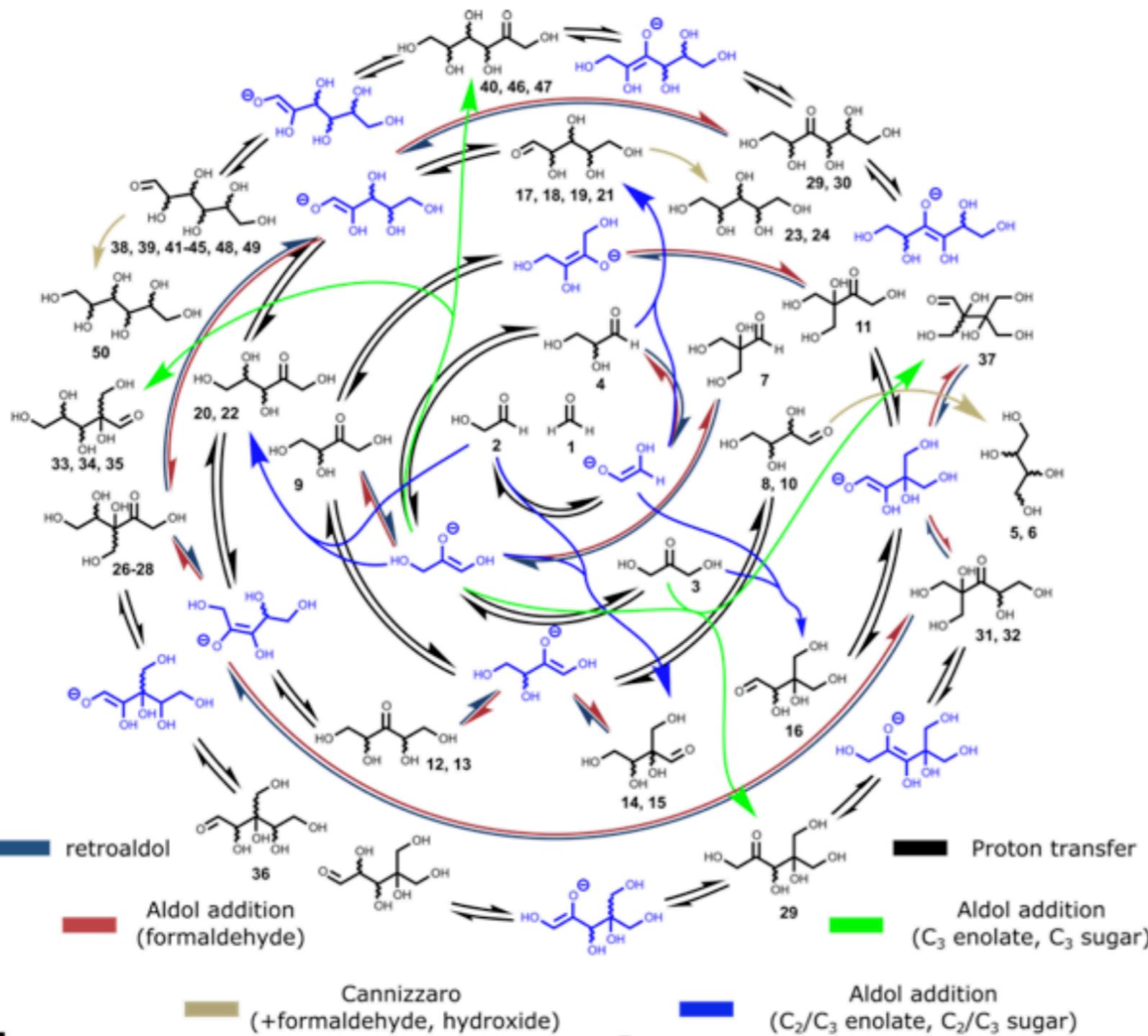
$$\mathbf{f} = -\nabla V + \nabla \times \mathbf{A}$$

Conservative Non-conservative



What if the space of configurations is a network?

Chemical reaction networks



Topological complexity

Biological functions

Nonequilibrium

Chemical reaction networks

How to identify nonequilibrium (chemical) forces in complex topology?

Rev. Mod. Phys. (1976)

**Network theory of microscopic and macroscopic behavior
of master equation systems**

J. Schnakenberg

Institut für Theoretische Physik, Rheinisch-Westfälische Technische Hochschule, Aachen, West Germany

Chemical reaction networks

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- Microscopic description: **population dynamics**
- Decomposition of forces based on **graph theory**
- Importance of **chemical cycles** for non-equilibrium steady states

Chemical reaction networks

How to identify nonequilibrium (chemical) forces in complex topology?

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Problem: graph at the level of population is impractical!

Deterministic rate equations

- For large systems: evolution at the average level

$$x_i \equiv \lim_{N,V \rightarrow \infty} \frac{N_i}{V}$$

Deterministic rate equations

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\mathbb{S} : Stoichiometric matrix encodes the topology

Deterministic rate equations

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\mathbb{S} : Stoichiometric matrix encodes the topology

- Pair of current and affinity for each reaction ρ

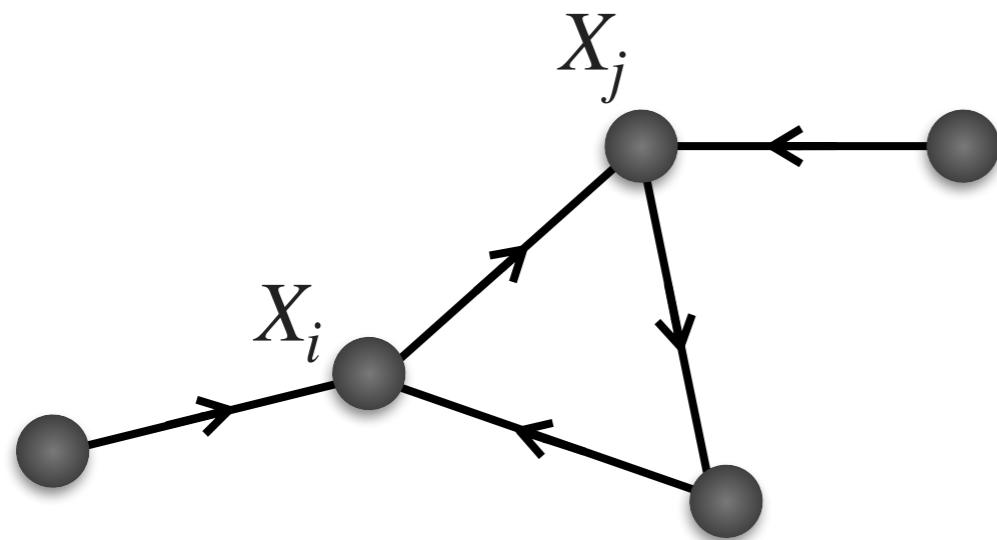
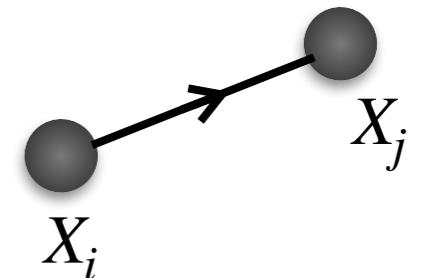
$$J_\rho(x), A_\rho(x)$$

Noninteracting CRNs

Unimolecular reactions: $X_i \rightleftharpoons X_j$

Noninteracting CRNs

Unimolecular reactions: $X_i \rightleftharpoons X_j$



Planar graph representation:

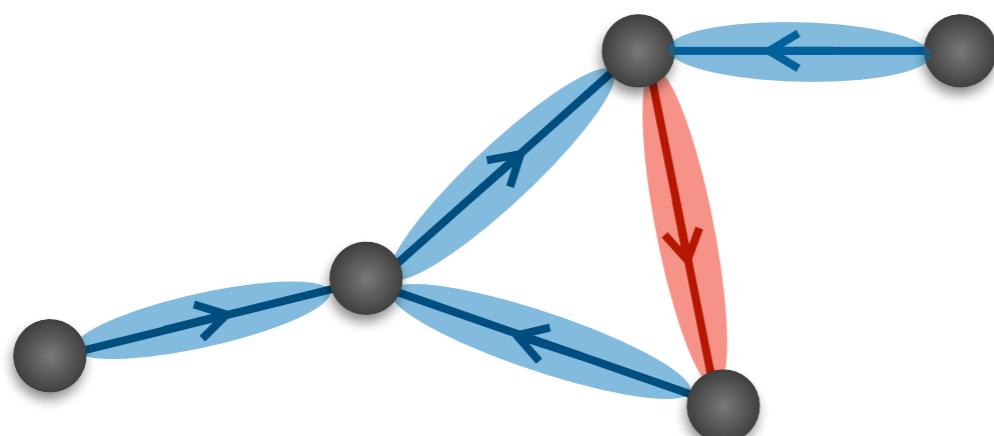
Nodes \iff Species

Edges \iff Reactions

\neq graph at the population level!

Noninteracting CRNs

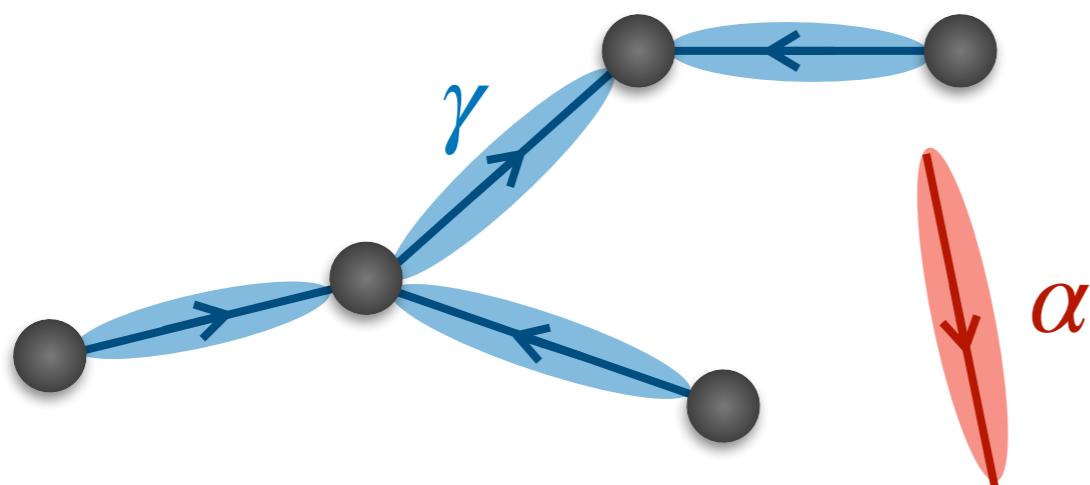
Unimolecular reactions: $X_i \rightleftharpoons X_j$



Choosing a spanning tree

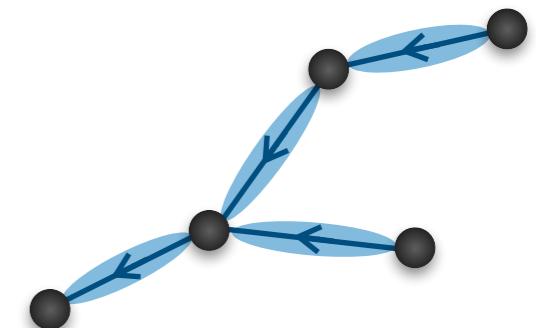
Noninteracting CRNs

Unimolecular reactions: $X_i \rightleftharpoons X_j$

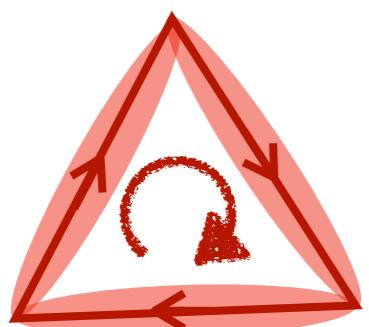
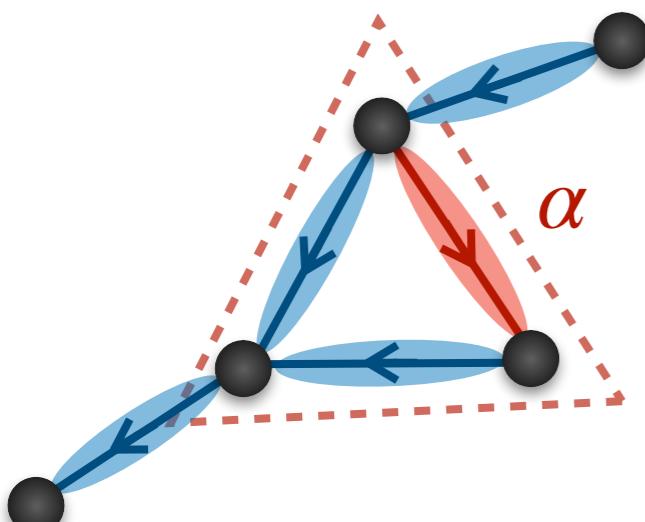


Choosing a spanning tree

Cycles & cocycles

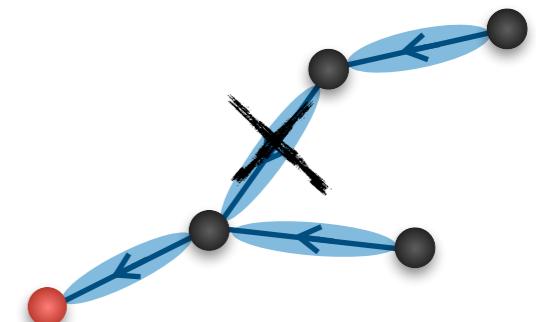


Adding an edge back:

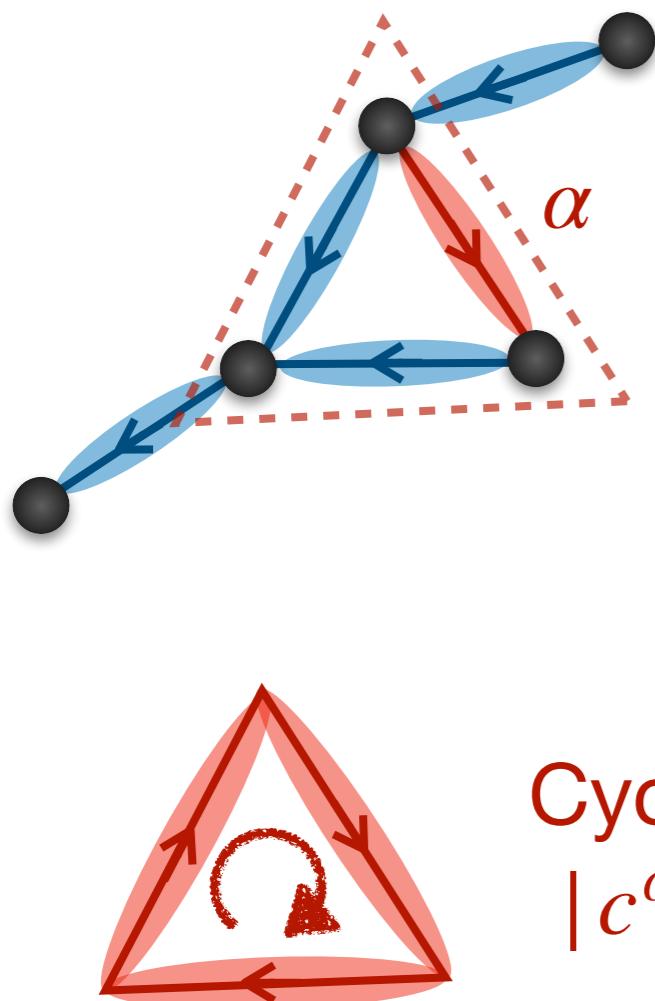


Cycle
 $|c^\alpha\rangle$

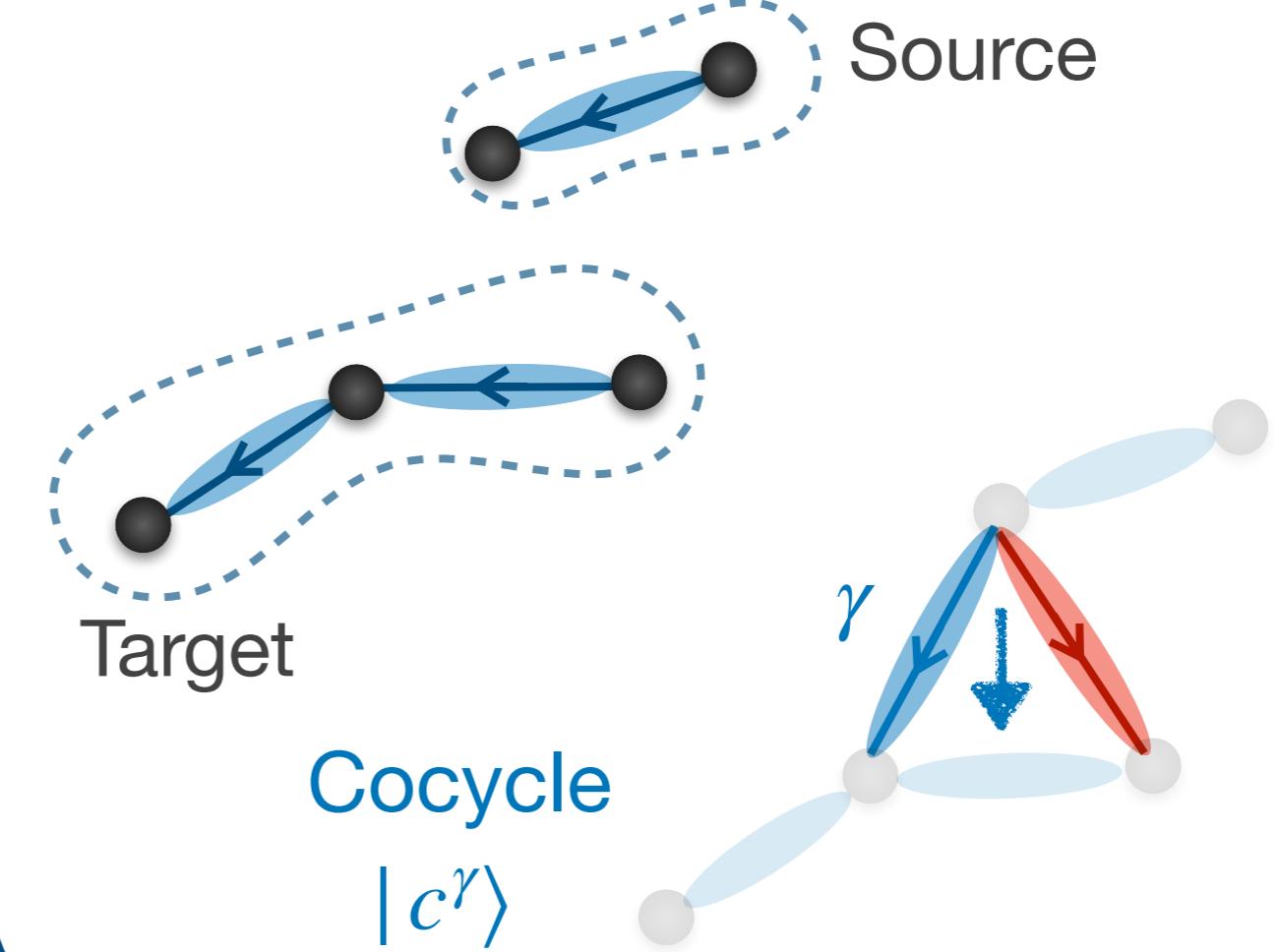
Cycles & cocycles



Adding an edge back:



Removing a edge:

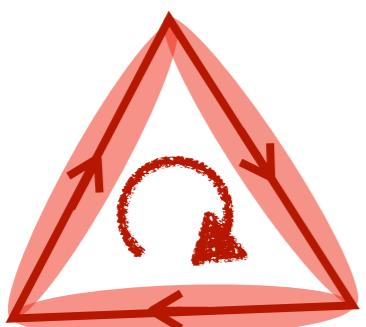


Cycles & cocycles

Adding an edge back:

Removing a edge:

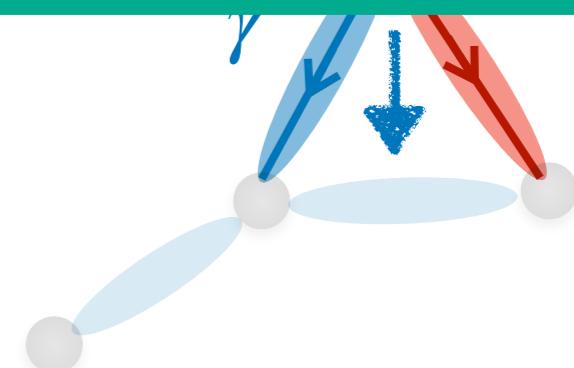
How can they be used to decompose the chemical affinities?



Cycle
 $|c^\alpha\rangle$

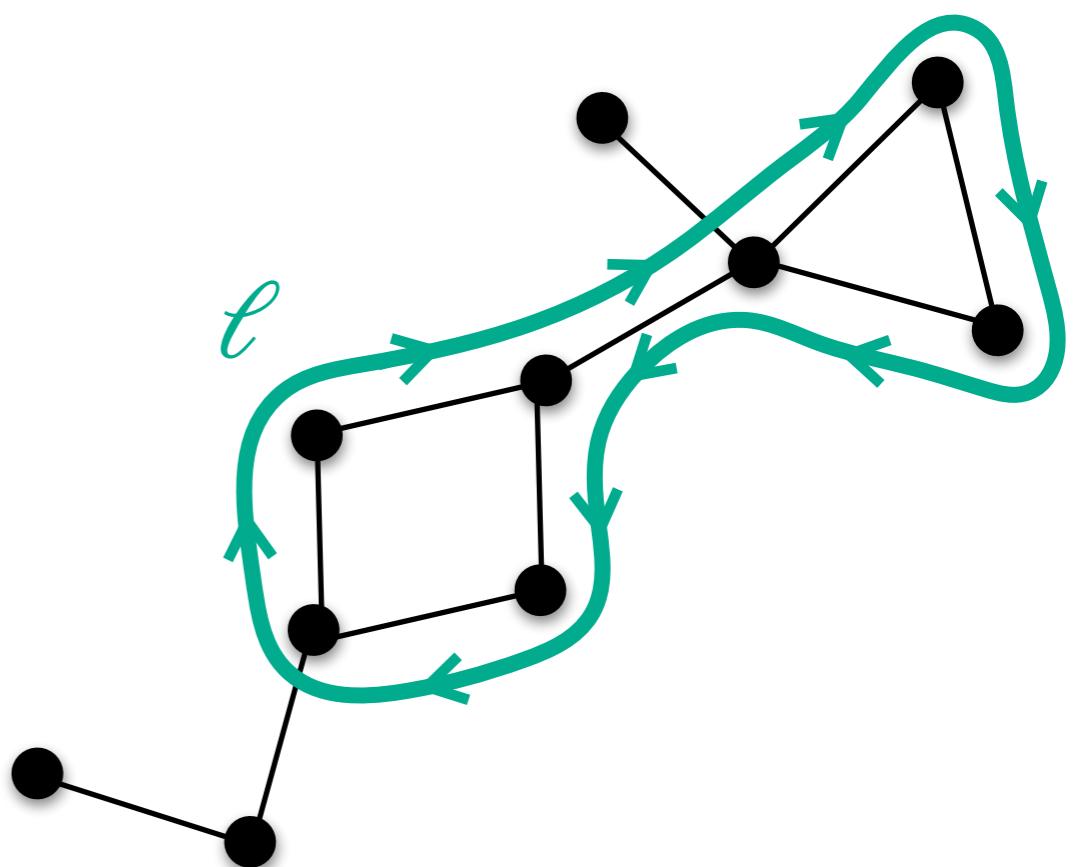
Target

Cocycle
 $|c^\gamma\rangle$



Non-conservative affinities

A generic loop on the graph is a linear combination of cycles



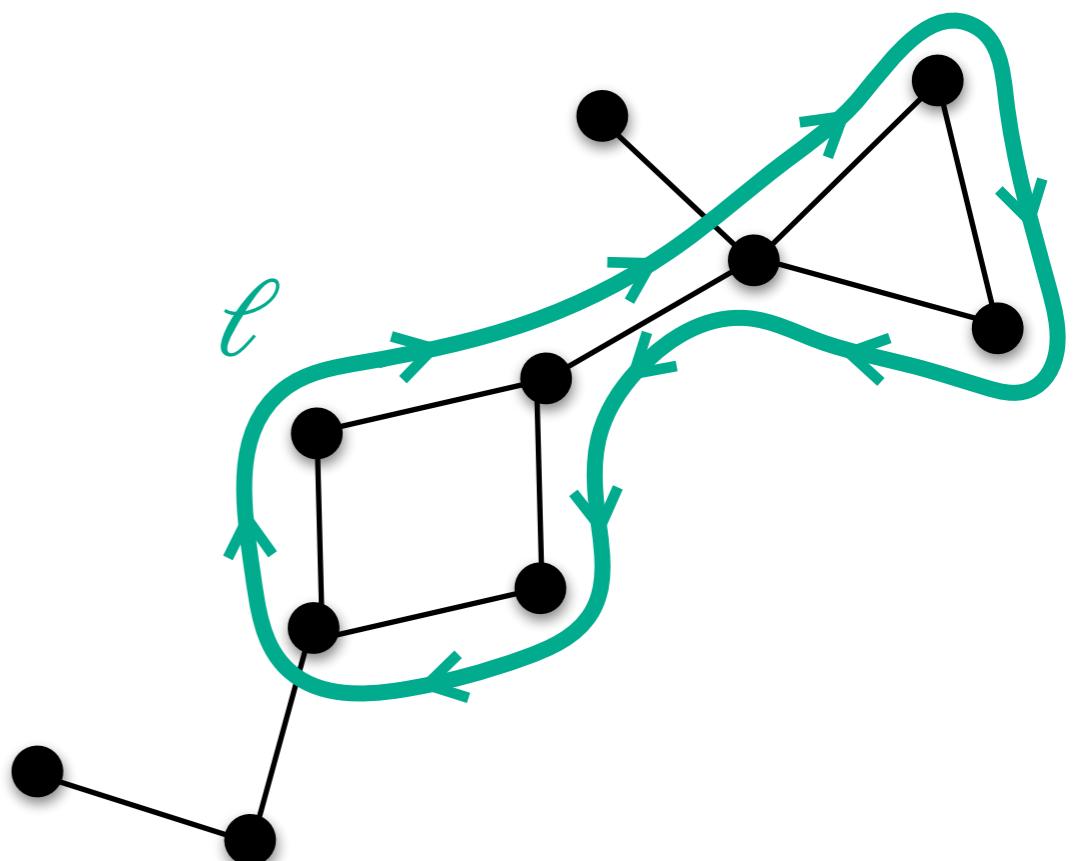
Non-conservative affinities

A generic loop on the graph is a linear combination of cycles

Circulation of the affinity:

$$\langle A | c^\alpha \rangle = A_\alpha$$

Non-conservative affinities



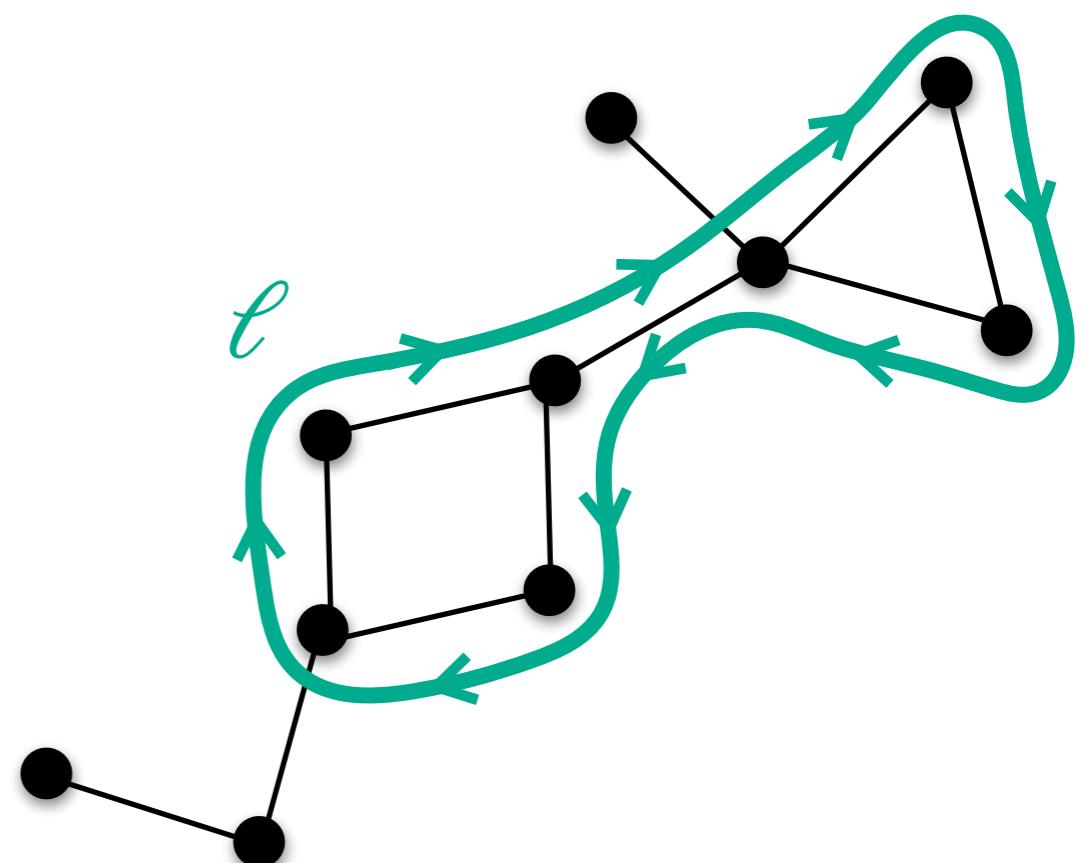
Non-conservative affinities

A generic loop on the graph is a linear combination of cycles

Circulation of the affinity:

$$\langle A | c^\alpha \rangle = A_\alpha$$

Non-conservative affinities



$$A_\alpha = 0 \quad \forall \alpha \iff \text{Conservative affinity (KCL)}$$

[Schnakenberg]

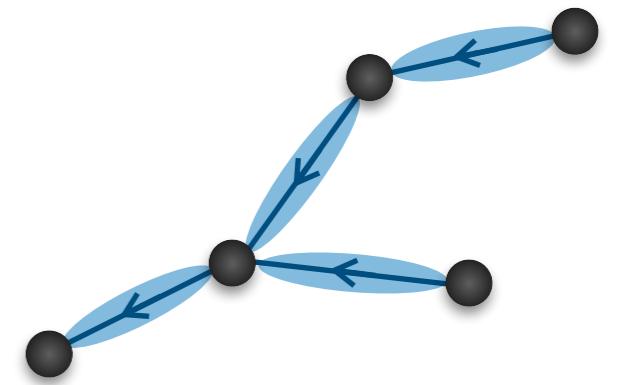
Conservative affinities

The cocycles form a basis for the conservative affinities

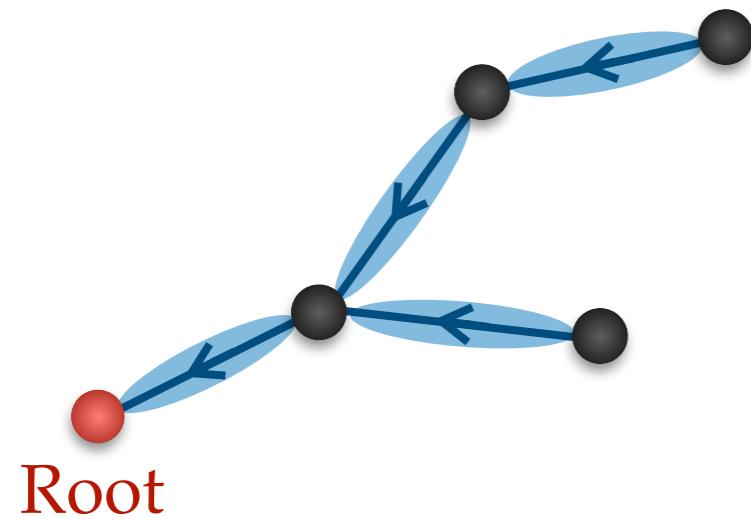
In the case of conservative dynamics:

$$|A\rangle = \sum_{\gamma} A_{\gamma} |c^{\gamma}\rangle$$

Affinities of the reactions which
form the spanning tree



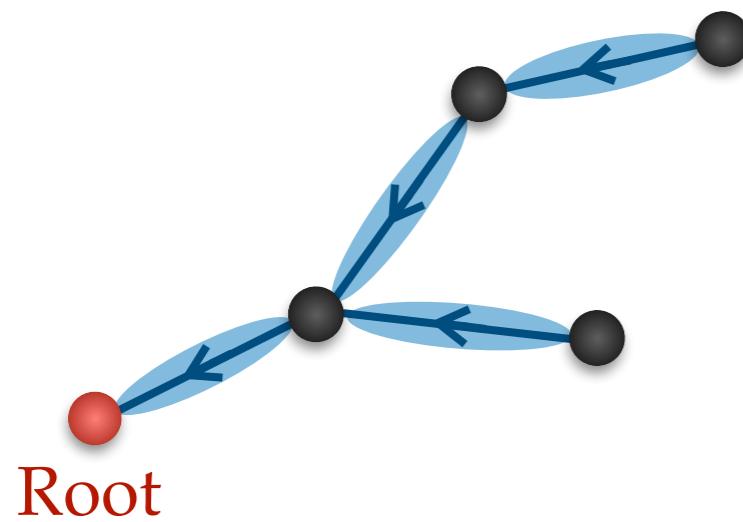
Conservative affinities



Integrating on the spanning tree:

$$G^T = \begin{pmatrix} 1 & 0 & 0 & 0 & 0 \\ 1 & 1 & 0 & 0 & 0 \\ 1 & 1 & 1 & 0 & 0 \\ 1 & 1 & 1 & 1 & 0 \\ 1 & 1 & 0 & 0 & 1 \end{pmatrix}$$

Conservative affinities



Integrating on the spanning tree:

$$G^T = \begin{pmatrix} 1 & 0 & 0 & 0 & 0 \\ 1 & 1 & 0 & 0 & 0 \\ 1 & 1 & 1 & 0 & 0 \\ 1 & 1 & 1 & 1 & 0 \\ 1 & 1 & 0 & 0 & 1 \end{pmatrix}$$

...Underlying potential landscape?

$$\sum_{[i \rightarrow \text{root}]} A_\gamma = \mu_i$$

Chemical potential

$$\sum_{[i \rightarrow \text{root}]} \log \left(\frac{k_\gamma^-}{k_\gamma^+} \right) = \mu_i^\Theta$$

Standard chemical potential

A physical decomposition

From graph theory:

- A criterion for the affinity to be conservative
- A basis for the conservative affinities

A physical decomposition

From graph theory:

- A criterion for the affinity to be conservative
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$\{A_\gamma\}$ Complete set of conservative affinities

$\{A_\alpha\}$ Complete set of non-conservative affinities

Interacting CRNs

Example



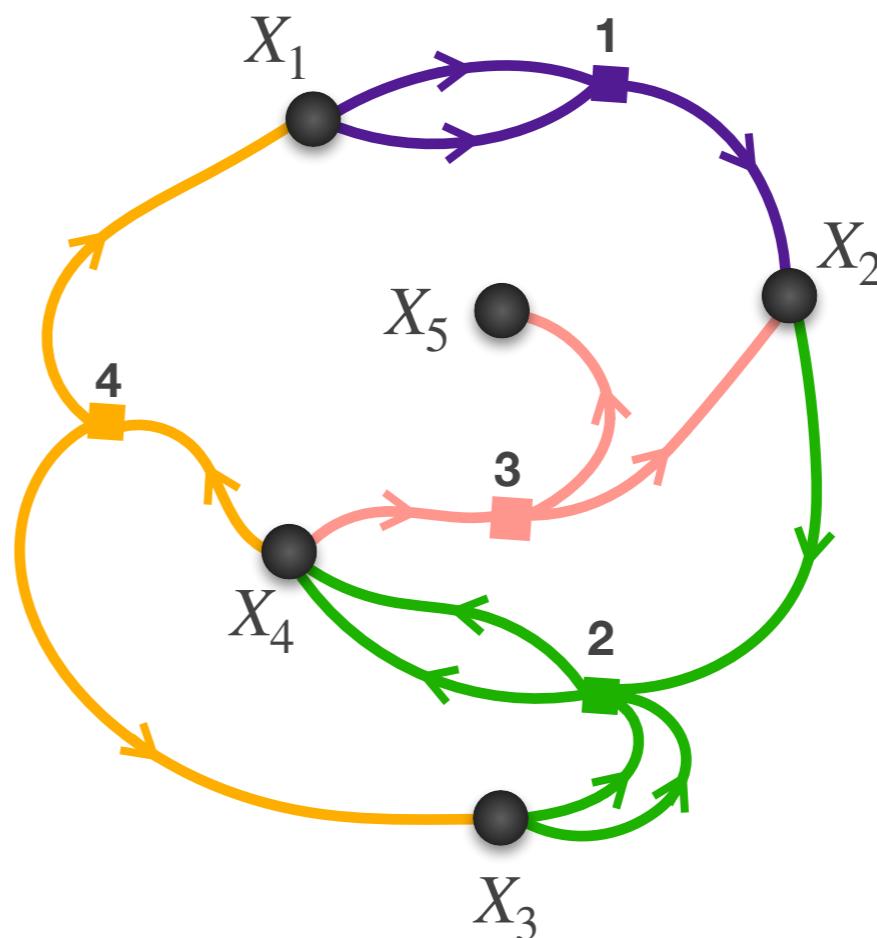
Ex. Autocatalytic networks, epidemic dynamics...

Interacting CRNs

Example



Hypergraph representation:



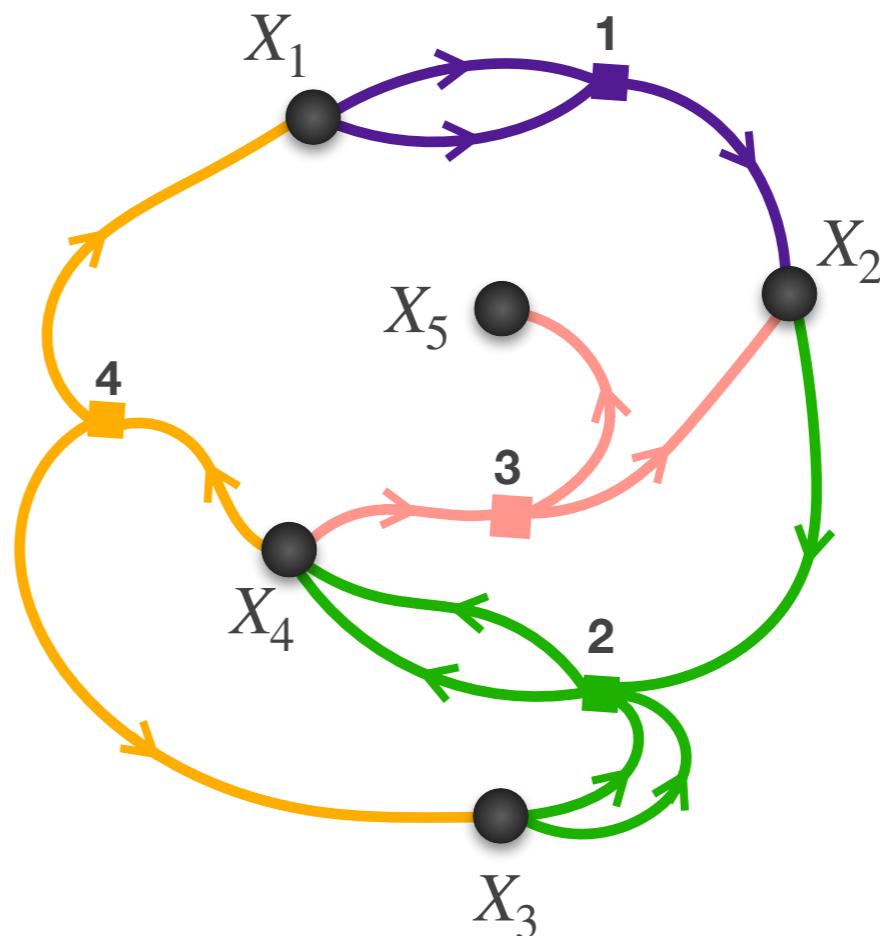
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Interacting CRNs

Example



Hypergraph representation:



Ex. Autocatalytic networks, epidemic dynamics...

No longer a spanning tree... what about cycles and cocycles?

Gauss-Jordan elimination

Row reduced Echelon form for \mathbb{S} :

$$G\mathbb{S} = \begin{pmatrix} \mathbf{1}_M & \mathbb{T} \\ 0 & 0 \end{pmatrix}$$

Algebraic tool

Gauss-Jordan elimination

Row reduced Echelon form for \mathbb{S} :

$$G\mathbb{S} = \begin{pmatrix} \mathbf{1}_M & \mathbb{T} \\ 0 & 0 \end{pmatrix}$$

Fractional entries, unique

Algebraic tool



$$(|c^\gamma\rangle, |c^\alpha\rangle) = \begin{pmatrix} \mathbf{1}_M & -\mathbb{T} \\ \mathbb{T}^\top & \mathbf{1}_{\#cy} \end{pmatrix}$$

$|c^\alpha\rangle$ Still a basis for $\text{Ker } \mathbb{S}$

$|c^\gamma\rangle$ Still a basis for $\text{Im } \mathbb{S}^\top$

Geometry of hypergraphs

Integrator operator on the hypergraph:

$$G^T = \left(\begin{array}{c|ccc} \text{Conservation laws} & 0 & \cdots & 0 \\ \hline 0 & \cdots & & 0 \\ 0 & & & 0 \end{array} \right) \quad \begin{matrix} \uparrow \\ \downarrow \end{matrix} \quad \# \text{ csv} = \# \text{ roots}$$

Integration paths

Geometry of hypergraphs

Integrator operator on the hypergraph:

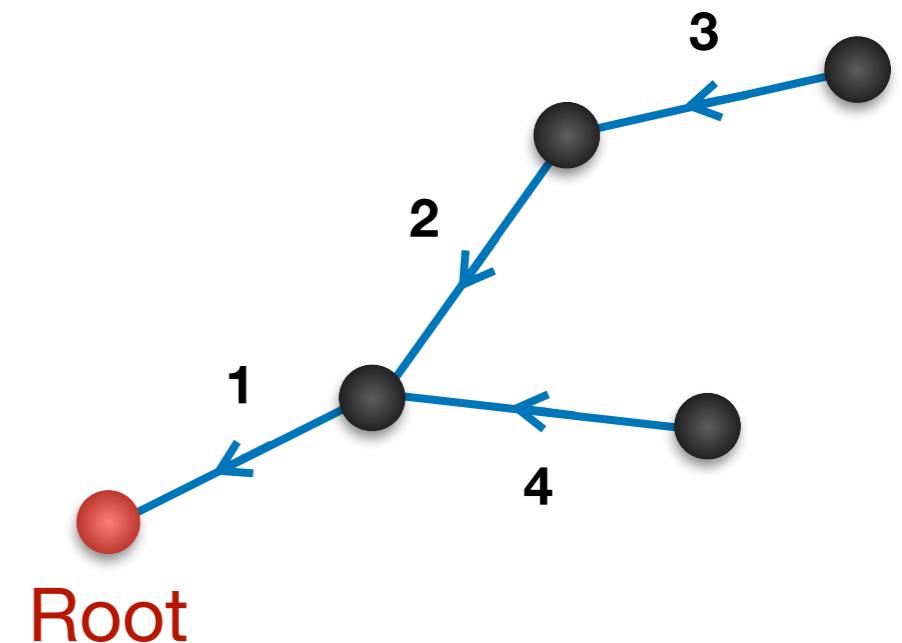
$$G^T = \left(\begin{array}{c|cccc} \text{Conservation laws} & 0 & \dots & 0 \\ \hline 0 & \dots & & 0 \\ 0 & & & 0 \end{array} \right) \quad \# \text{ csv} = \# \text{ roots}$$

Integration paths

For the simple graph:

$$G^T = \left(\begin{array}{c|ccccc} 1 & 0 & 0 & 0 & 0 \\ \hline 1 & 1 & 0 & 0 & 0 \\ 1 & 1 & 1 & 0 & 0 \\ 1 & 1 & 1 & 1 & 0 \\ 1 & 1 & 0 & 0 & 1 \end{array} \right) \quad 1 \text{ root}$$

Mass conservation law



Geometry of hypergraphs

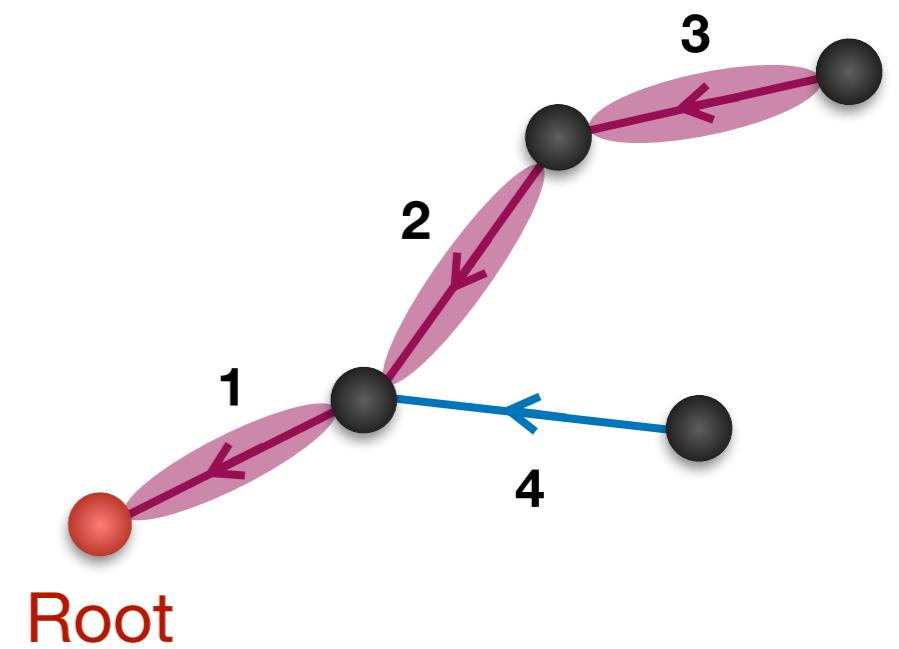
Integrator operator on the hypergraph:

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Integration paths

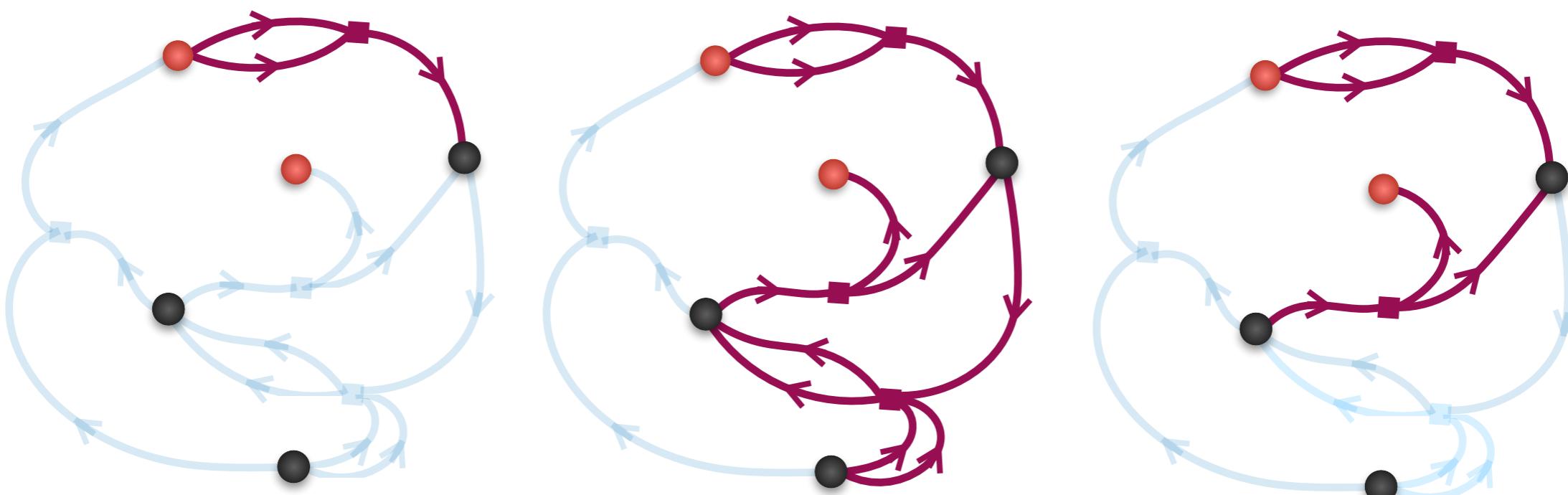
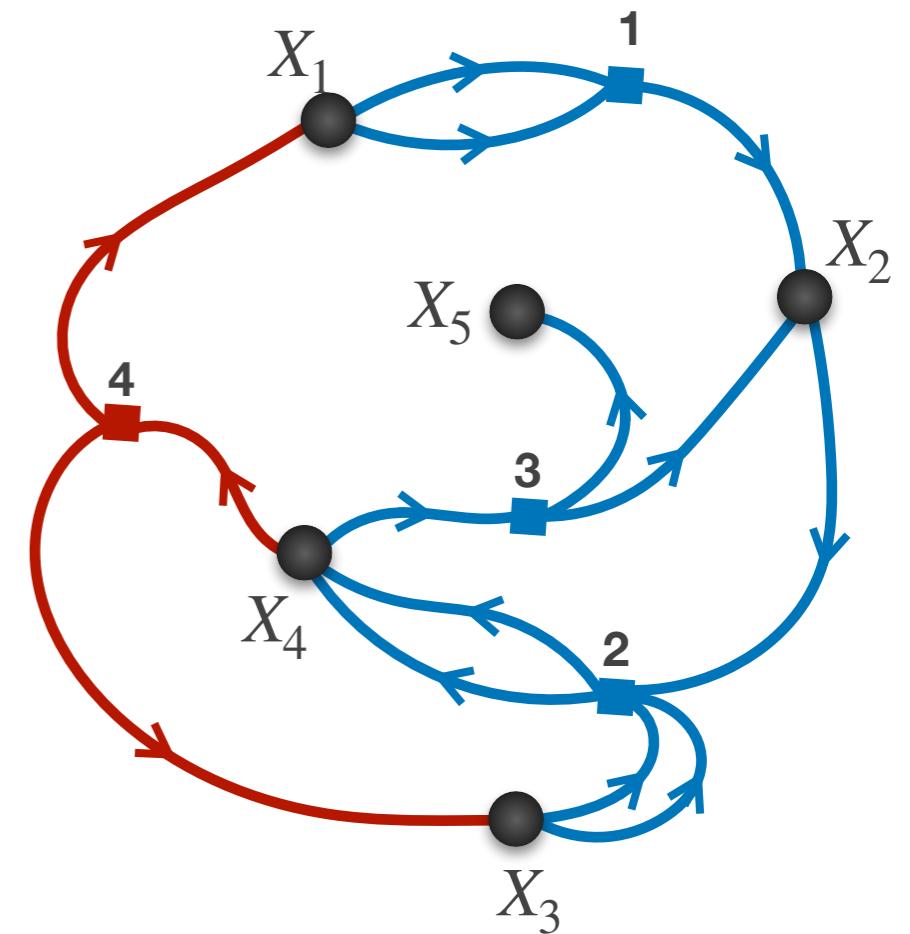
For the simple graph:

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Geometry of hypergraphs

$$G^T = \begin{pmatrix} 1 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 0 \\ 2 & 0 & 1 & 0 & 0 \\ 1 & 1 & 1/2 & -1/2 & -1 \\ 2 & 1 & 1 & 0 & -1 \end{pmatrix}$$



Onsager reciprocities

Natural framework to identify Onsager cross-couplings in linear response

Onsager reciprocities

Natural framework to identify Onsager cross-couplings in linear response

→ **Transient response in reversible dynamics:**

$$J_\gamma \equiv \langle c^\gamma | J \rangle = \langle c^\gamma | \Lambda | A \rangle = \langle c^\gamma | \Lambda | c^{\gamma'} \rangle A_{\gamma'}$$

Onsager reciprocities

Natural framework to identify Onsager cross-couplings in linear response

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M transient currents

M conservative affinities

Response matrix of relaxation

The diagram illustrates the mathematical expression for transient currents J_γ . It is shown as a product of three components: a blue circle on the left labeled "M transient currents" containing the term $\langle c^\gamma | J \rangle$, a central bracket under the term $\langle c^\gamma | \Lambda | c^{\gamma'} \rangle$ labeled "Response matrix of relaxation", and a blue circle on the right labeled "M conservative affinities" containing the term $A_{\gamma'}$.

Onsager reciprocities

Natural framework to identify Onsager cross-couplings in linear response

→ **Transient response in reversible dynamics:**

$$J_\gamma \equiv \langle c^\gamma | J \rangle = \langle c^\gamma | \Lambda | A \rangle = \langle c^\gamma | \Lambda | c^{\gamma'} \rangle A_{\gamma'}$$

→ **Steady-state response due to external drive (chemostatting)**

$$A_\alpha \equiv \langle c^\alpha | A \rangle = \langle c^\alpha | \Lambda^{-1} | J \rangle = \langle c^\alpha | \Lambda^{-1} | c^{\alpha'} \rangle J_{\alpha'}$$

Onsager reciprocities

Natural framework to identify Onsager cross-couplings in linear response

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Non-conservative forces

Steady-state currents

Onsager reciprocities

Natural framework to identify Onsager cross-couplings in linear response

→ Transient response in reversible dynamics:

$$J_\gamma \equiv \langle c^\gamma | J \rangle = \langle c^\gamma | \Lambda | A \rangle = \langle c^\gamma | \Lambda | c^{\gamma'} \rangle A_{\gamma'}$$

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How are the two matrices related?

Onsager reciprocities

The two response matrices, after a change of variables:

$$L^{\text{rel}} = \begin{pmatrix} \mathbf{1}_M + \tilde{\mathbb{T}}\tilde{\mathbb{T}}^\top & (0) \\ \hline (0) & \mathbf{0}_{\#\text{cy}} \end{pmatrix}$$

$$L^{\text{ext}} = \begin{pmatrix} \mathbf{0}_M & (0) \\ \hline (0) & \mathbf{1}_{\#\text{cy}} + \tilde{\mathbb{T}}^\top\tilde{\mathbb{T}} \end{pmatrix}$$

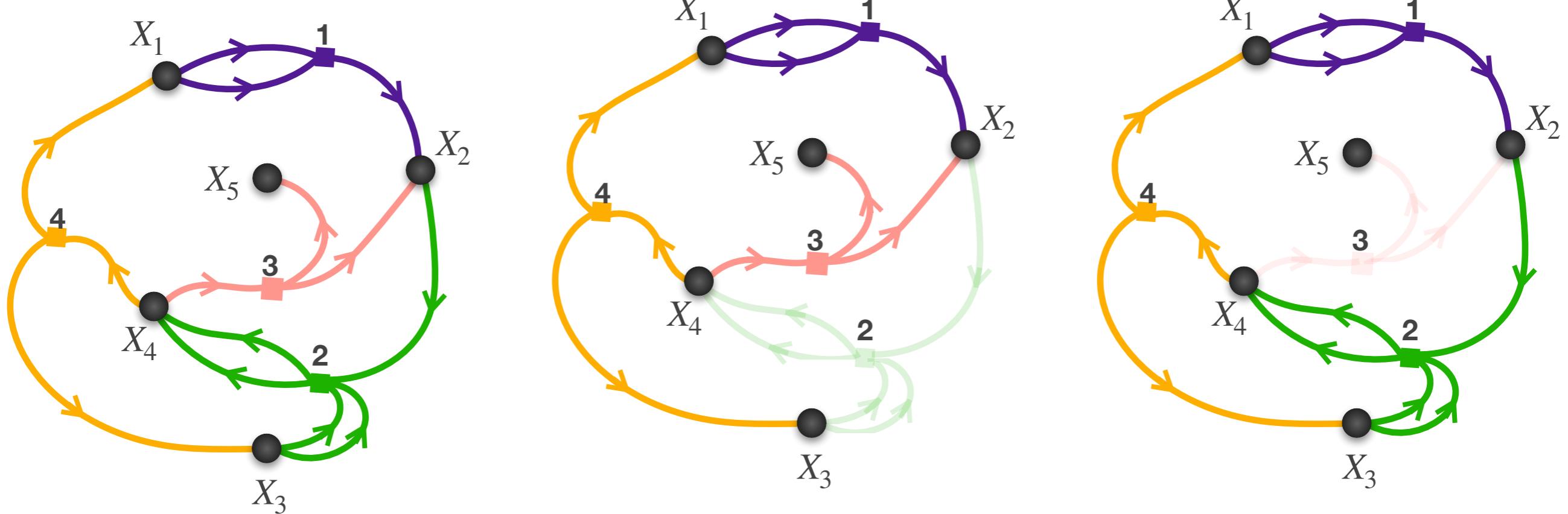
- Spectrum fully controls by the matrix $\tilde{\mathbb{T}}$
- They **share the same spectrum** up to the multiplicity of 0's and 1's eigenvalues
- Spectral Einstein relation

L^{rel} \iff Noise matrix

L^{ext} \iff Mobility

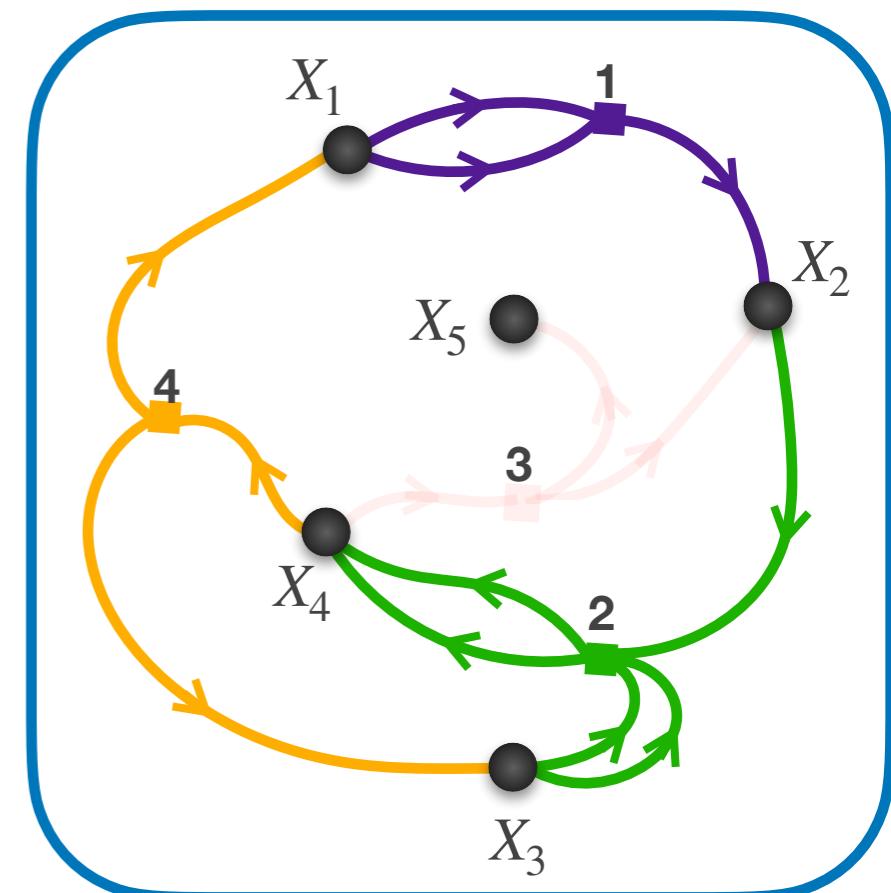
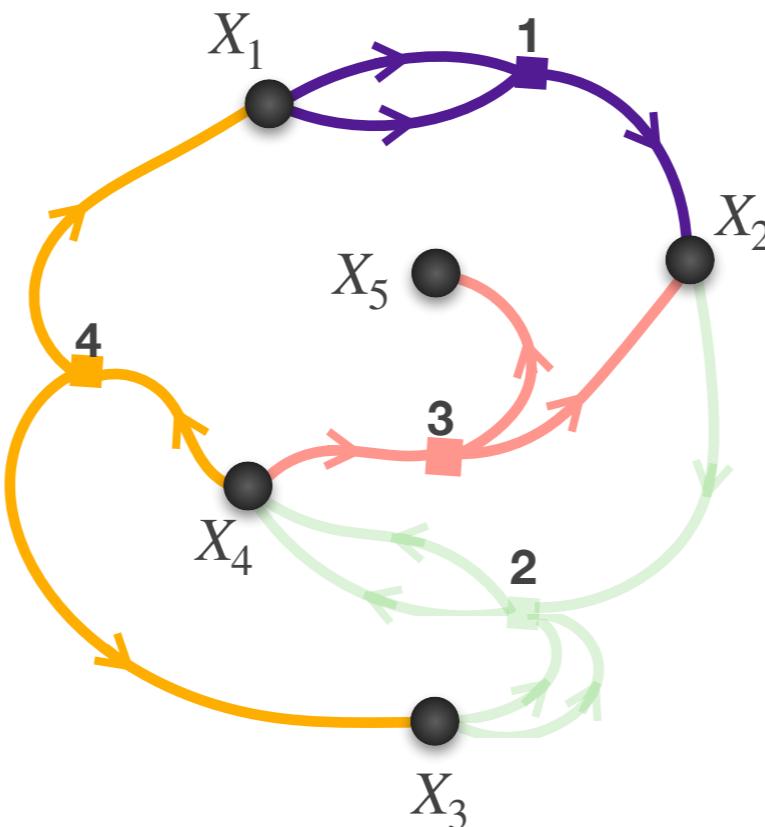
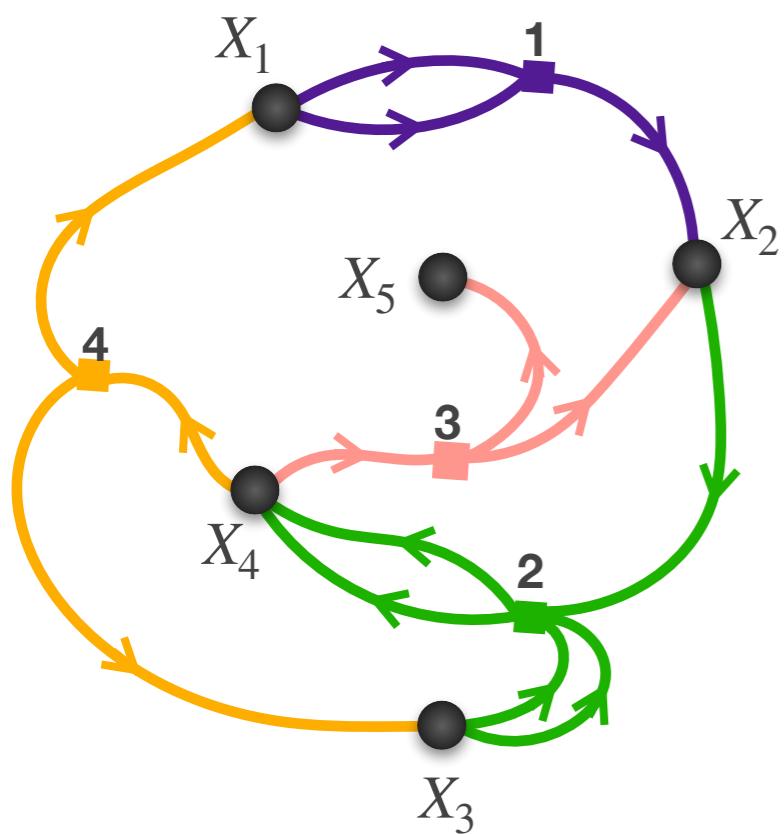
Beyond linear response

Response of the system to the suppression of a reaction?



Beyond linear response

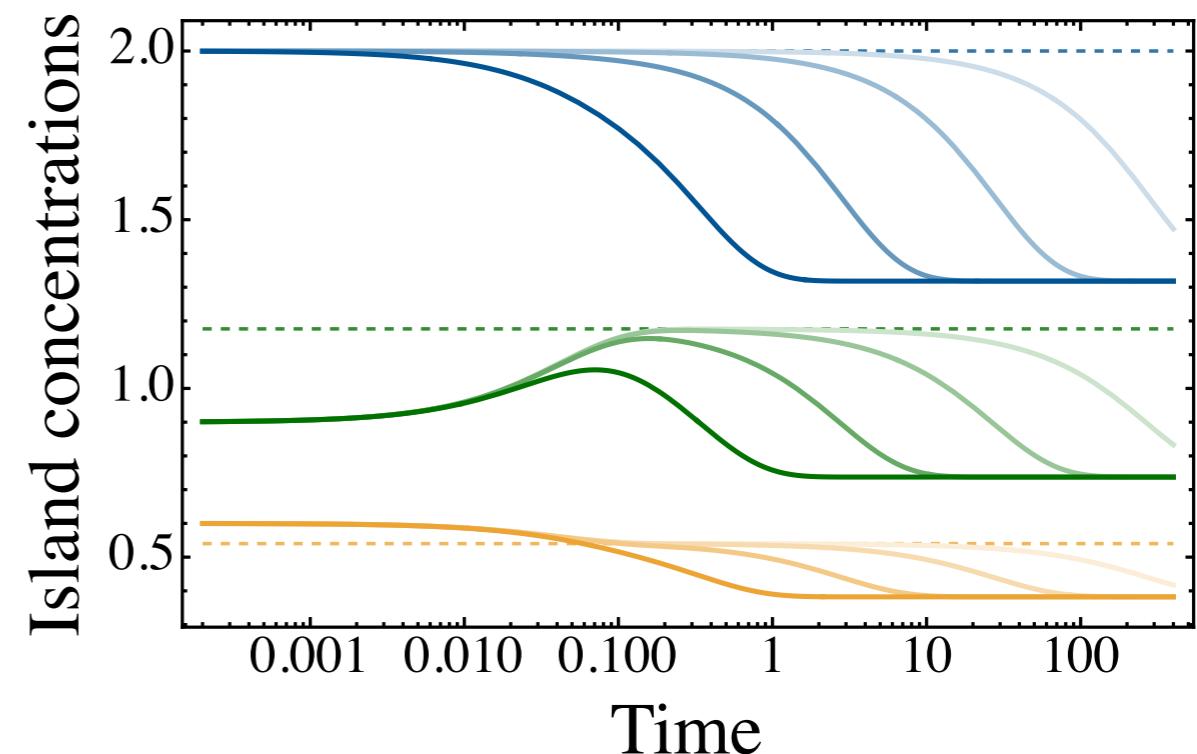
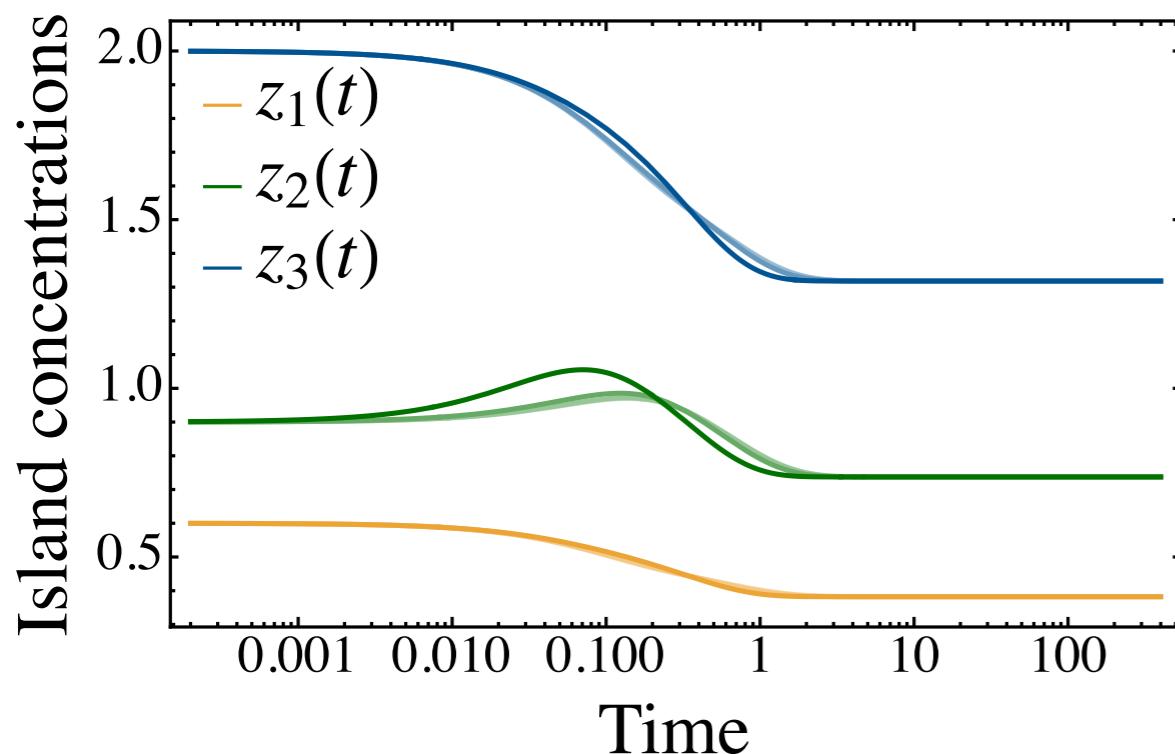
Response of the system to the suppression of a reaction?



Reaction 3 is a cocycle,
i.e. a mode of relaxation

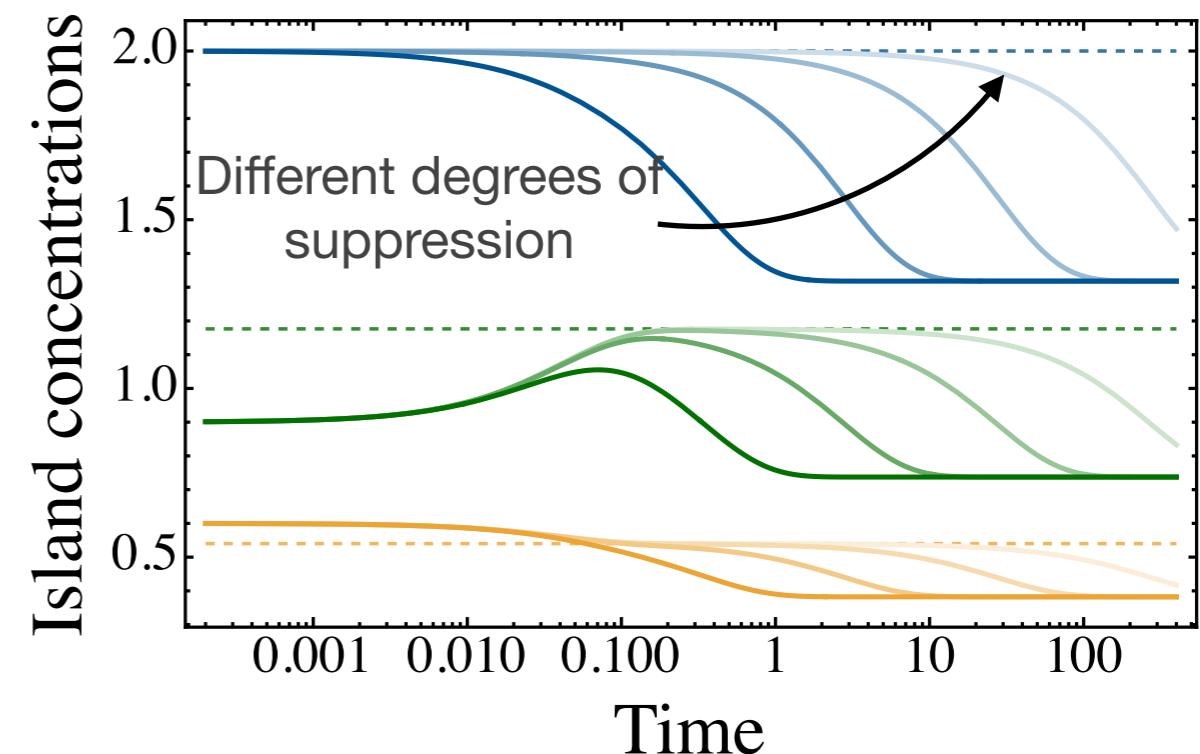
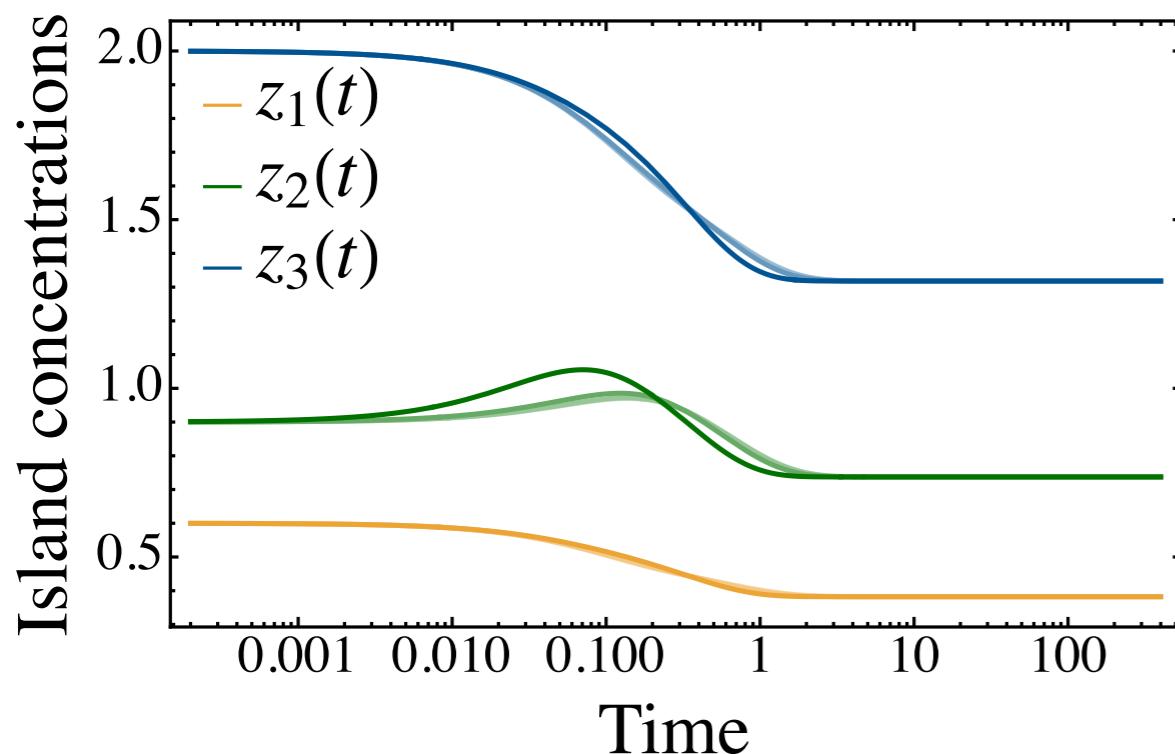
Beyond linear response

Emergent timescale in the relaxation by suppressing a cocycle



Beyond linear response

Emergent timescale in the relaxation by suppressing a cocycle



Dynamical control of complex CRNs?

Conclusions

Geometrically-informed decomposition of affinities and currents

Framework describing both relaxation and steady-state

Generalization to interacting CRNs using linear algebra

Application to linear and non-linear response

Conclusions

Geometrically-informed decomposition of affinities and currents

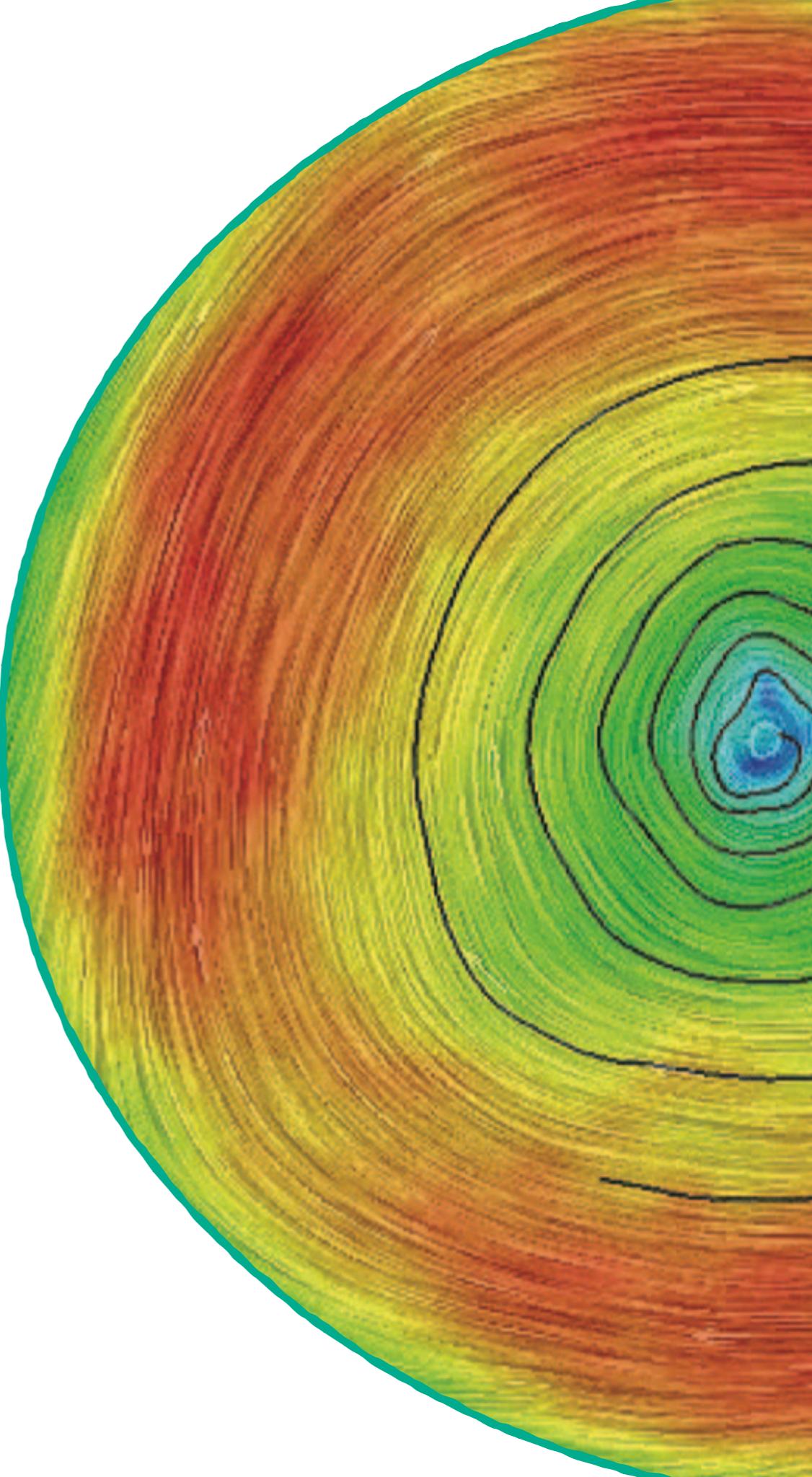
Framework describing both relaxation and steady-state

Generalization to interacting CRNs using linear algebra

Application to linear and non-linear response

Collaborators: Vivien Lecomte, Matteo Polettini

Thank you!



Algebraic interpretation

Stoichiometric matrix encodes the topology of the network

$$\mathbb{S} = \left(\begin{array}{c} \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \end{array} \right) \quad \longleftrightarrow \quad N \text{ species}$$
$$R \text{ reactions} \quad \longleftrightarrow \quad \left(\begin{array}{c} \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \end{array} \right)$$

The cocycles are a basis
for the coimage of the
stoichiometric matrix

$\text{Im } \mathbb{S}^T$

The cycles are a basis for
the kernel of the
stoichiometric matrix

$\text{Ker } \mathbb{S}$

Algebraic interpretation

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The cocycles are a basis
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$$\# \text{ cocycles} = M = \text{Rank } \mathbb{S}$$

The cycles are a basis for
the kernel of the
stoichiometric matrix

$$\# \text{ cycles} = R - M$$