### Topology protects long cycles in stochastic systems



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### How do long and stable timescales and rhythms emerge?

Given heterogeneity and randomness, how does robust function emerge?

Long time-scales such as circadian cycles or memory, persist despite consisting of many faster reactions

Can we predict such global dynamics analytically?





# Biological systems show many types of robust self-organization



Many parts acting together to produce more than the sum of each part

Similar dynamics all hinge on the edge state of a topological phase, that consists of many non-reciprocal cycles

### Living things show strong non-Hermiticity and dissipation which appears wasteful

Living systems exhibit many microscopic out-of-equilibrium transitions, e.g. by consuming fuel such as ATP or GTP

"Futile cycles" consume a lot of energy; ubiquitous in biology (metabolism, sensory systems, muscular contraction, protein synthesis)



Could these "wasteful" motifs contribute to system robustness?

# A topological mechanism for robust function in stochastic systems



Model for emergent oscillations



New physics and underlying topology



Non-reciprocity is a necessary condition

### A two-dimensional reaction space with four internal states



→ Phoshorylation → Dephosphorylation

Another process primes system for next transition, i.e. 4 internal states which we can label

- A: Pink phosphorylation
- B: Orange phosphorylation
- C: Pink dephosphorylation
- D: Orange dephosphorylation

Transitions between internal states of the same phosphorylation level

## Repeating this motif of four internal states allows the tiling of a lattice



### When large external transitions dominate, the system will stay on the edge



Edge current is robust to inaccessible or missing states

Random matrix theory: robustness to random perturbations

ET et al., Phys Rev X 2021



### Can describe emergent cycles in biological oscillators

24hr cycle in KaiC hexamers arises from sequential phoshorylation of T and S sites

KaiC monomers can take different conformations, e.g., with exposed or buried A-loops

Cohen & Golden, Microbio & Mol Bio Rev 2015

Given the large reaction space, why does a global and robust cycle emerge?



## A large phase space of many possible different reactions

Monomers can autophosphorylate or assemble into hexamers



Brettschneider et al., Mol Syst Biol 2010

Can take different conformations in same hexamer Han et al., Nat Comm 2023

With this large phase space of monomer transitions, why do they phosphorylate in a concerted way?



### Within the large space of transitions: we propose 4-state directed cycles



### Experimental evidence for this model

Specific internal states are needed for phosphorylation steps

- KaiA binding promotes phosphorylation Tomita et al., Science 2005
- KaiB binding promotes dephosphorylation Rust et al., Science 200
- KaiA only binds to exposed A-loop Kim et al., PNAS 2008

#### Difference in transition rates

- Phosphorylation reactions are faster Paijmans et al., PLoS comp bio 2017
- KaiB binding (and possibly its unbinding) are slower steps
  Abe et al., Science 2015, Kageyama et al., Mol. Cell 2006, Since et al., 2000 are al.

S phosphorylati







## Edge cycle reproduces observed dynamics with fewer fine-tuned parameters



#### Edge state produces cycles of

- all T phosphorylation,
- all S phosphorylation,
- all T dephosphorylation,
- all S dephosphorylation

Zheng and ET, Nat Comm 2024

k<sub>ps</sub> Increasing phosphorylation к<sub>dps</sub> k<sub>ps</sub> ƙ<sub>dps</sub> k<sub>ps</sub> Active Inactive state state

In MWC paradigm: ordering put in by hand; intermediate transitions assumed negligible

van Zon et al., PNAS 2007

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## To see if the system is in a topological state, analyze ${\cal W}\,$ the transition matrix

Systems described by the Master equation  $\frac{d}{dt}p = \mathcal{W}p$ 

Lattice description allows for calculation of the Berry connection, which is calculated from the eigenvectors of  $\ensuremath{\mathcal{W}}$ 



Integration over reciprocal space gives 0 (trivial phase) or  $\pi$  (topological phase)

#### First developed for quantum Hamiltonians

$$-i\frac{d}{dt}\psi(t) = \mathcal{H}\psi(t),$$

Identical to the Master equation up to a *i* 

Nat Rev Phys 2019

### Verified using stochastic simulations



Gillespie algorithm,  $\gamma_{ex} = 10^3 \gamma_{in}$ 



### Edge flows drive macroscopic re-organization in driven colloids



We identify and analyze edge currents around voids and clusters

Building on theory first developed in Dasbiswas et al., PNAS 2018 17

## Edge flows create stress patterns that drive coarsening on different time scales



We obtain physical properties of edge flows



Coarsening happens on much longer timescale for voids as compared to clusters

Nelson, Lobmeyer, Biswal and ET, arXiv 2024

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## Out-of-equilibrium living systems require new theoretical tools

Stochastic systems are described by Master equation  $\frac{d}{dt} \boldsymbol{p} = \mathcal{W} \boldsymbol{p}$ 

Out-of-equilibrium or non-reciprocal transitions make  $\mathcal{W}$  non-Hermitian

Non-reciprocal interactions are legion in living and active systems, e.g. between predators and their prey

New properties in stochastic topological systems



Nelson and ET, Phys Rev B 2024

## Quantum and stochastic networks are mapped only in the bulk

Network has the same operator under periodic boundary conditions

Since 
$$W = A - D$$
,  $A_{ij} = \langle i | j \rangle$ , transition rate from state  $p_j$  to  $p_i$   
 $D_{ij} = \delta_{ij} \sum_k \langle k | i \rangle$ , a diagonal matrix

Then  ${\mathcal D}$  is proportional to the identity, so the spectrum is just shifted





State of interest is also different: zero mode vs steady-state

## Stochastic systems break the bulk boundary correspondence

Quantum and stochastic networks have same operator and topological invariant under periodic boundaries, or in the bulk

With open boundaries, longest-lived states in each system look different



Despite same bulk topological invariant, reciprocal stochastic systems do not have edge states even when the quantum one does



More generally: unlike in quantum systems, stochastic systems require non-Hermiticity to have edge states

We can prove this for any geometry and dimension

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A topological mechanism for robust function in biological networks

1. Using simple repeated motifs such as phosphorylation cycles, we propose a topological framework that predicts the emergence of robust global dynamics, such as a global clock.

ET et al., Phys Rev X 2021

2. We demonstrate the relevant biophysical mechanisms in KaiC which regulates the circadian rhythm.

Zheng and ET, Nat Comm 2024

3. Topological edge flows can drive macroscopic re-organization in colloids.

Nelson, Lobmeyer, Biswal and ET, arXiv 2024

Invited review: Agudo-Canalejo and ET, arXiv 2024

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Student and postdoc positions available!