

Nonequilibrium Organization and Fluctuations in Active Interfaces and Polymers

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Pre-PhD Seminar

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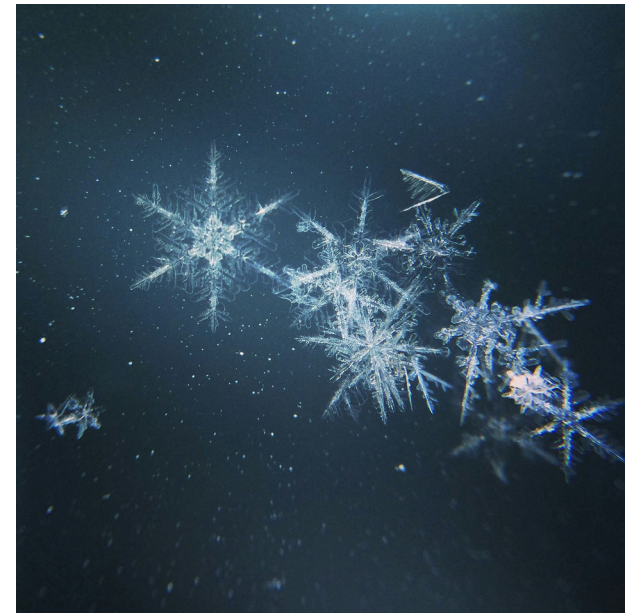
Non-equilibrium Organization and Fluctuations



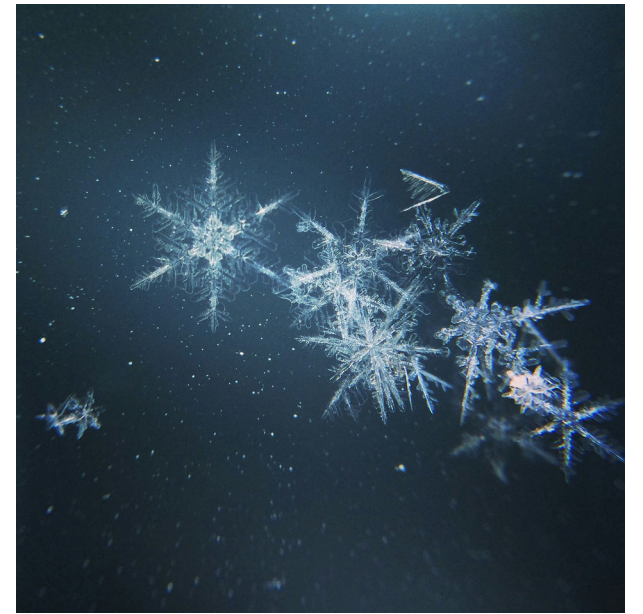
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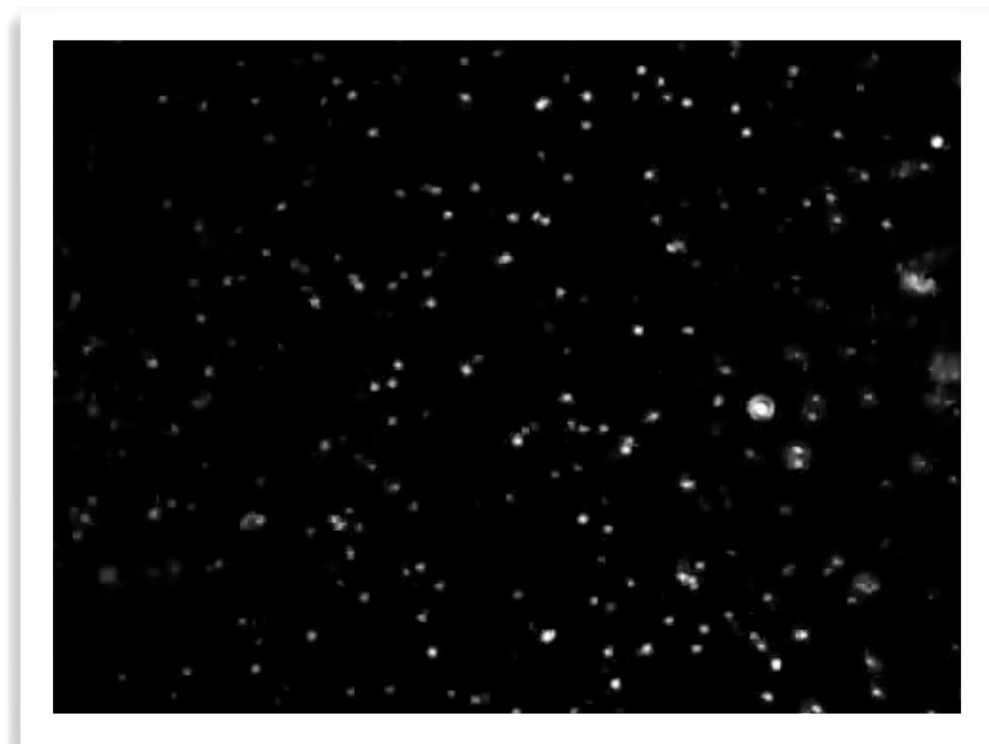
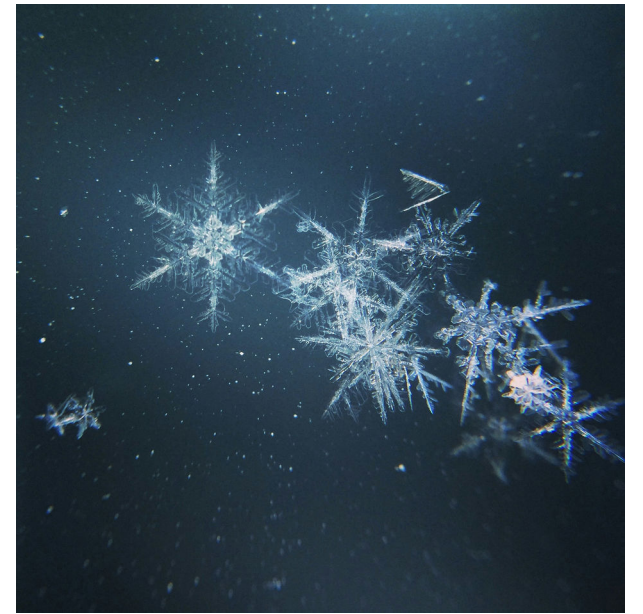
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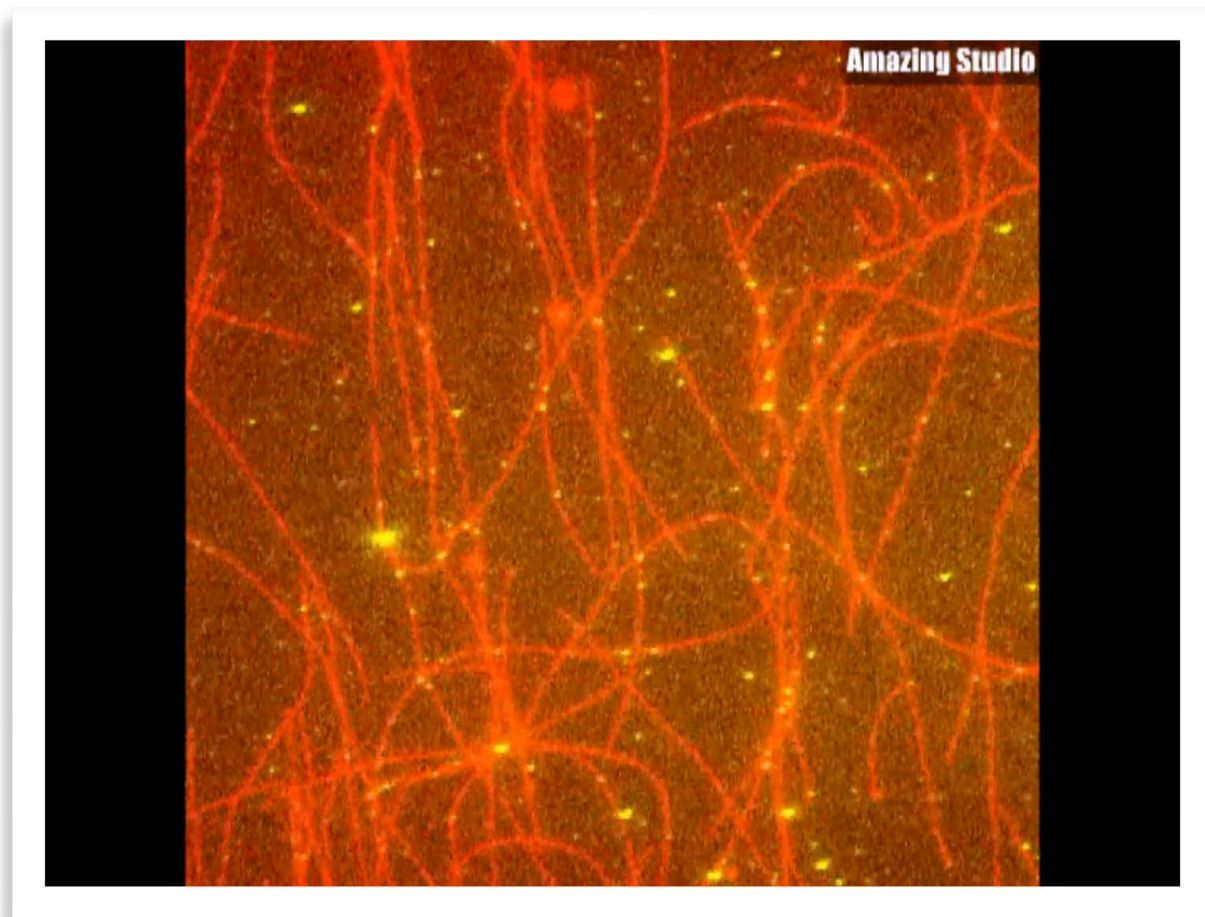
Non-equilibrium Organization and Fluctuations



Non-equilibrium Organization and Fluctuations



Active Systems as Non-equilibrium systems



GFP tagged Kinesin walking on TRITC labeled microtubules.

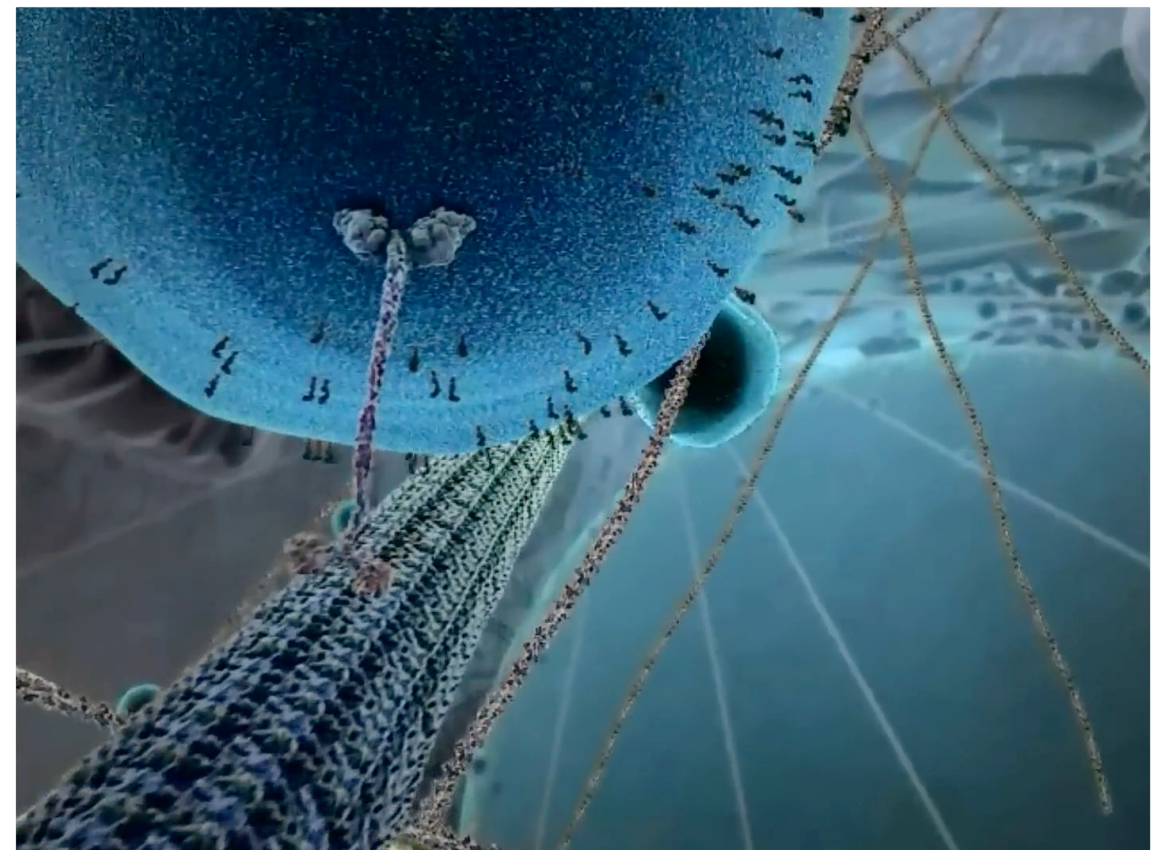
(Video courtesy of Cytopirate Labs, an educational community biolab).

Active Systems as Non-equilibrium systems



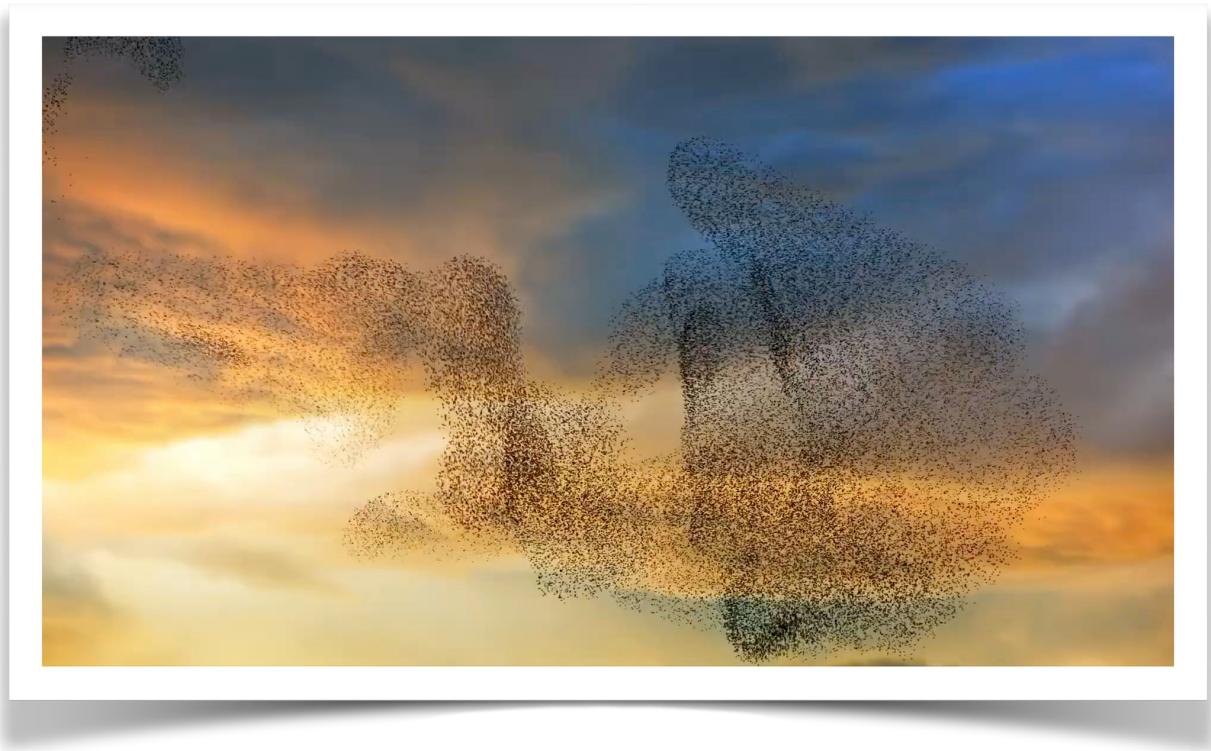
GFP tagged Kinesin walking on TRITC labeled microtubules.

(Video courtesy of Cytopirate Labs, an educational community biolab).



Kinesin protein walking on a microtubule in a cell

Collective behaviour in active matter



A flock of birds

Collective behaviour in active matter



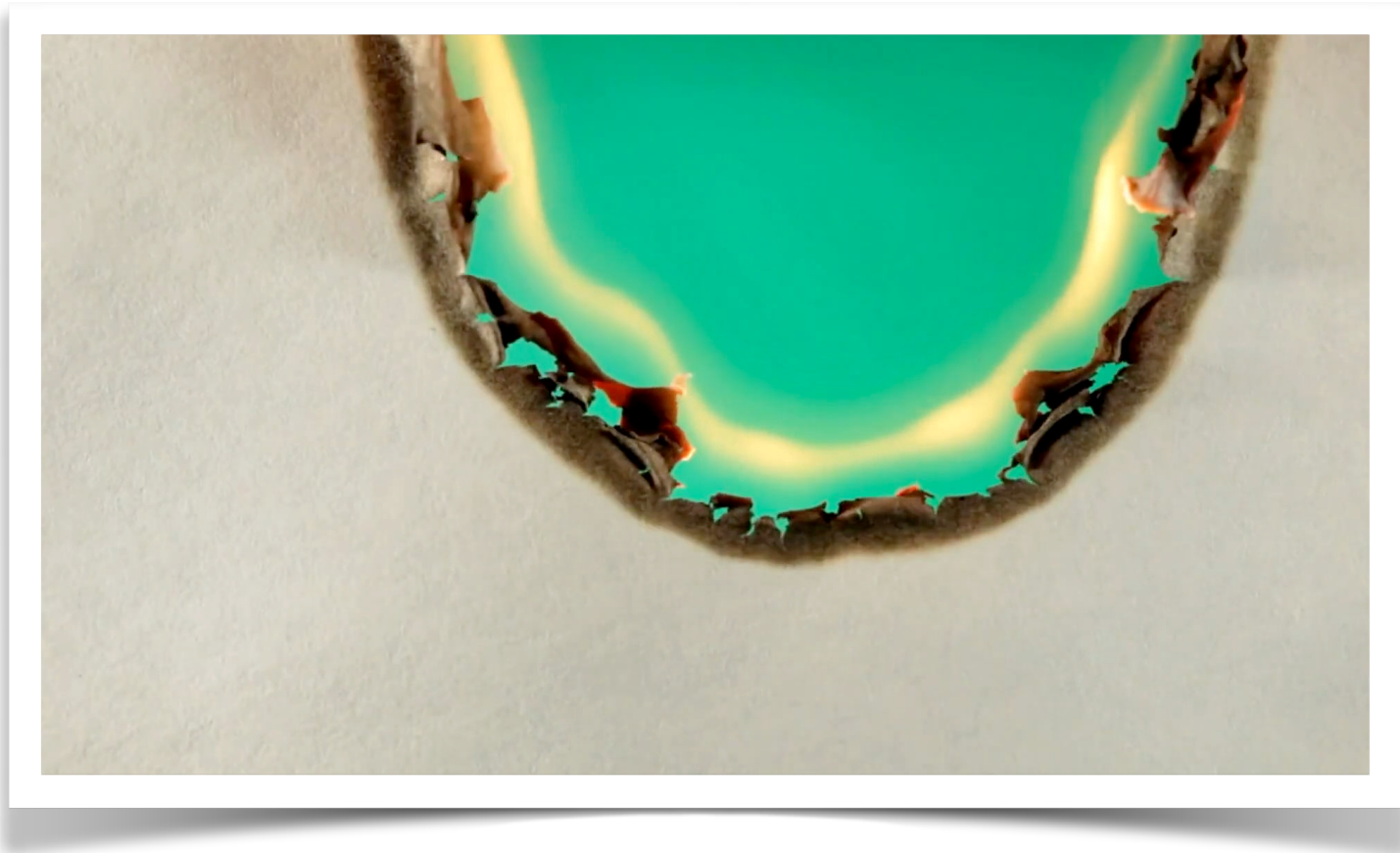
A flock of birds

Jeremie Palacci et al. ,Living Crystals of Light-Activated Colloidal Surfers.Science339,936-940(2013).DOI:10.1126/science.1230020

Living Crystals of Light-Activated Colloidal Surfers

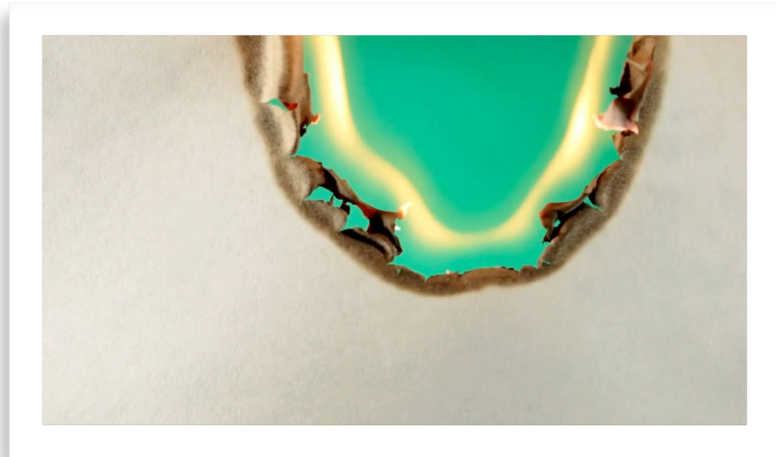


Growth Vis-à-Vis: The Non-Active vs. Active

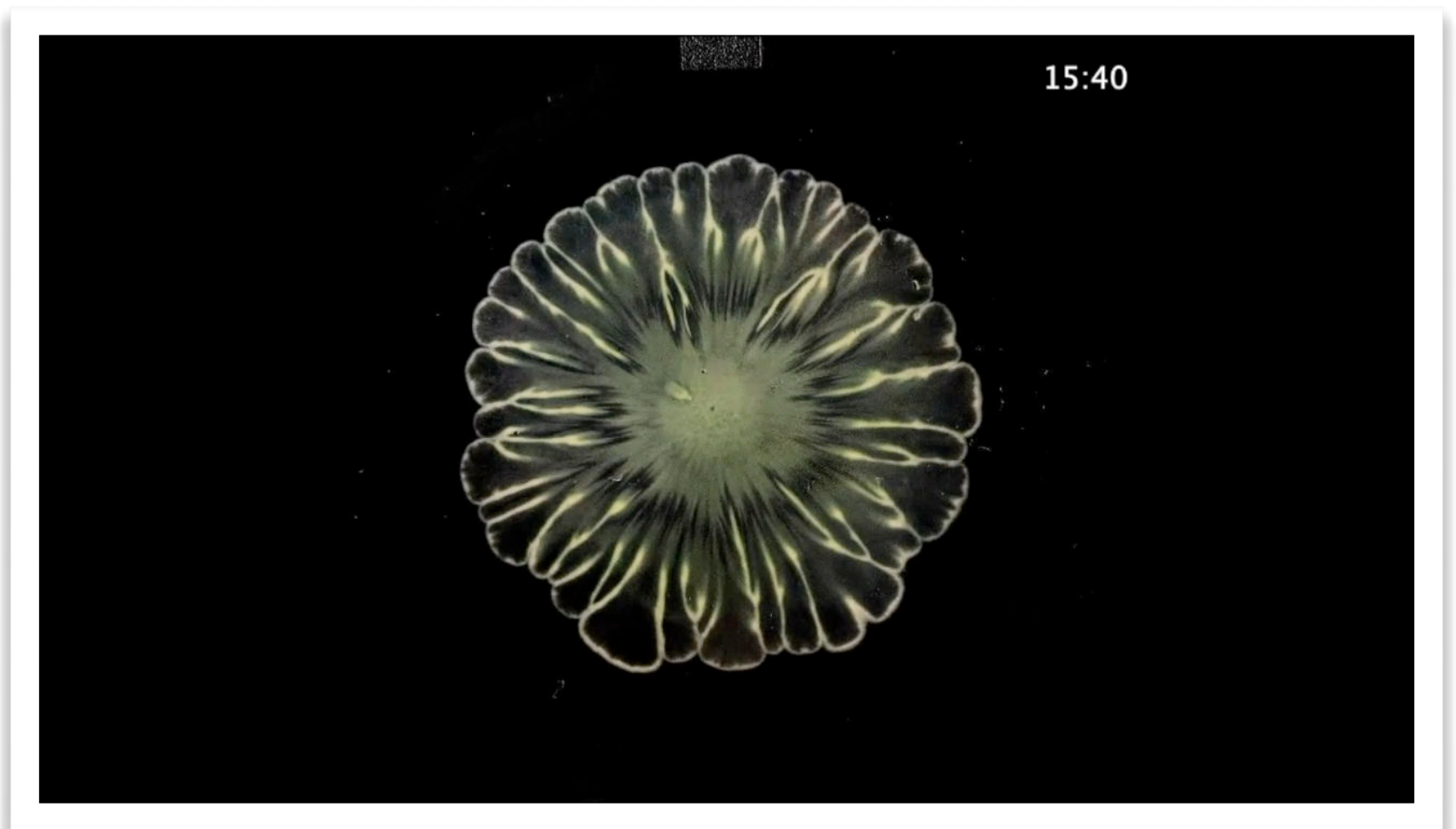


A non-living KPZ interface.

Growth Vis-à-Vis: The Non-Active vs. Active



A non-living KPZ interface.

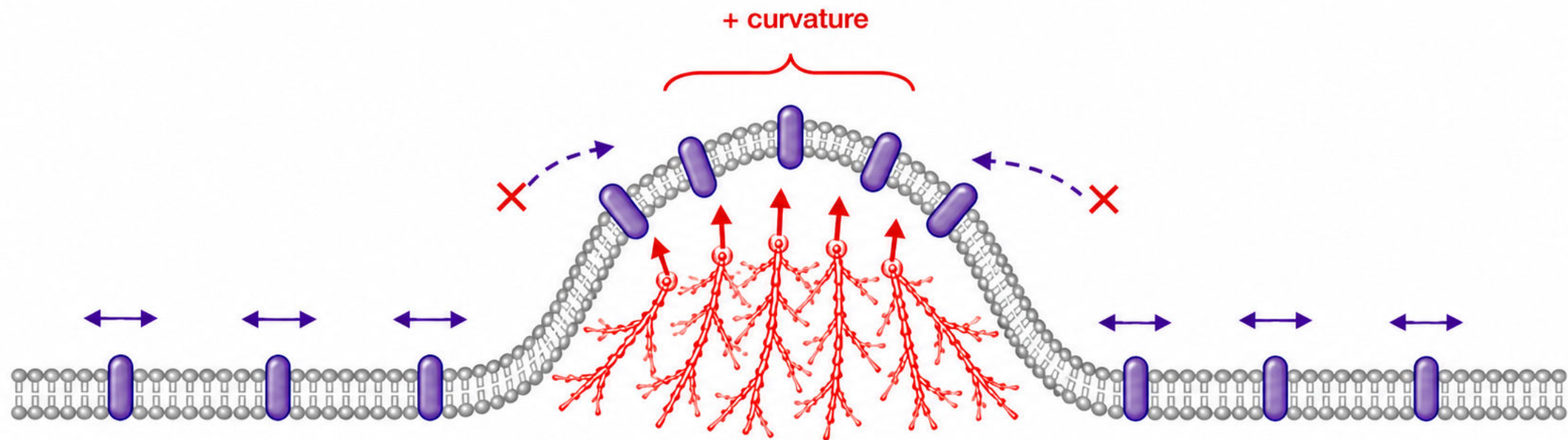


Time-lapse video of flower-like patterns that emerge when *E. coli* and *A. baylyi* grow over a 24-hr. period.

Part A

Active particles interacting with an interface

A model for an active interface



Membrane
bound proteins



Actin



Actin polymerization



Random diffusion
(on flat membrane)



Doesn't like
+ curvature

A model for an active interface

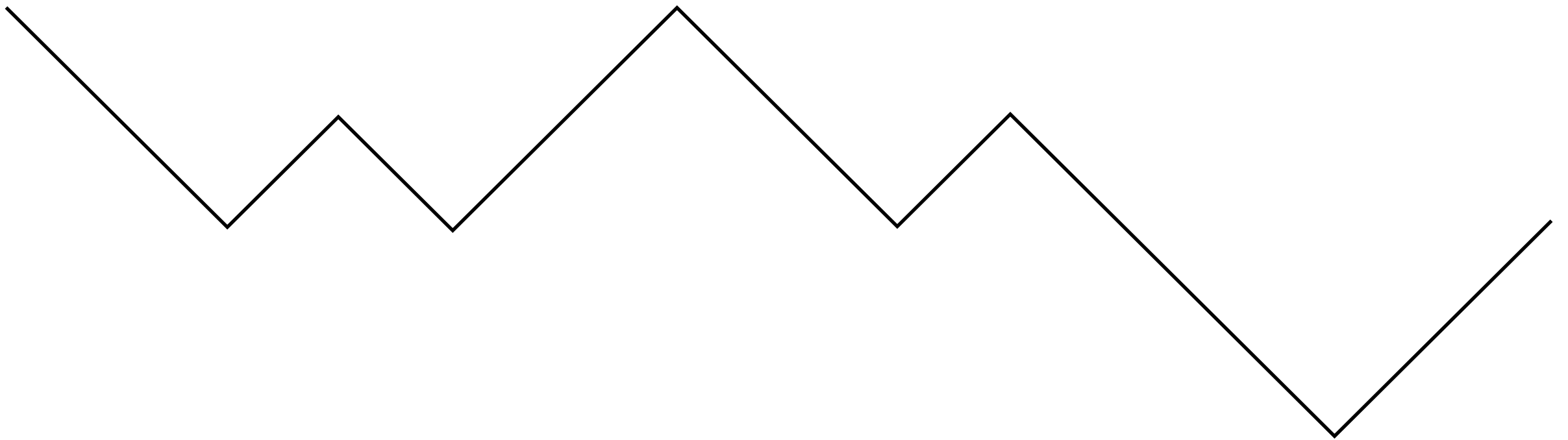
Model Setup

- **The Single-Step Model:** We model the interface as a 1D lattice where the local height difference between adjacent sites is strictly restricted to slopes of $\tau = \pm 1$ (a zig-zag membrane).

A model for an active interface

Model Setup

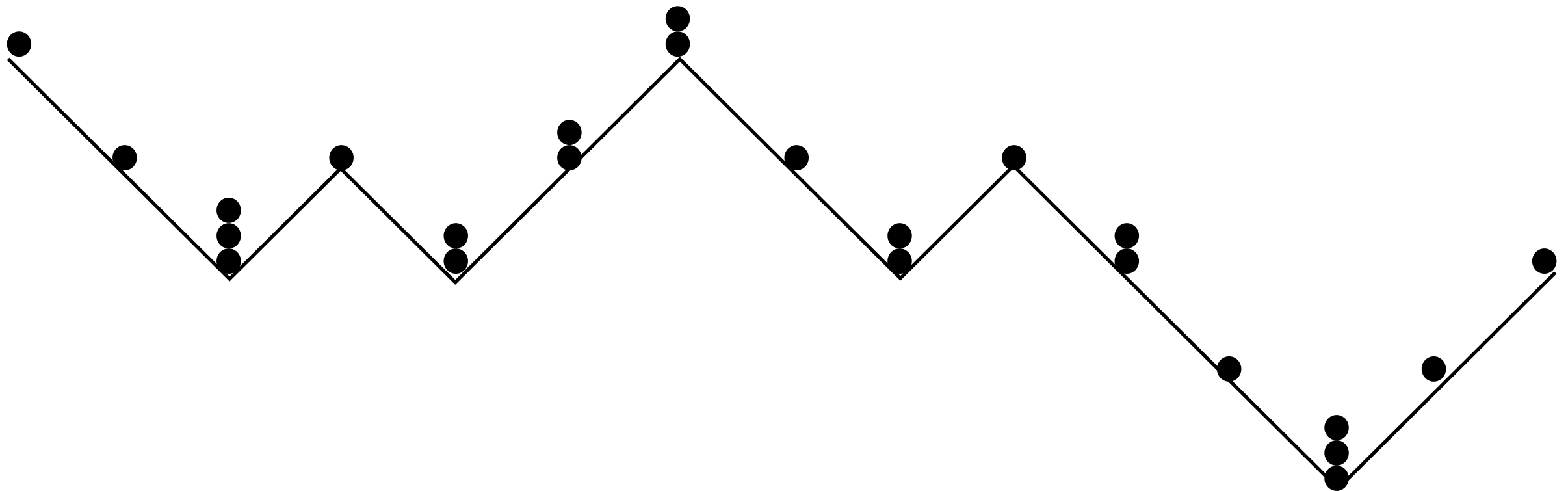
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A model for an active interface

Model Setup

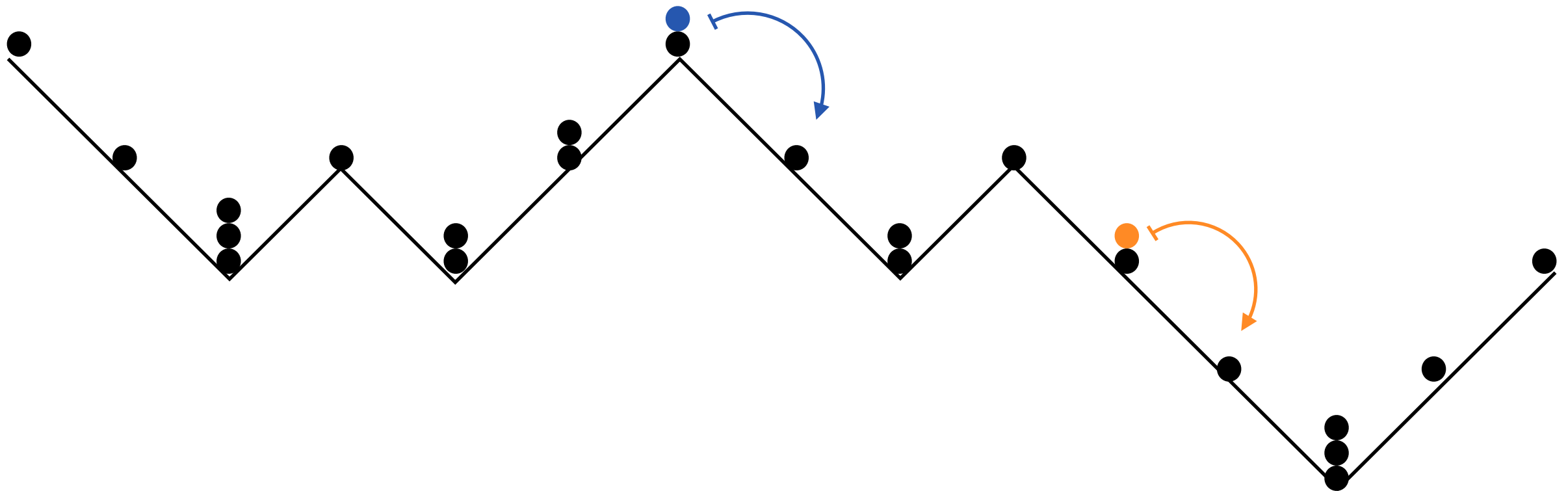
- **Particles Residing on Membrane:** We introduce particles into this environment. These particles occupy the nodes (the joints between the slopes) along the membrane.



Particle Kinetics

Moving on the Lattice

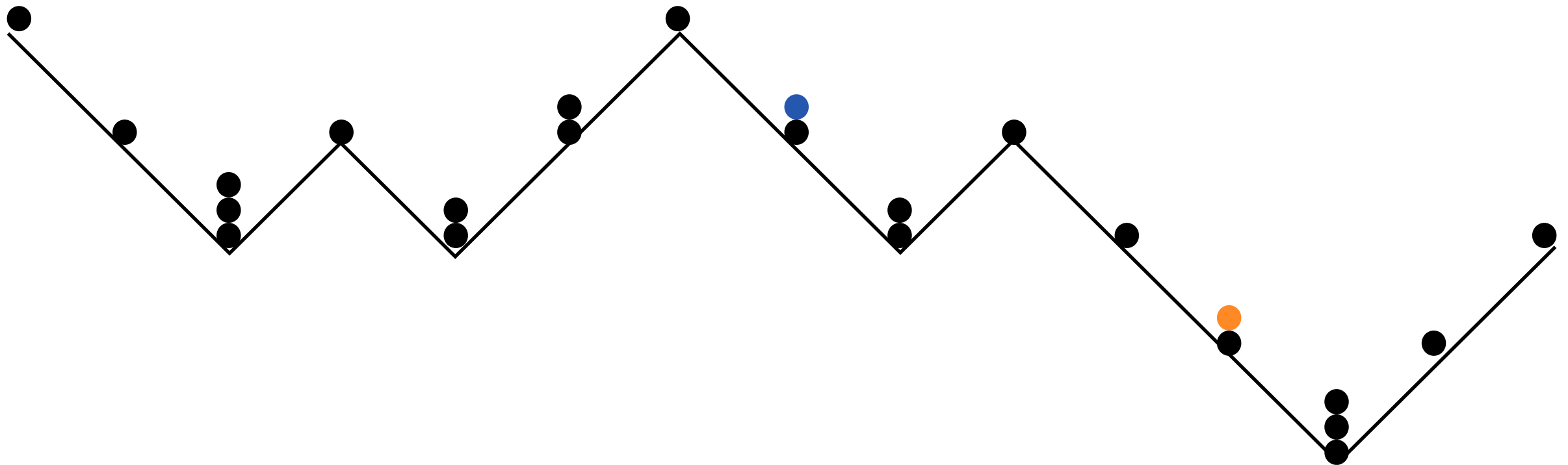
- **Topology-Dependent Hopping:** Particles do not just move blindly; they sense the local slopes. They hop to right adjacent nodes with a base rate $q^R \equiv q^-$ (and $q^L \equiv q^+$), but preferentially slide down slopes due to local "gravity".



Particle Kinetics

Moving on the Lattice

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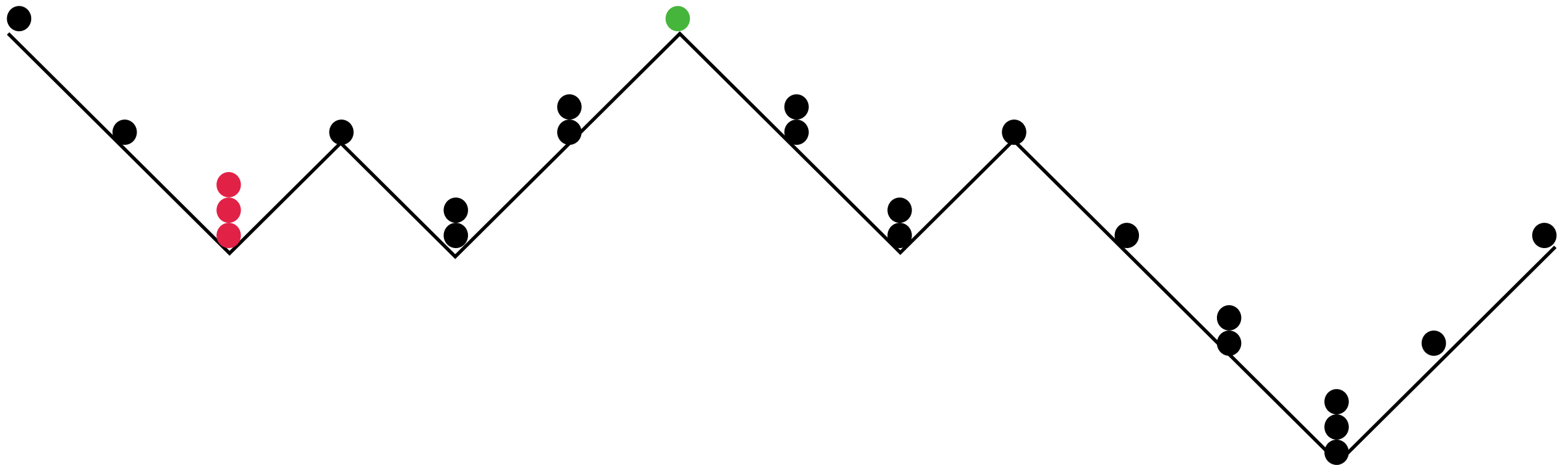


$$q^{\pm} = q_0 \left(1 \pm \frac{\gamma}{2} \nabla h \right)$$

Membrane Kinetics

EW vs KPZ vs Active

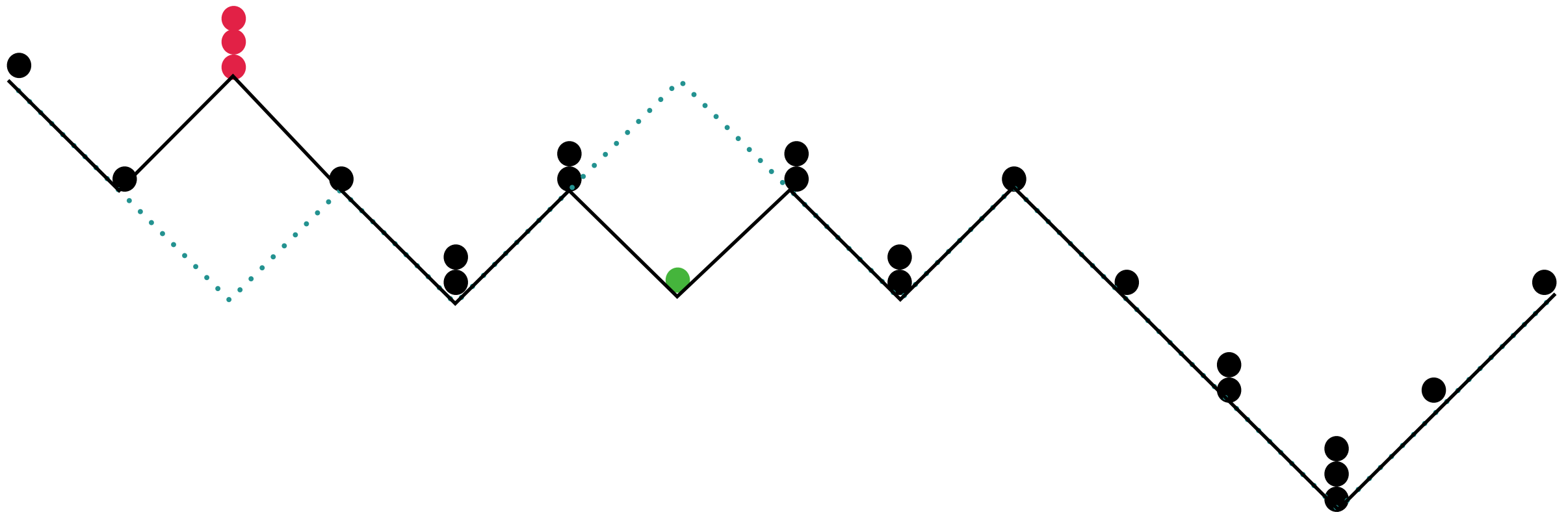
- **The Local Flips:** The membrane evolves through localized inversions. A valley ($\vee$) can invert into a peak ($\wedge$) with probability p^+ , and a peak can invert into a valley with probability p^- .



Membrane Kinetics

EW vs KPZ vs Active

- **The Local Flips:** The membrane evolves through localized inversions. A valley (V) can invert into a peak ($\wedge$) with probability p^+ , and a peak can invert into a valley with probability p^- .



Membrane Kinetics

EW vs KPZ vs Active

- **EW:** $p^+ = p^-$
- **KPZ:** $p^+ \neq p^-$
- **Active:** $p_n^\pm = f(n, \nabla^2 h)$

p^+ is the rate at which
the $\vee$ is converted to a $\wedge$.

Membrane Kinetics

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$$p_n^\pm = u \frac{e^{\pm n\beta}}{e^{n\beta} + e^{-n\beta}}$$

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$$p_n^\pm = u \frac{e^{\pm n\beta}}{e^{n\beta} + e^{-n\beta}}$$

$$p_{\{n_1, n_2\}}^\pm = u \frac{e^{\pm(n_1\beta_1 + n_2\beta_2)}}{e^{(n_1\beta_1 + n_2\beta_2)} + e^{-(n_1\beta_1 + n_2\beta_2)}}$$

Single Species Organization

Competition of parameters

- **The Two parameters:** β and γ are the two parameters in active membrane.

- β defines the activity generated by particles on membrane:

$$p_n^{\pm} = u \frac{e^{\pm n\beta}}{e^{n\beta} + e^{-n\beta}}$$

- γ defines the gravity in the system for the particles:

$$q^{\pm} = q_0 \left(1 \pm \frac{\gamma}{2} \nabla h \right)$$

Single Species Organization

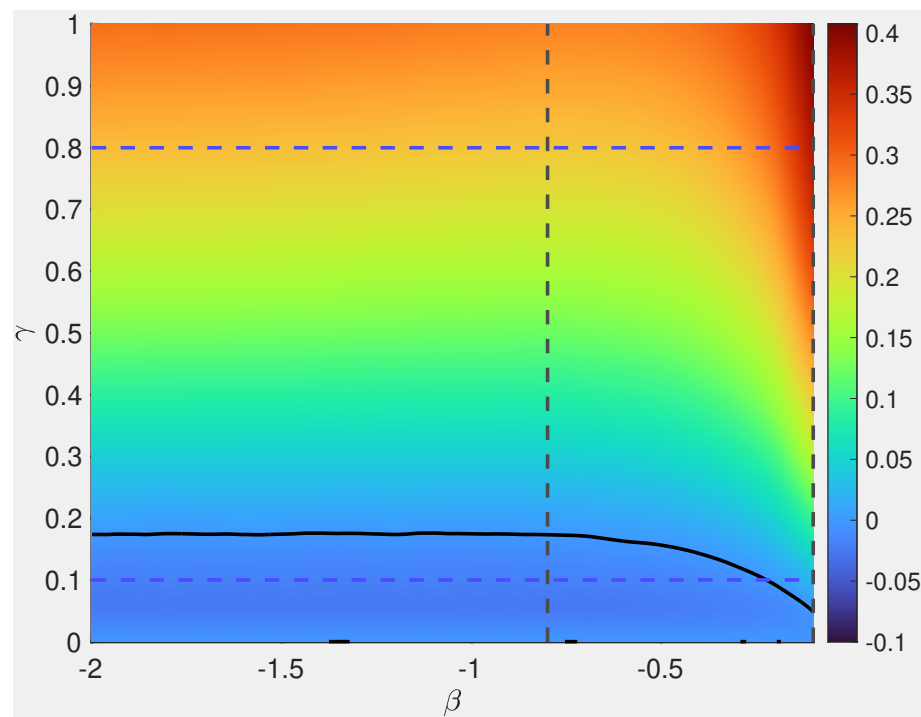
Competition of parameters

- $\beta < 0$: Pulls up the membrane
- $\gamma > 0$: particle falls in gravity downwards
 - ▶ Particles gets collected due to gravity
 - ▶ As particles gets collected more and more the valley becomes more and more unstable due to activity and eventually converted to hill.
 - ▶ Particles collected found themselves on hill which is an unstable situation due to gravity and local microcluster breaks as the particle moves to find a new valley.

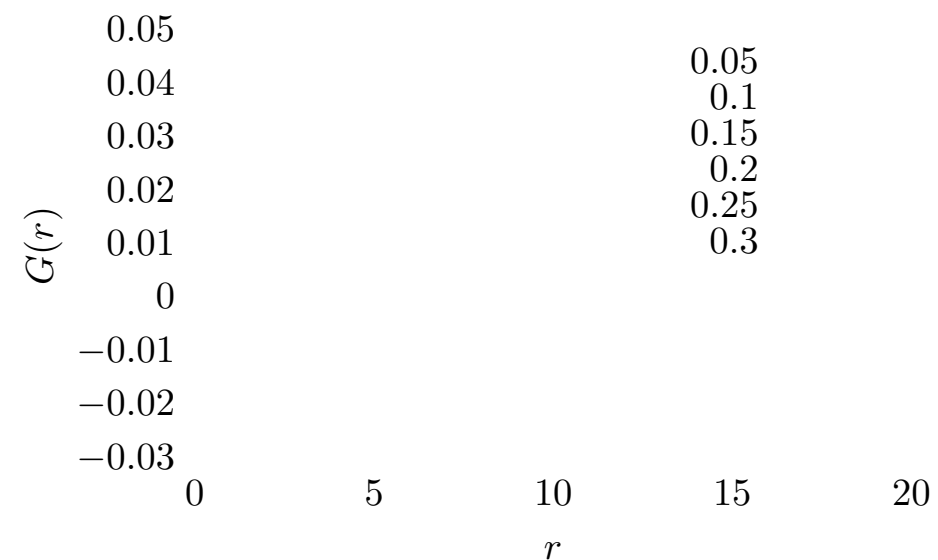
Competition

Anticorrelation, Zero-Correlation and Correlation Regime

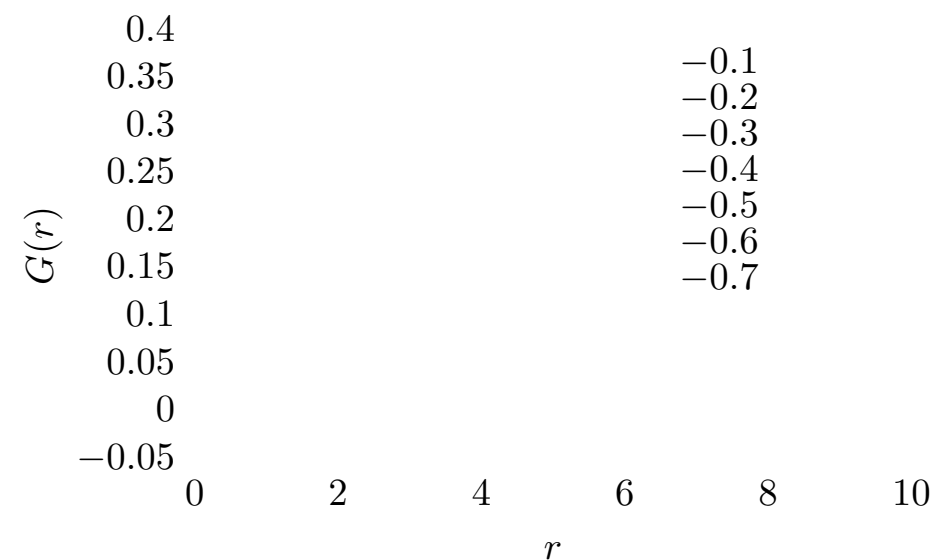
- $$G(r) = \frac{\langle s_i s_{i+r} \rangle - \langle s_i \rangle^2}{\langle s_i^2 \rangle - \langle s_i \rangle^2}$$



- $G(1)$ in β and γ phase-space



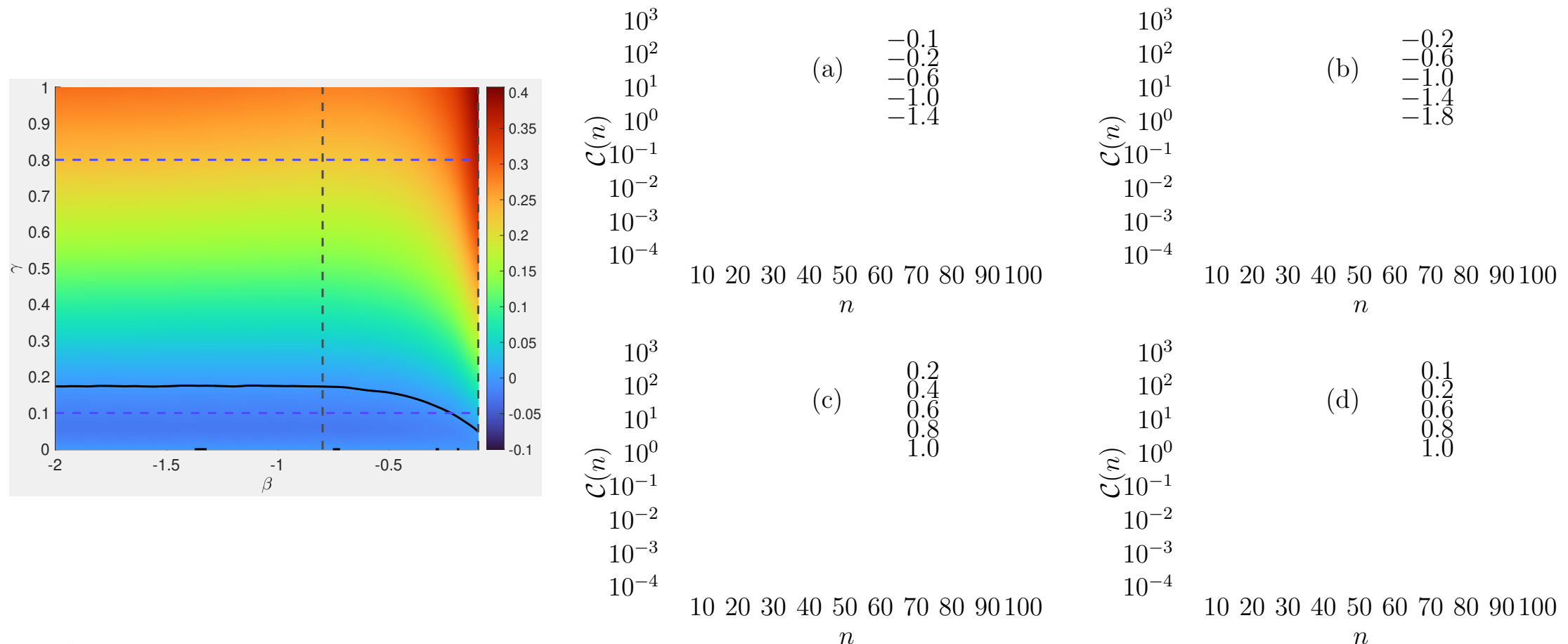
$\beta = -0.8$ and varying γ



$\gamma = 0.8$ and varying β

Cluster Distribution Function

- A cluster is defined as a contiguous group of particles such that empty sites on both ends bound the group.



- $\triangle$ is for zero-gravity case; $\bigcirc$ is for passive case (zero activity).
- [(a) $\gamma = 0.1$; (b) $\gamma = 0.8$] (β variable);
- [(c) $\beta = -0.1$; (d) $\beta = -0.8$] (γ variable)

Two types of active species

4 Cases

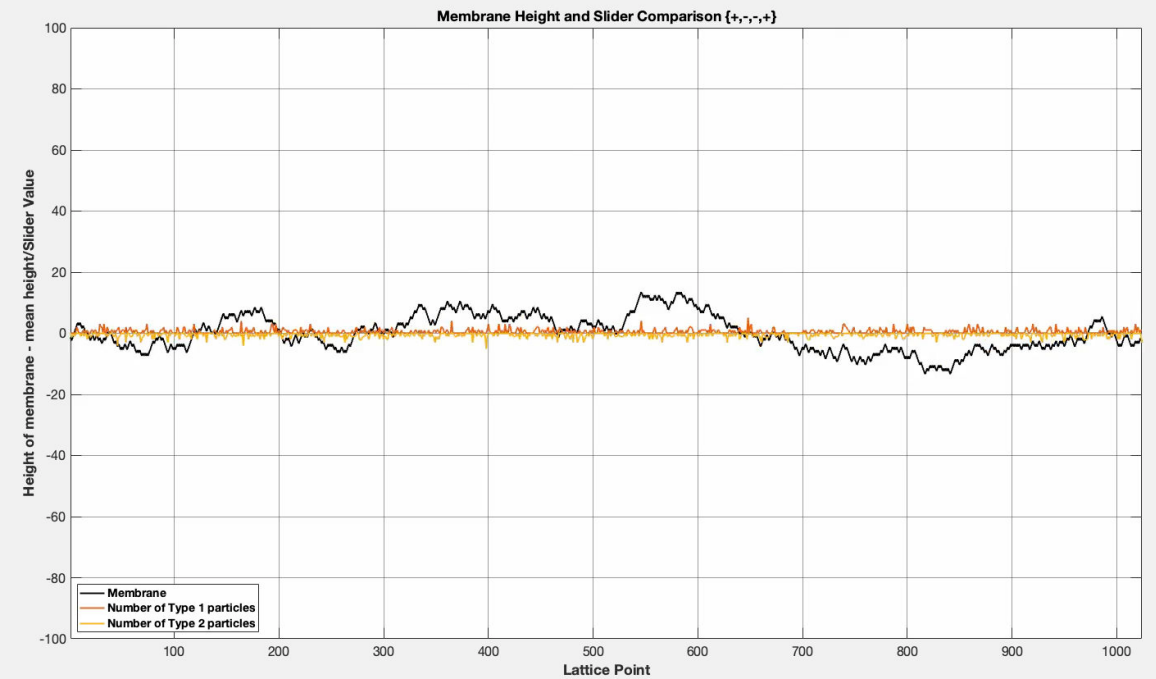
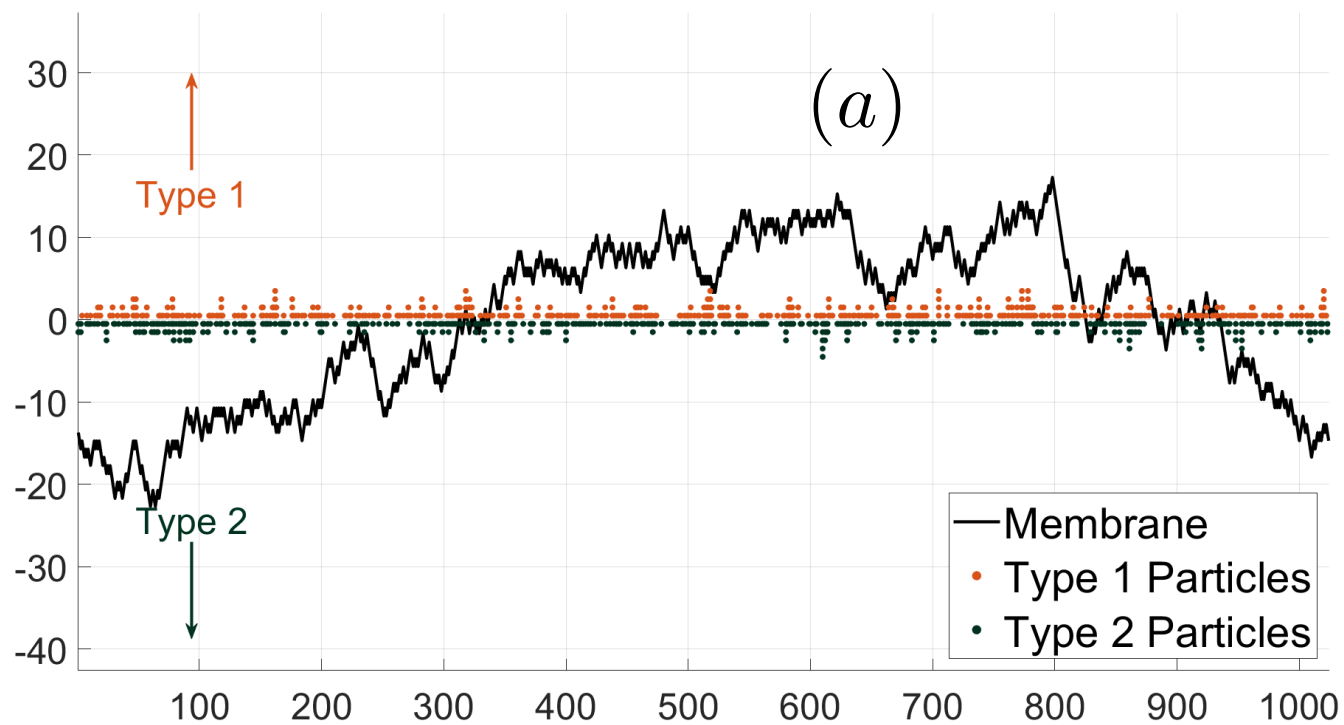
- **Sign Convention:** Gravity and activity acting in downward direction is positive.

Cases\Symmetries	Reversal	Exchange	Both
$\{+,+,-,+\}$	$\{-,-,+,-\}$	$\{-,+,+,+\}$	$\{+,-,-,-\}$
$\{+,+,-,-\}$	$\{-,-,+,+\}$	C(2)	C(1)
$\{+,-,-,+\}$	$\{-,+,+,-\}$	C(2)	C(1)
$\{-,-,-,+\}$	$\{+,+,+,-\}$	$\{-,+,+,-\}$	$\{+,-,+,+\}$
Trivial cases			
$\{-,-,-,-\}$	$\{+,+,+,+\}$	C(1)	C(2)
$\{+,-,+,-\}$	$\{-,+,+,-\}$	C(1)	C(2)

Signs of $\{\beta_1, \gamma_1, \beta_2, \gamma_2\}$, whereas we kept $|\beta_i| = 2$ & $|\gamma_i| = 1$

Two types of particles

$\{+,-,-,+\}$



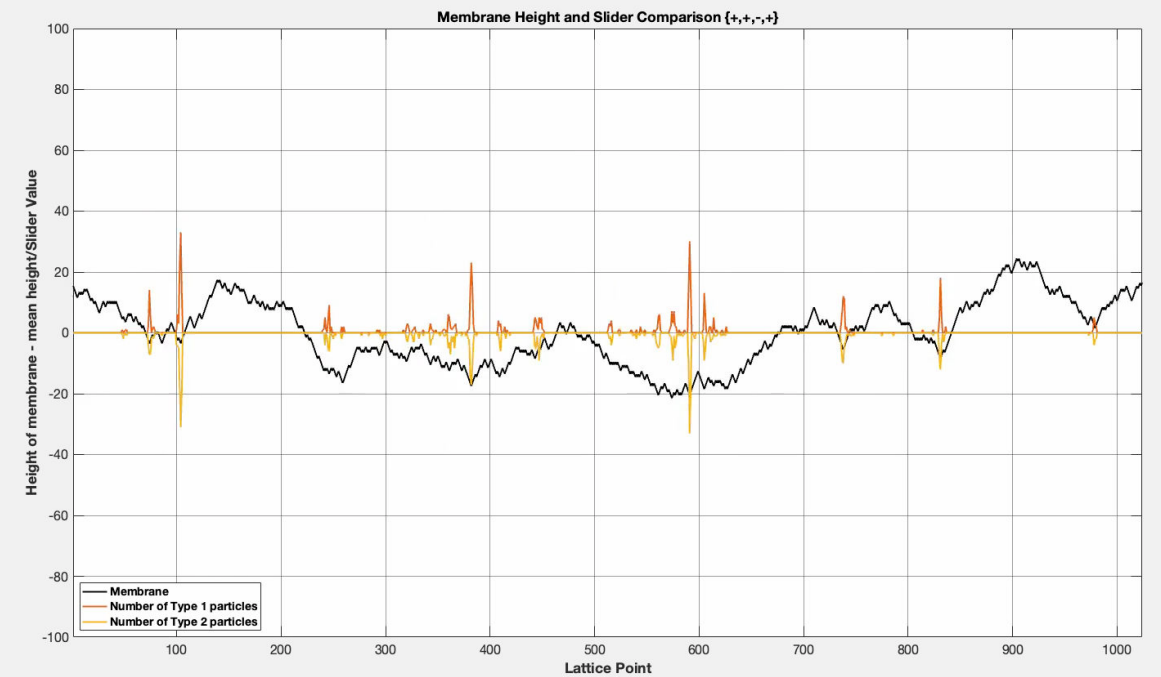
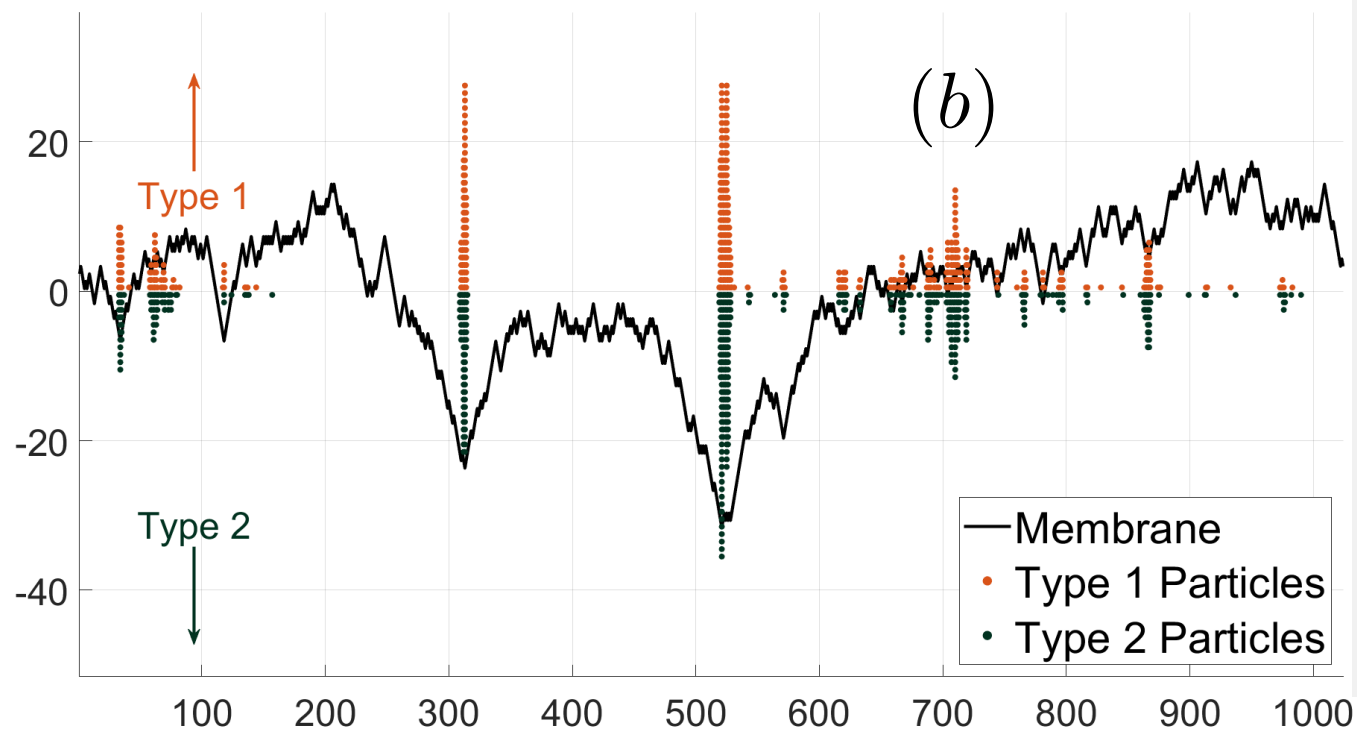
Type 1: Pulling membrane up and like valleys

Type 2: Pushing membrane down and like hills

Becomes like a single type of particle picture.

Two types of particles

$\{+,+,-,+\}$



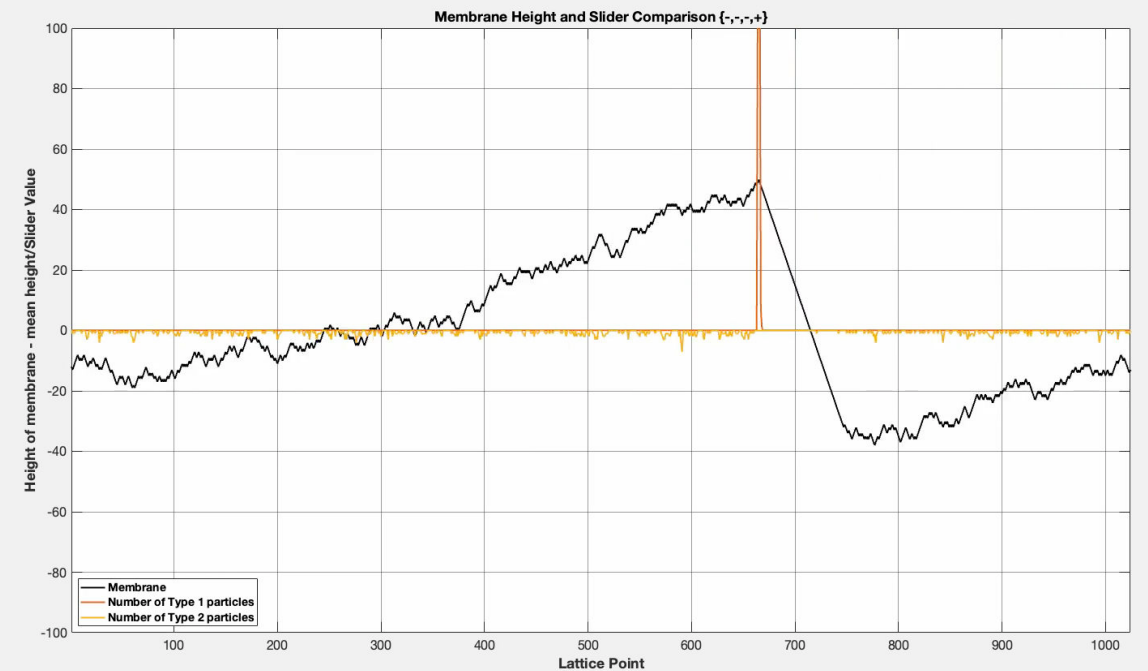
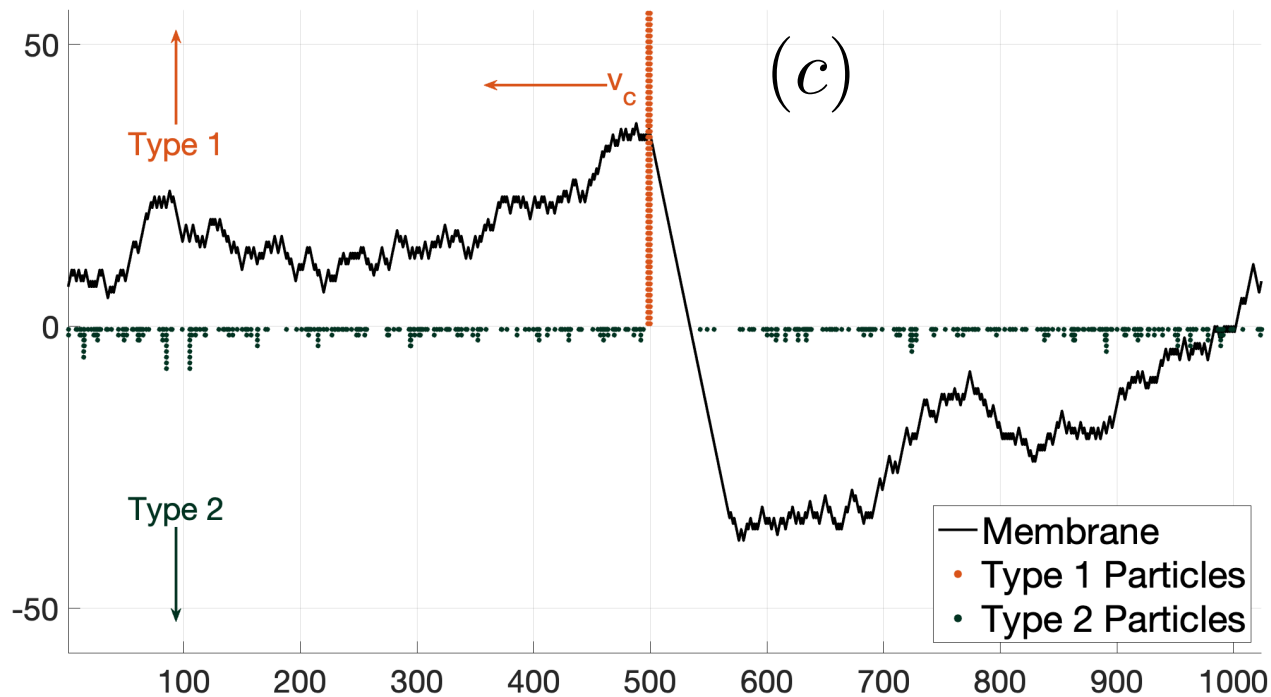
Type 1: Pulling membrane up and like hills

Type 2: Pushing membrane down and like hills

Becomes like a passive slider case, particles always stays balanced

Two types of particles

$\{-,-,-,+\}$



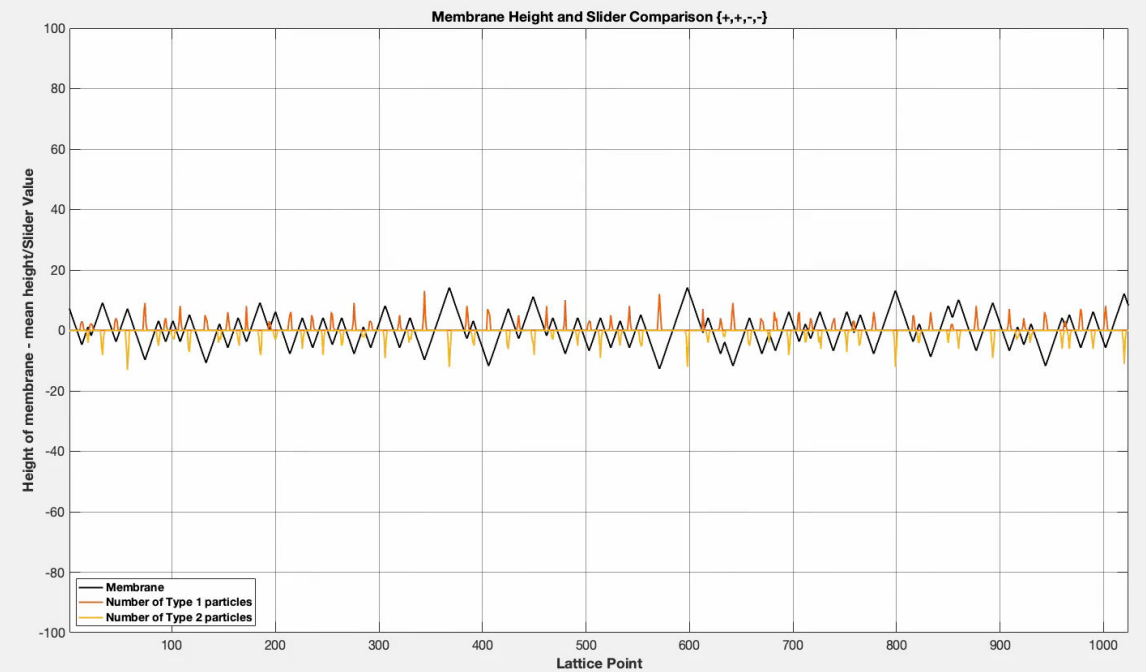
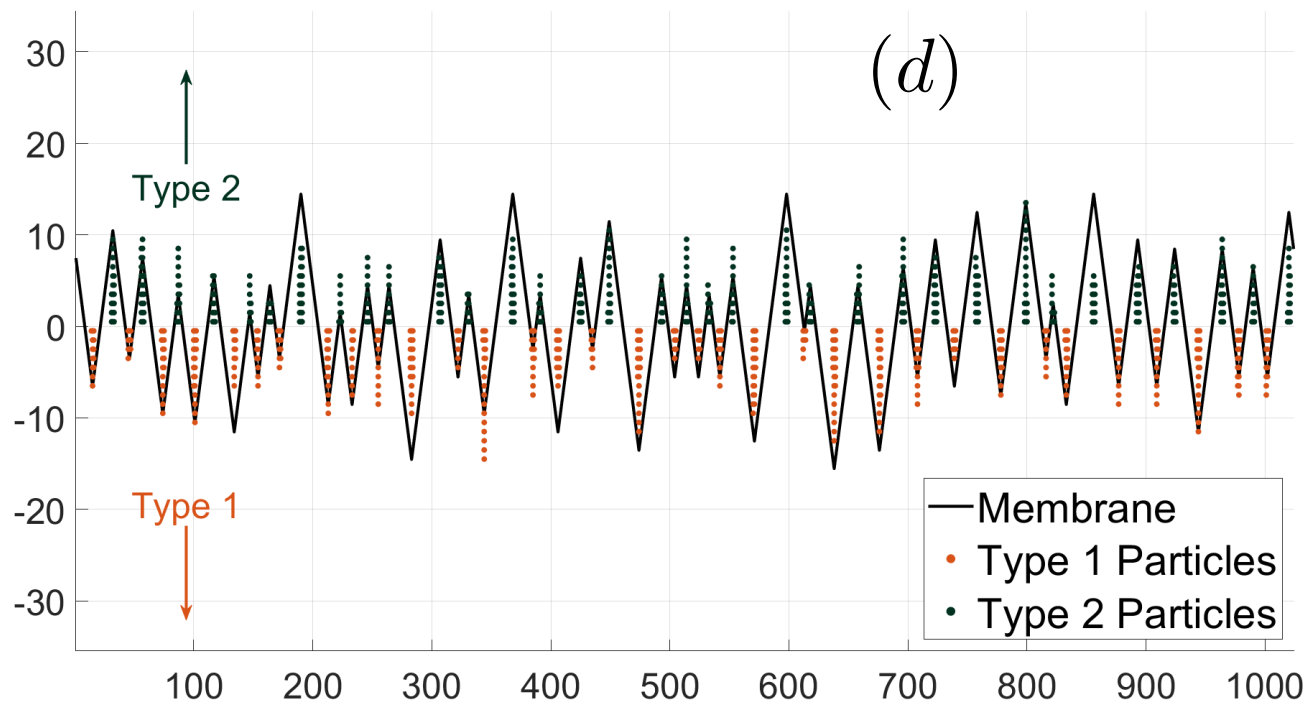
Type 1: Pushing membrane down and like valleys

Type 2: Pushing membrane down and like hills

CMAM like motion but with a memory. Non-Markovian motion of clusters.

Two types of particles

$\{+,+,-,-\}$



Type 1: Pulling membrane up and like hills

Type 2: Pushing membrane down and like valleys

CMAM like motion but with a memory. Non-Markovian motion of clusters.

Cluster Distribution function

Two types of particles

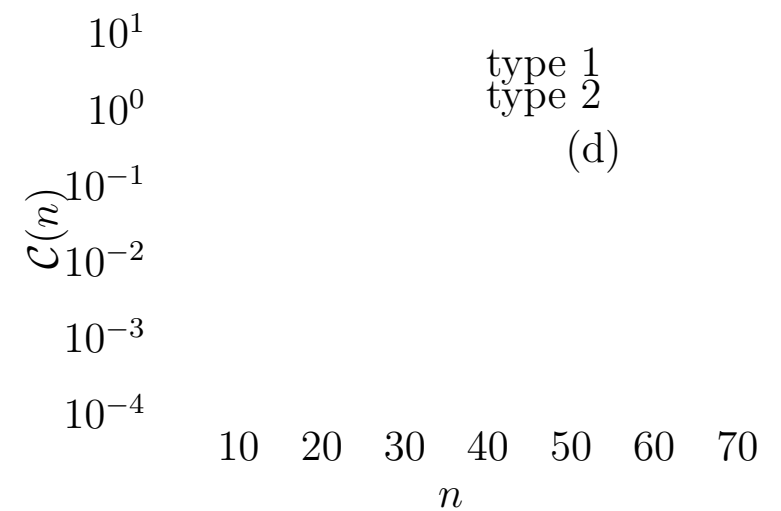
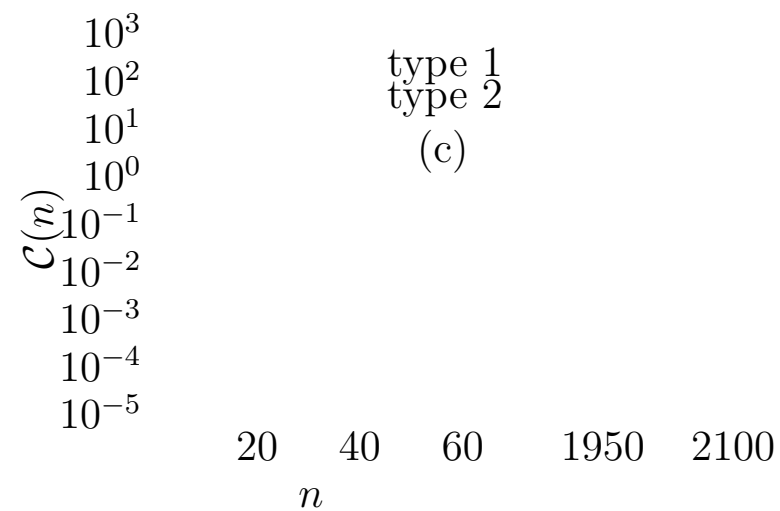
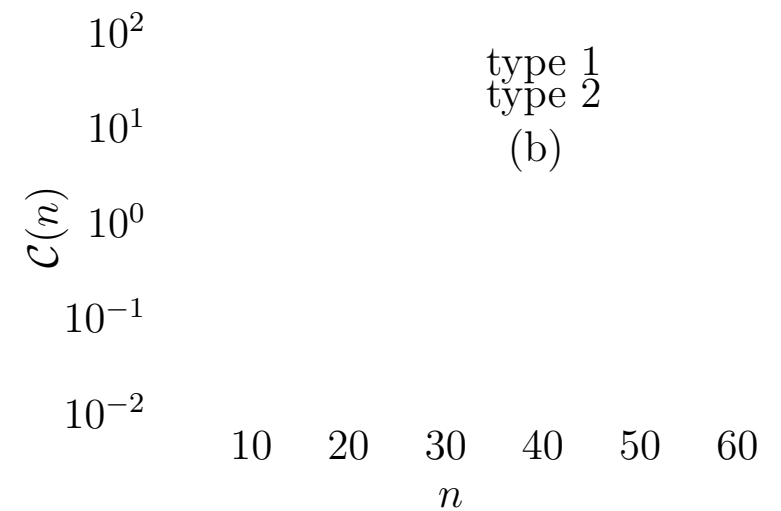
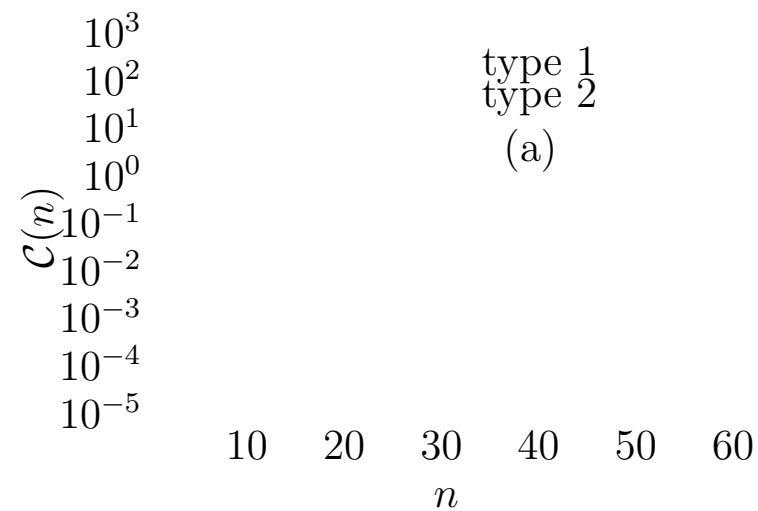
Reference: $\{\beta_1, \gamma_1, \beta_2, \gamma_2\}$

- (a) $\{+, -, -, +\}$

- (b) $\{+, +, -, +\}$

- (c) $\{-, -, -, +\}$

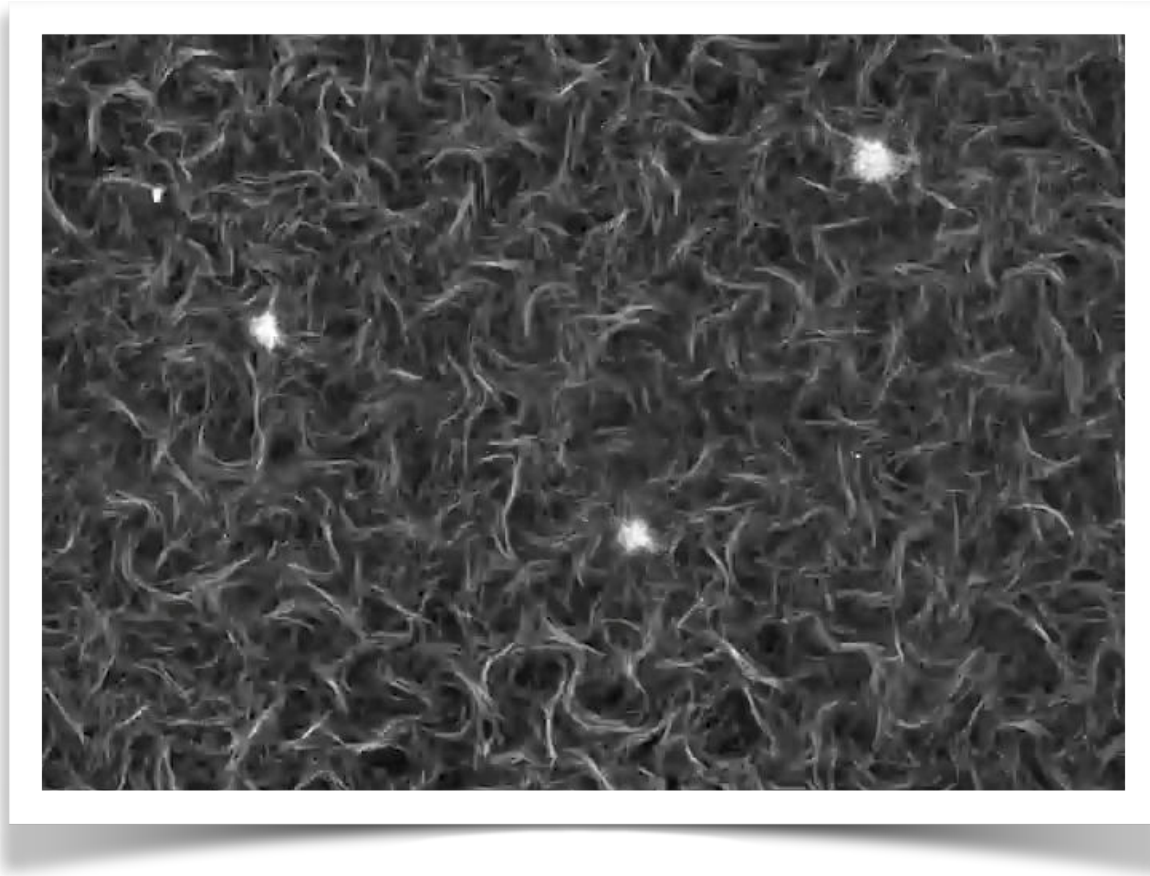
- (d) $\{+, +, -, -\}$



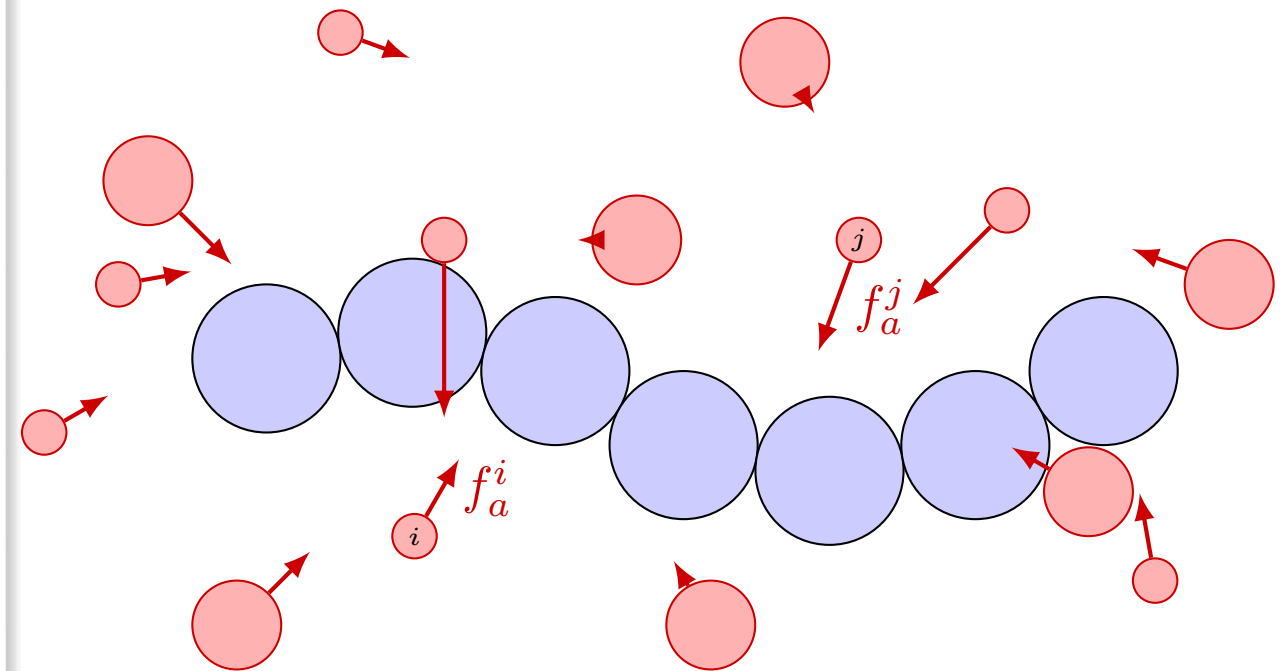
Part B

Polymer in active bath

Polymer in an active environment



Active collection of microtubules and kinesin motors.



A single polymer immersed in an active environment.

"Percolating Active Network of Microtubules."
YouTube, uploaded by Dogic Lab, 9 Jan. 2014,
<https://www.youtube.com/watch?v=NuFMloTVTsg>.

Model

Semiflexible Polymer

- Harmonic bonds.

$$V_b = \frac{k_s}{2} \sum_{i=2}^N (|\mathbf{r}_i - \mathbf{r}_{i-1}| - r_0)^2$$

- Rigidity: Semiflexibility.

$$V_a = \kappa \sum_{i=2}^{N-1} (1 - \cos \theta_i)$$

- Excluded volume interactions.

$$V_{WCA}(r) = 4\epsilon \left[\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^6 \right] + \epsilon; \text{ for } r \leq 2^{1/6}\sigma; 0 \text{ otherwise}$$

- Equations of motion of monomers:

$$m \frac{d\mathbf{v}_i}{dt} = -\gamma_t \mathbf{v}_i + \mathbf{F}_i^c + \sqrt{2\gamma_t k_B T} \boldsymbol{\xi}_i(t),$$

Model

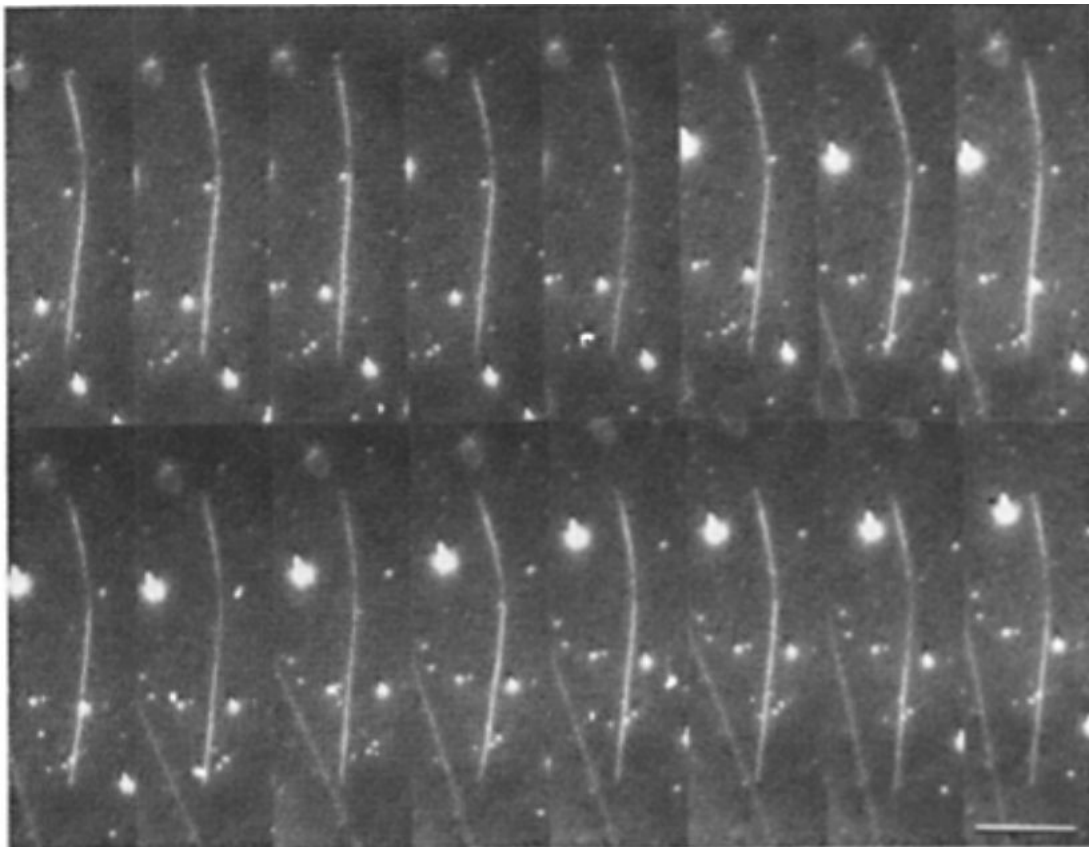
Active particles

- Active forcing: f_a
- Persistence time: $\tau = 1/D_r$
- Excluded volume interactions.
- **Equations of motion:**

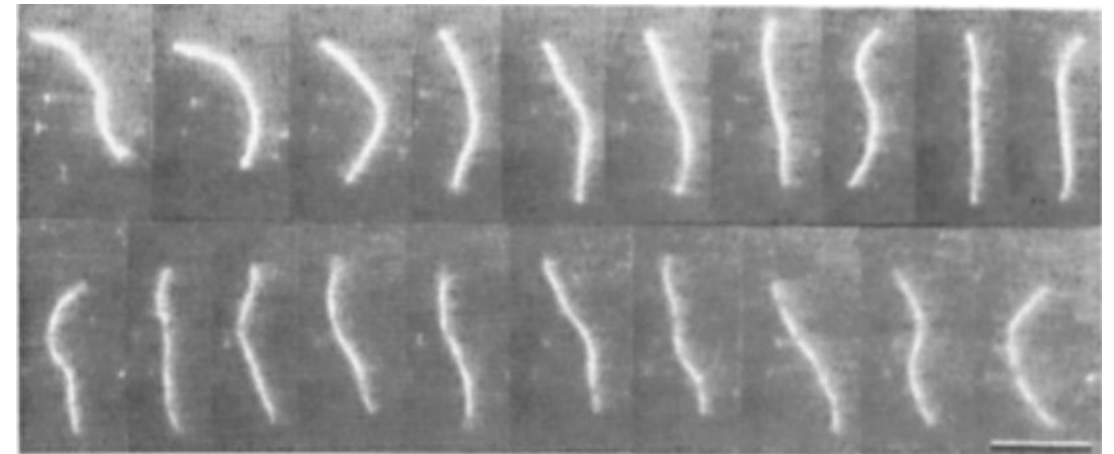
$$m_a \frac{d\mathbf{V}_j}{dt} = -\gamma_t \mathbf{V}_j + \mathbf{F}_j^c + f_a \hat{\mathbf{n}}_j + \sqrt{2\gamma_t k_B T} \boldsymbol{\Xi}_j(t),$$

$$I \frac{d^2\theta_j}{dt^2} = -\gamma_r \frac{d\theta_j}{dt} + \sqrt{2D_r} \eta_j(t),$$

Mode Amplitude



Microtubule undergoing
thermal fluctuations



Actin filaments undergoing
thermal fluctuations

Gittes, Frederick, et al. "Flexural rigidity
of microtubules and actin filaments measured
from thermal fluctuations in shape."
The Journal of cell biology 120.4 (1993): 923-934.

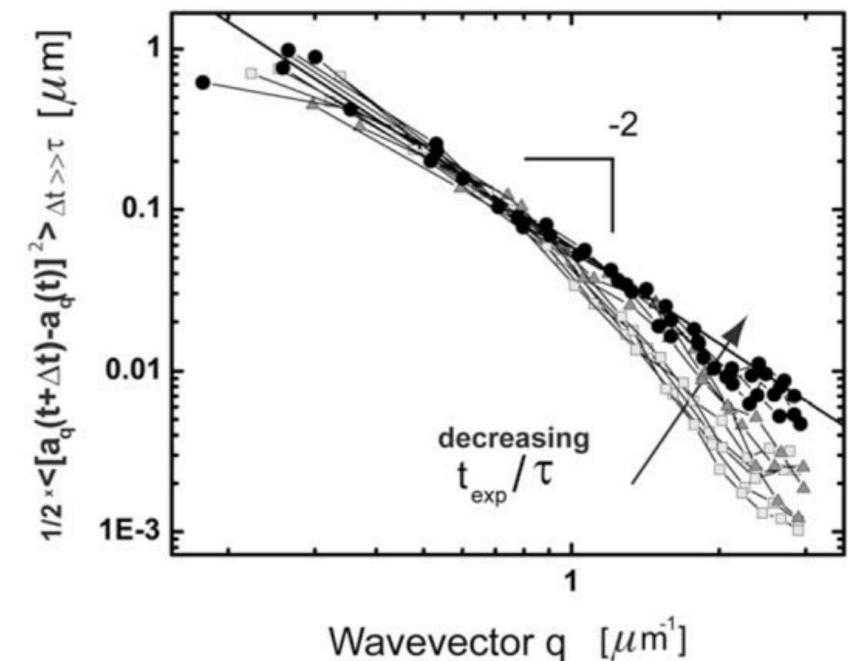
Mode Amplitude

- Bond vectors: $\Delta \mathbf{r}_k = \mathbf{r}_{k+1} - \mathbf{r}_k$
- We decompose the bending modes:
- Then we calculated the variance of these modes:
- What do we expect in purely thermal bath:

$$\mathbf{b}_n = \sqrt{\frac{2}{N}} \sum_{k=1}^{N-1} \Delta \mathbf{r}_k \cos \left[\frac{n\pi}{N} \left(k + \frac{1}{2} \right) \right]$$

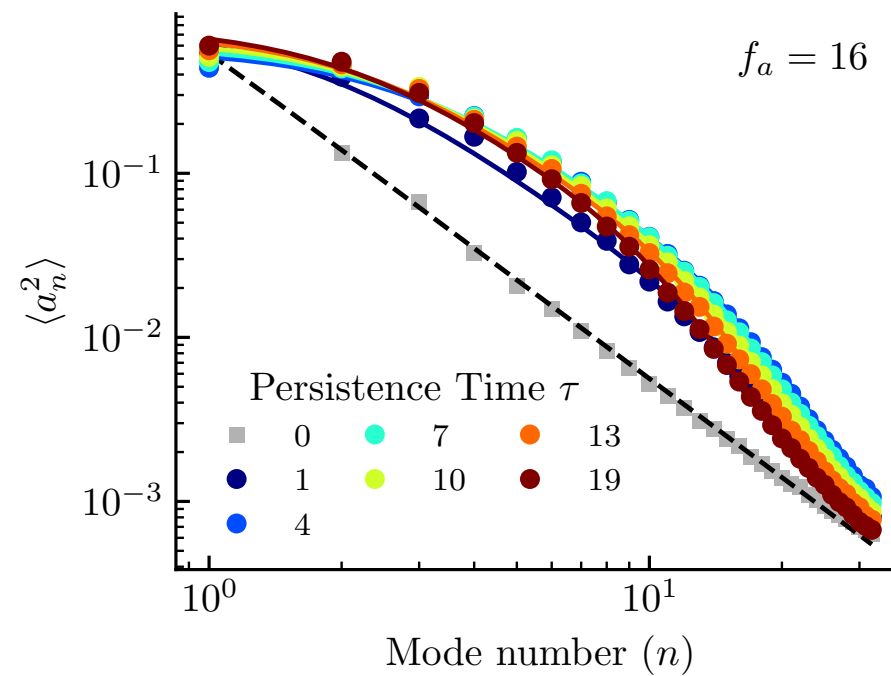
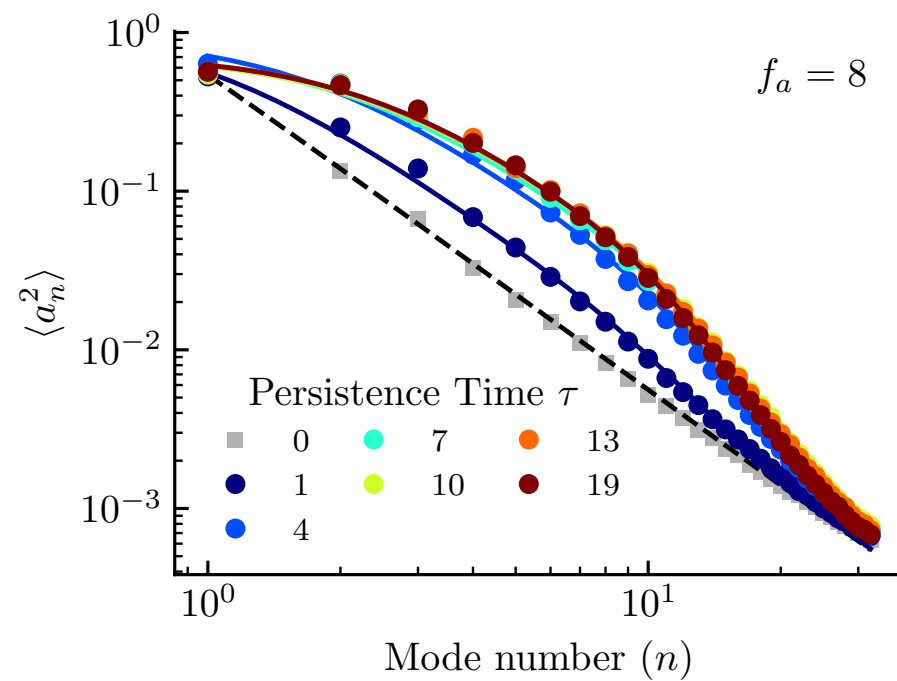
$$\langle |a_n|^2 \rangle = \langle |\mathbf{b}_n|^2 \rangle$$

$$\langle a_n^2 \rangle = \frac{k_B T}{\kappa q_n^2}$$

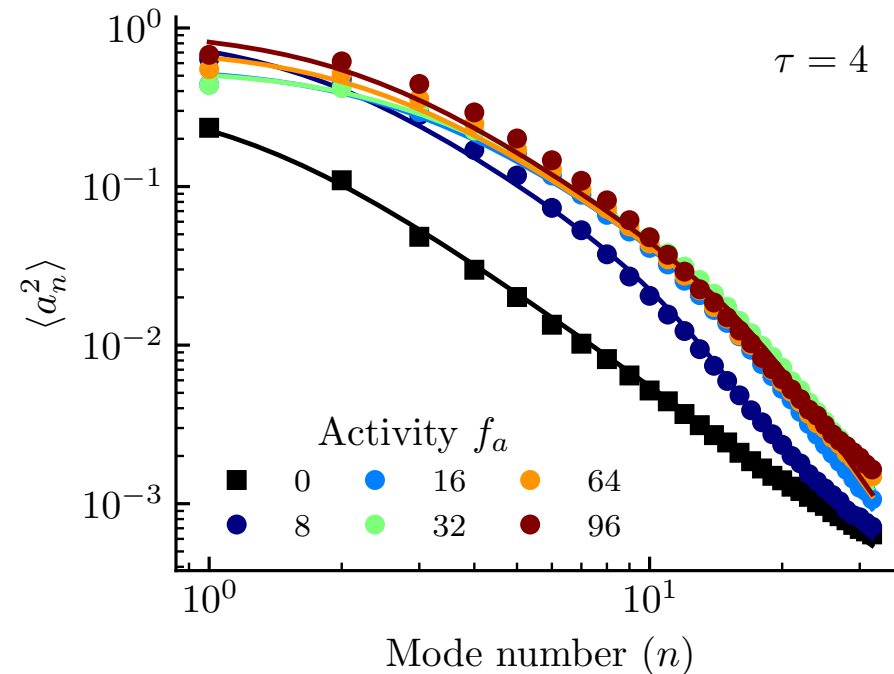
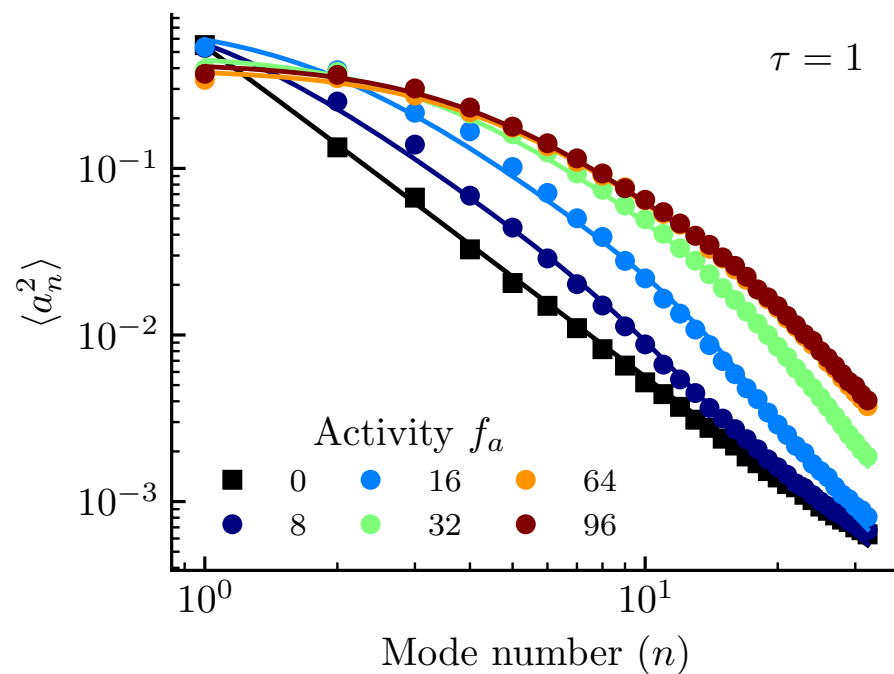
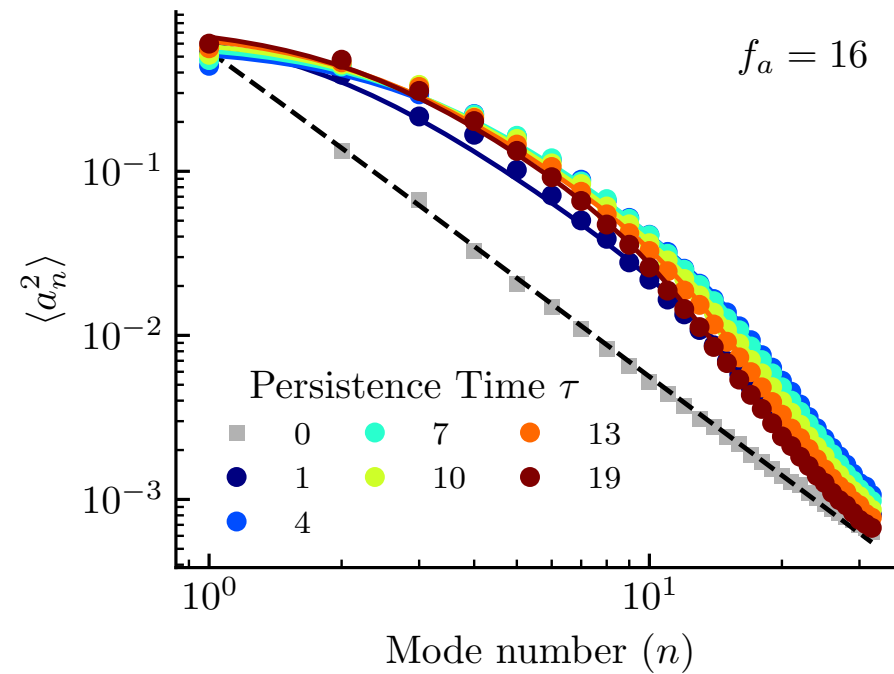
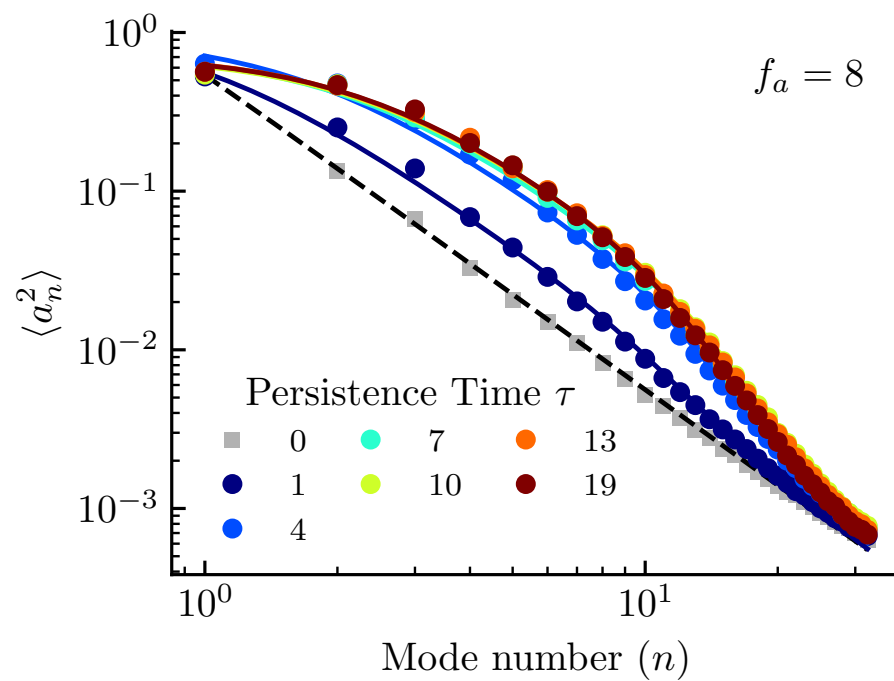


Brangwynne, Clifford P., et al. "Bending dynamics of fluctuating biopolymers probed by automated high-resolution filament tracking." *Biophysical journal* 93.1 (2007): 346-359.

Semiflexible polymer in active bath



Semiflexible polymer in active bath



Theoretical framework

- Semiflexible polymer in weak bending limit

$$\zeta \partial_t u_{\perp} = -\kappa \partial_s^4 u_{\perp} + \eta_u^{th} + \eta_u^{act}$$

- Decompose in cosine modes:

$$u_{\perp}(s, t) = \sum_{n \geq 1} a_n(t) \phi_n(s)$$

- We can immediately write the equation of motion of a_n

$$\zeta \dot{a}_n(t) + \kappa q_n^4 a_n(t) = \xi_n^{th}(t) + \xi_n^{act}(t),$$

- Fourier transform leads to the form of $\langle a_n^2 \rangle$

$$\langle |a_n|^2 \rangle = \int_{-\infty}^{\infty} \frac{d\omega}{2\pi} |H_n(\omega)|^2 [S_n^{th}(\omega) + S_n^{act}(\omega)]$$

- If we can calculate the active force-force correlations we will be able to get a form of variance of tangent mode amplitude.

Theoretical framework

- From microscopic picture of active particles

$$C_{\alpha\beta}(\Delta\mathbf{r}, \Delta t) = \rho_0 V_0^2 \sigma^{2d} \left(\frac{\pi}{A(\Delta t)} \right)^{d/2} \left[\frac{\delta_{\alpha\beta}}{2A(\Delta t)} - \frac{\Delta r_\alpha \Delta r_\beta}{4A(\Delta t)^2} \right] \exp \left[-\frac{|\Delta\mathbf{r}|^2}{4A(\Delta t)} \right],$$

- With some further approximations

$$\langle \xi_n^{u,\text{act}}(t) \xi_m^{u,\text{act}}(t') \rangle = \delta_{nm} q_n^2 D_0 e^{-q_n^2 \sigma^2} e^{-|\Delta t|/\tau_{\text{eff}}}.$$

- And therefore

$$S_n^{\text{act}}(\omega) = q_n^2 D_0 e^{-q_n^2 \sigma^2} \frac{2\tau_{\text{eff}}}{1 + \omega^2 \tau_{\text{eff}}^2}.$$

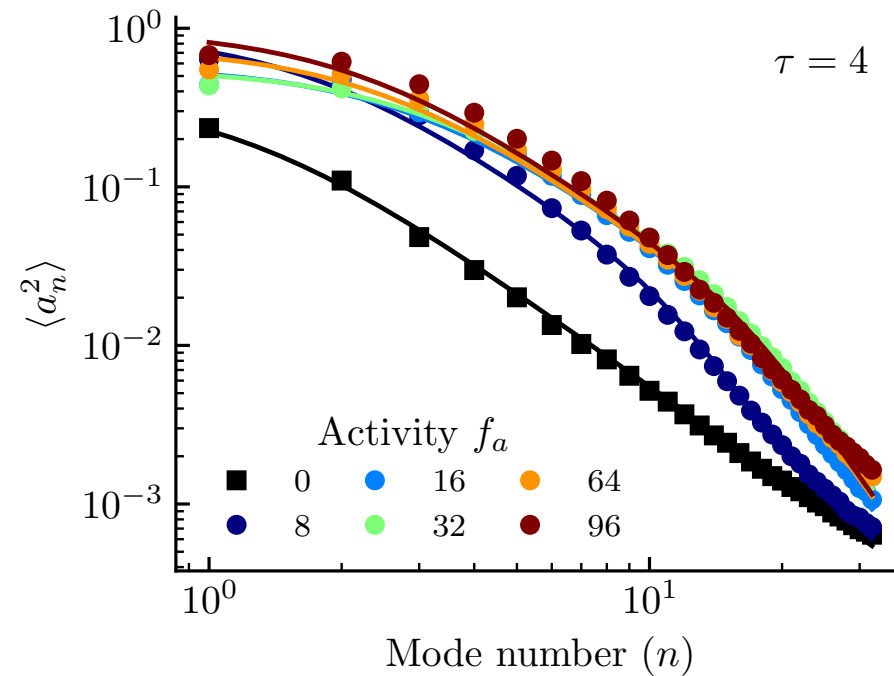
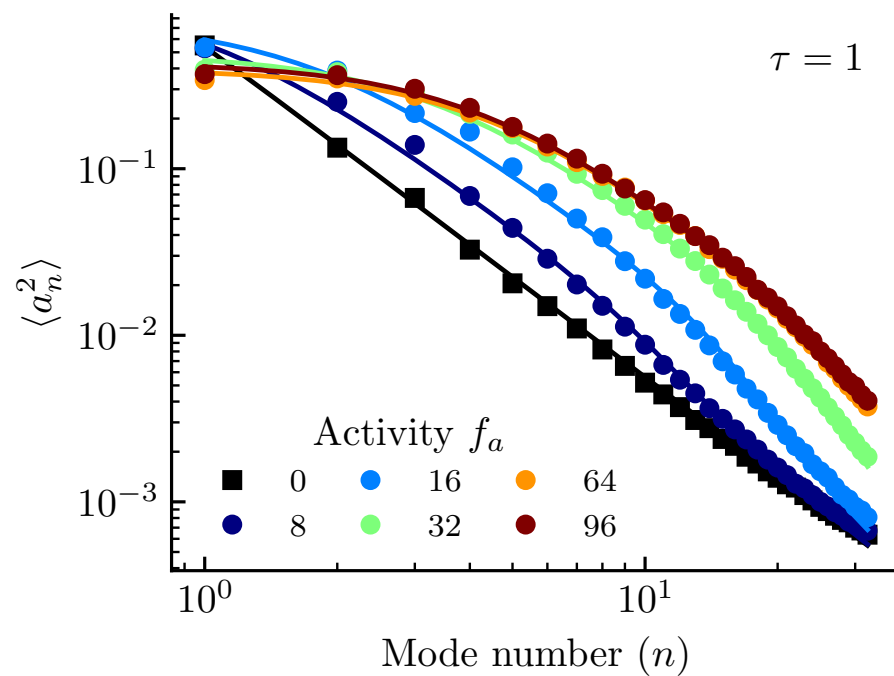
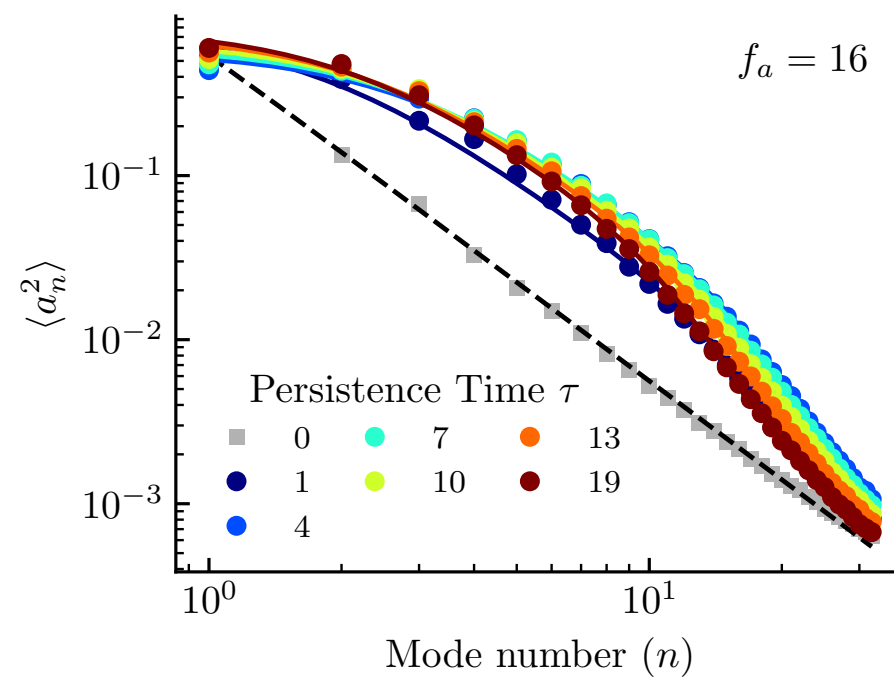
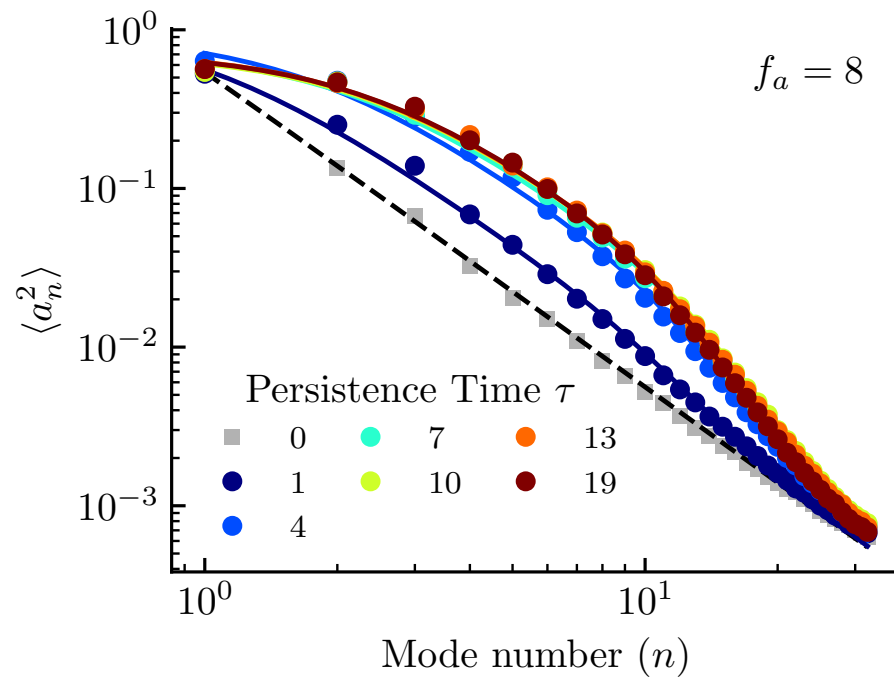
- The variance is finally:

$$\langle |a_n|^2 \rangle = \frac{1}{\kappa q_n^2} \left[k_B T + \frac{D_0 \tau_{\text{eff}}}{\zeta [1 + \tau_{\text{eff}} \omega_n]} e^{-q_n^2 \sigma^2} \right]$$

Further a mode dependent temperature can also be defined here:

$$T_n^{\text{eff}} = T + \frac{D_0 \tau_{\text{eff}}}{k_B \zeta [1 + \tau_{\text{eff}} \kappa q_n^4 / \zeta]} e^{-q_n^2 \sigma^2}.$$

Semiflexible polymer in active bath



$$\langle |a_n|^2 \rangle = \frac{1}{\kappa q_n^2 + \mathcal{T}} \left[k_B T + \frac{D_0 \tau_{\text{eff}}}{\zeta [1 + \tau_{\text{eff}} \omega_n(\mathcal{T})]} e^{-q_n^2 \sigma^2} \right]$$

R_g reconstruction using mode variance

- Radius of gyration of polymer can be written as:

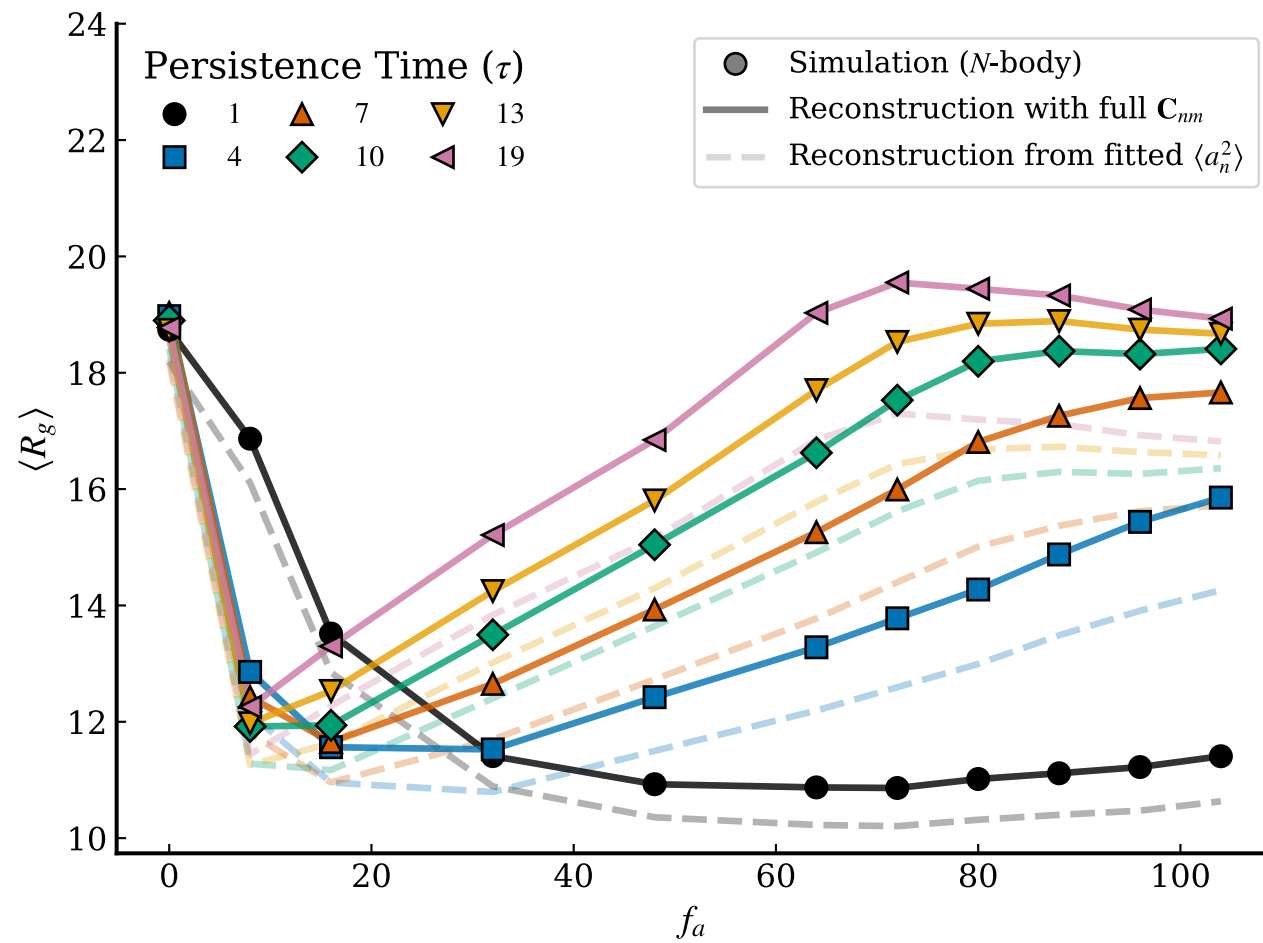
$$R_g^2 = \frac{1}{(N+1)^2} \sum_{i=0}^N \sum_{j=i+1}^N \left\langle |\mathbf{r}_j - \mathbf{r}_i|^2 \right\rangle.$$

- We can expand the bond vector in discrete cosine basis

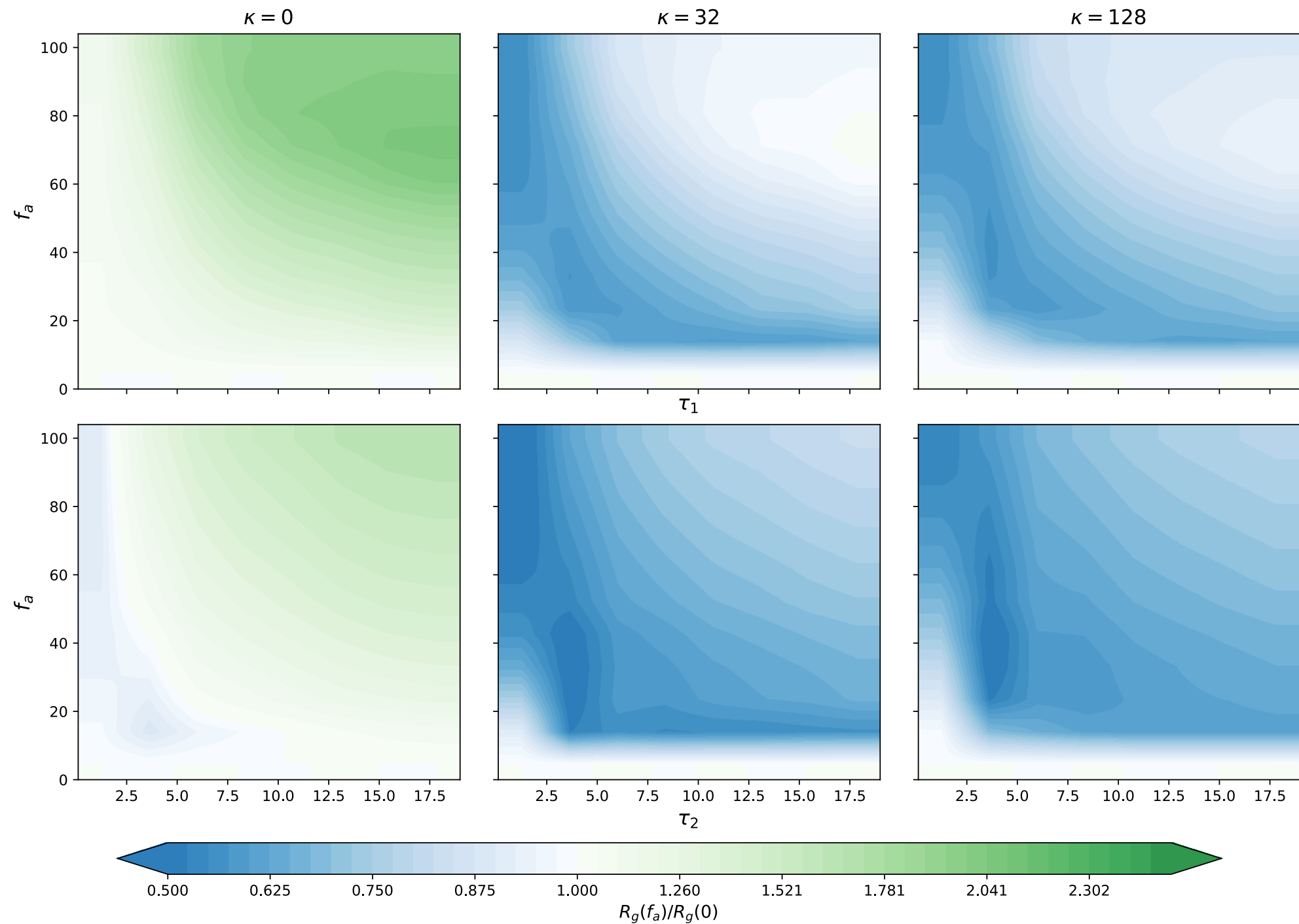
$$\Delta \mathbf{r}_k = \frac{1}{2} \mathbf{b}_0 + \sum_{n=1}^{N-1} \mathbf{b}_n \cos \left[\frac{n\pi}{N} \left(k + \frac{1}{2} \right) \right]$$

- $$R_g^2 = W_{00} \langle \mathbf{b}_0 \cdot \mathbf{b}_0 \rangle + \sum_{n \geq 1} W_{nn} \langle \mathbf{b}_n \cdot \mathbf{b}_n \rangle + \sum_{n \geq 1} W_{0n} \langle \mathbf{b}_0 \cdot \mathbf{b}_n \rangle + \sum_{n < m} W_{nm} \langle \mathbf{b}_n \cdot \mathbf{b}_m \rangle.$$

R_g reconstruction using mode variance



R_g comparison of explicit vs implicit bath



Limitations & Caveats

- Due to high active force and long persistent time the bond angle stretched is a factor that needed to be accounted further.
- The spatial correlations are not considered completely.
- With our theory which is with weak bending limit the reconstruction of R_g is systematically lesser than the actual R_g .
- Assumption that the different modes decouple.

Acknowledgement

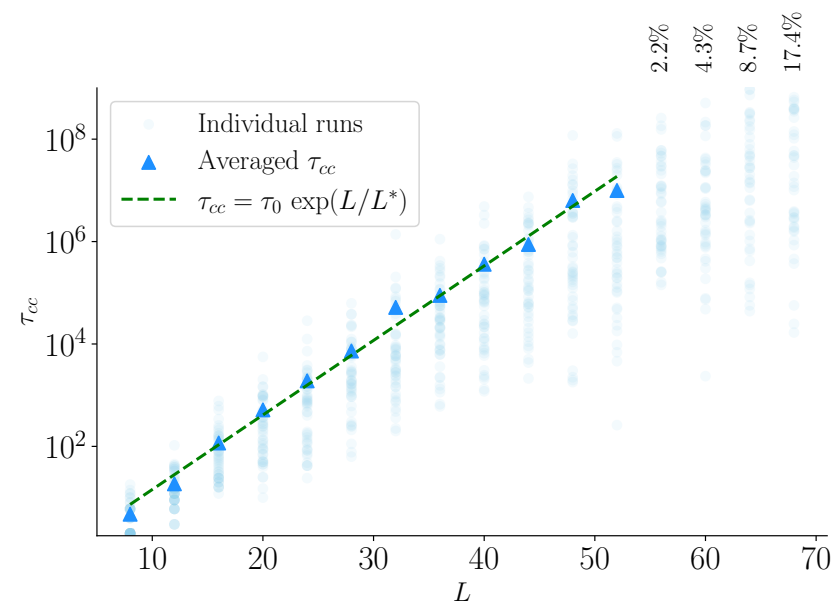
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Thank you :)

Questions?

Slow Coarsening and Large Cluster movements

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