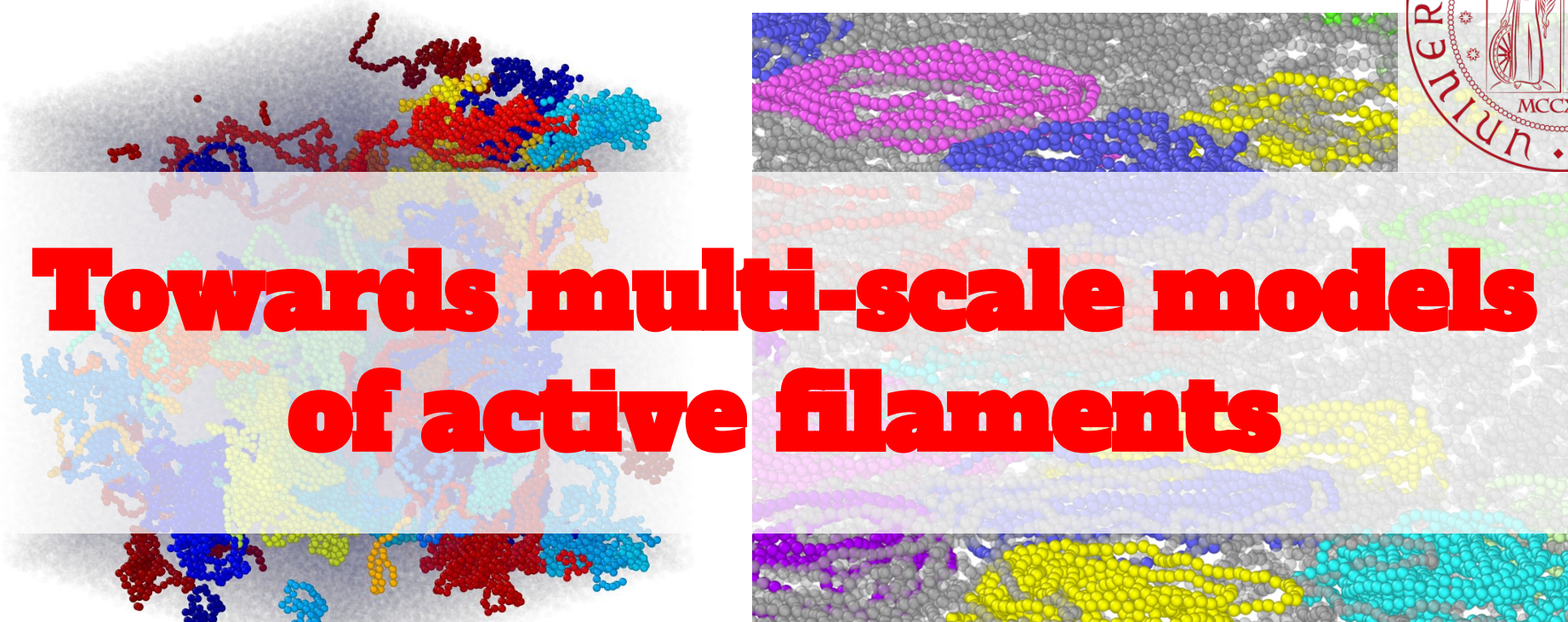




@elocatelli86



<http://statphys.dfa.unipd.it/people/locatelli.php>

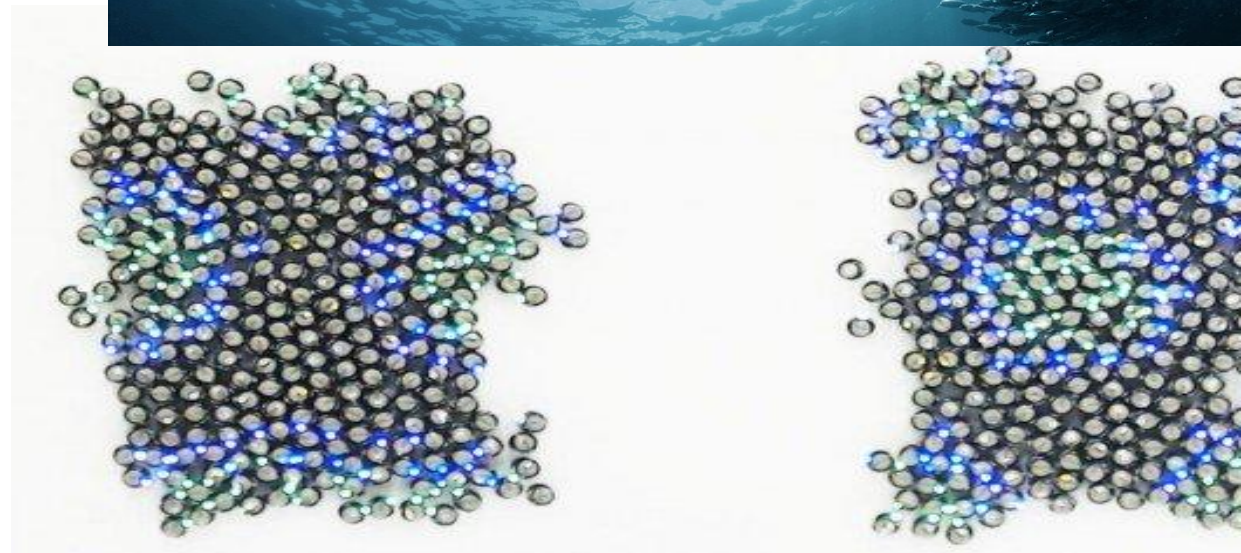
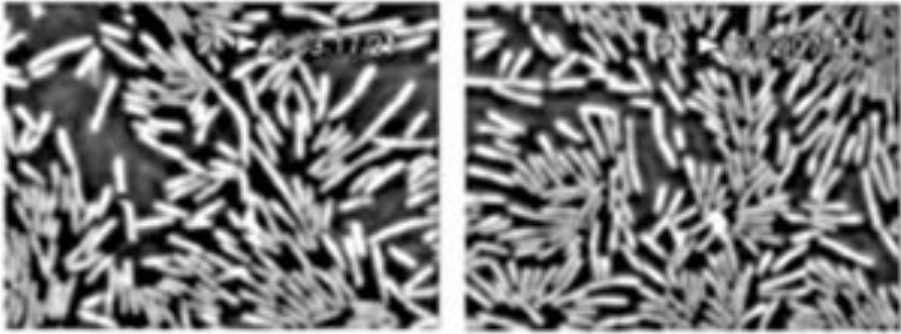


Towards multi-scale models of active filaments

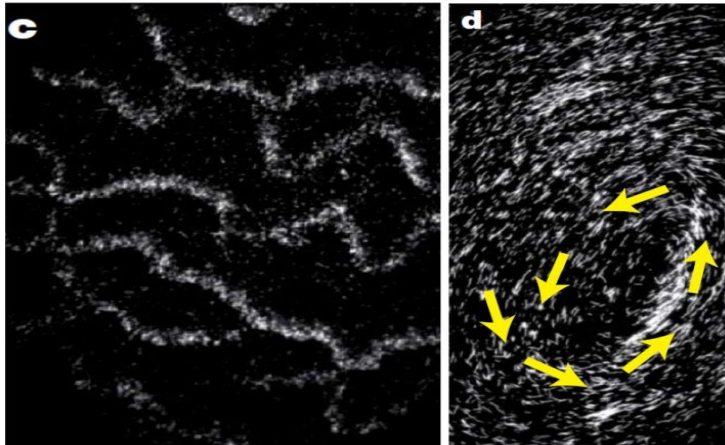
Emanuele Locatelli

**Department of Physics and Astronomy,
University of Padova
ECT* Workshop, Trento, 19-04-2024**

Active Matter

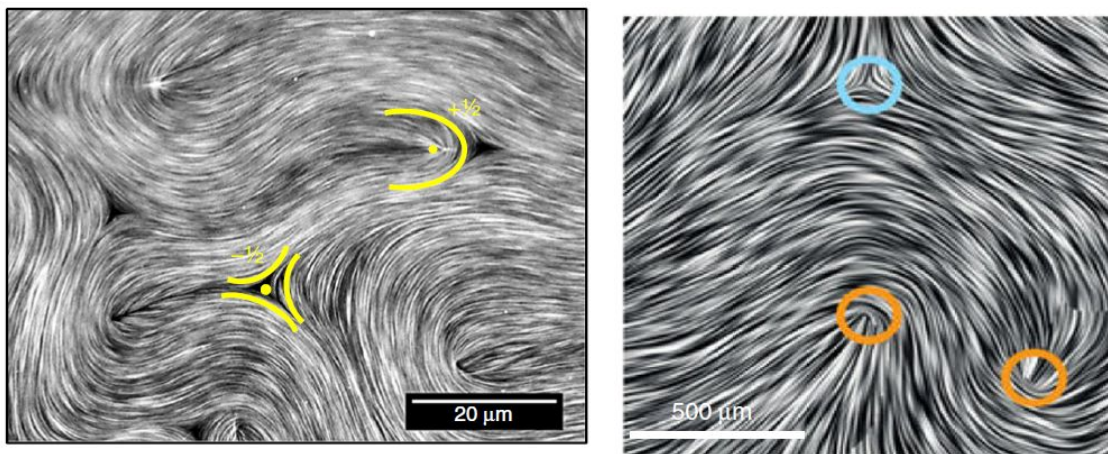


Active Polymeric Matter

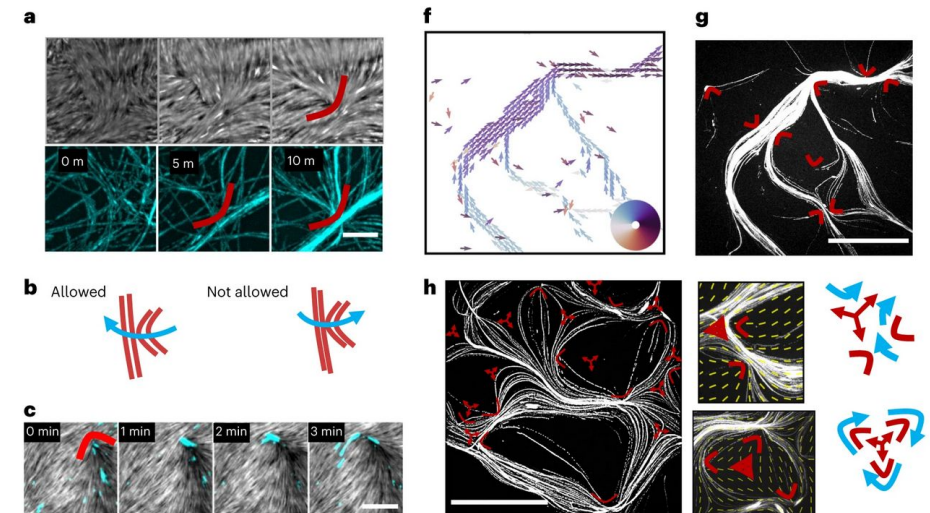


Active filaments (Active Nematics)
Nature, 467 (2010).

Nature Materials 22, 260–268 (2023)



Nat. Comm. 9, 1, 3246 (2018)

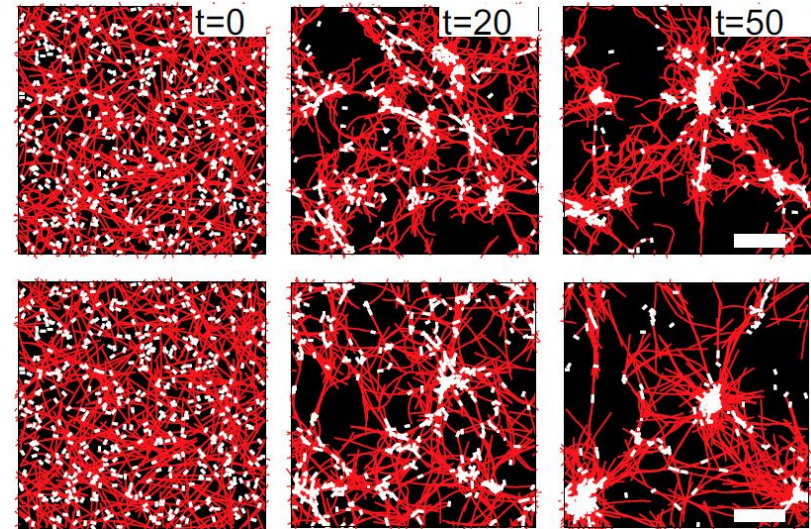


Active Polymeric Matter

Active intracellular processes

Nature, 416 (2002) | PNAS, 114 (2017)

Nature Physics 11, 2, 111 (2015).



Collective motion interphase chromatin

PNAS 110, 15555–15560 (2013) | PNAS 115, 45, 11442–11447

(2018) | PRX 12, 041033 (2022) | PRL 131, 048401 (2023)

Chromatin organization

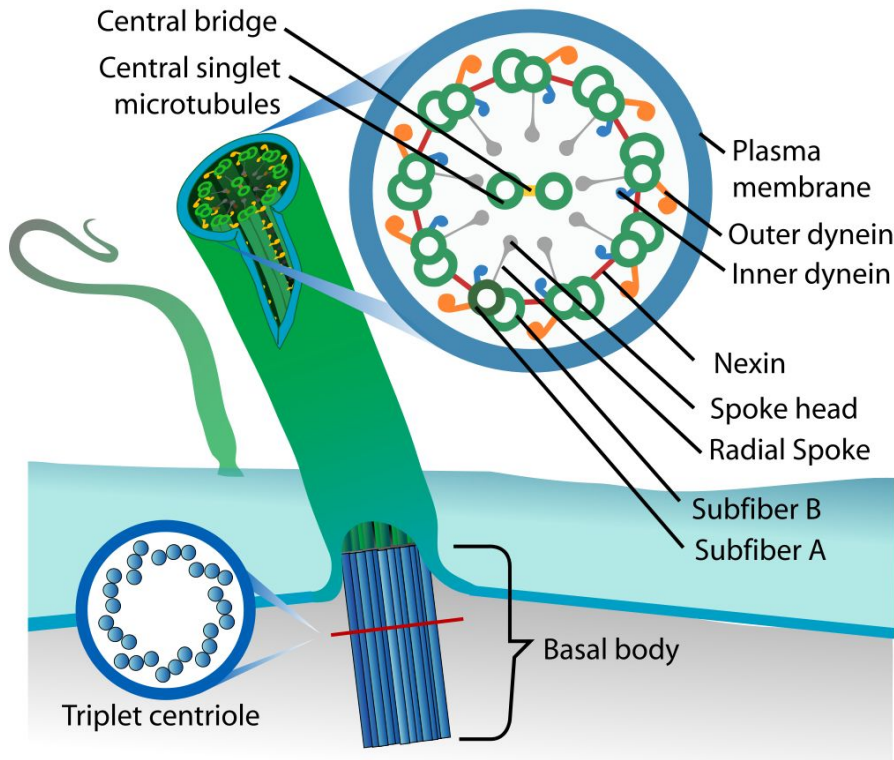
PRL 126, 158101 (2021) | Soft Matter, 19, 1348–1355 (2023) | Soft Matter,

18, 8134–8146 (2022) | Biophys. J. 106, 9, 1871–1881 (2014) | Phys. Rev.

E 99, 032421 (2019) | PNAS 120 e2221726120 (2023)



Active Polymeric Matter



Cilia in eukaryotic cells

Science 337: 937–941, (2012)| Annu. Rev. Physiol., 77:379 (2015)| Nat. Phys., 16 1158–1164 (2020)

Modelisation with active polymers

Sci. Rep. 7, 16758 (2017)| Soft Matter, 15, 7926 (2019)| J.Chem.Phys 146, 154901 (2017)| Phys. Rev. E 106, 054501 (2022)

J. R. Soc. Interface 11:20130884 (2013) | J. R. Soc. Interface 14: 20170491 (2017) | Sci. Rep. 3, 1964 (2013) | PRFluids 4, 043102 (2019)| PRL 123, 208101 (2019)

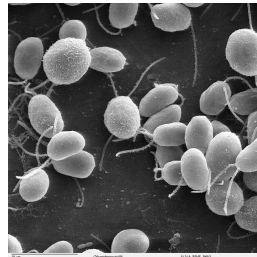
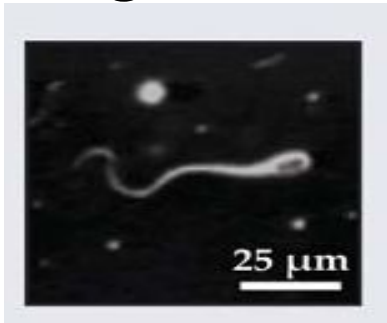
Source: Wikipedia

Cilia carpet in lung cells

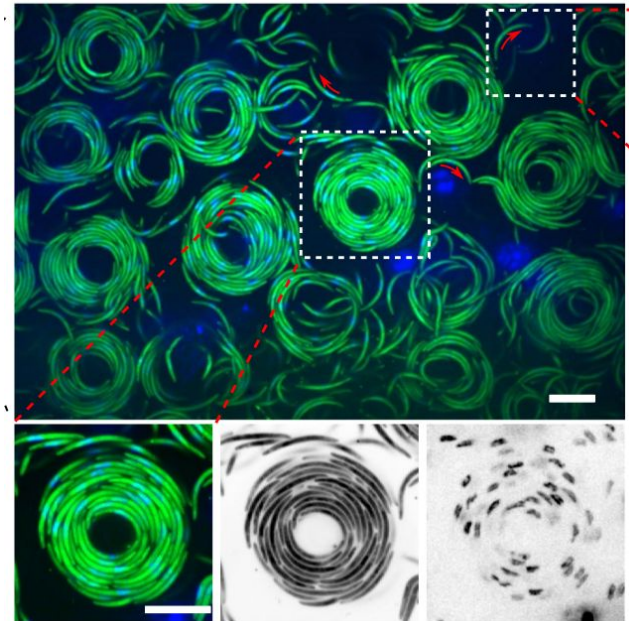
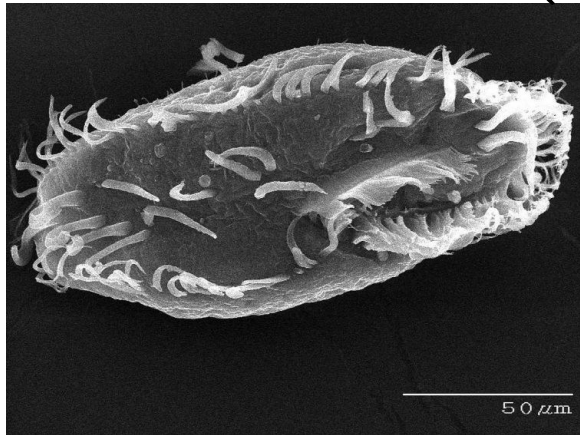
Active Polymeric Matter

Microswimmer locomotion

Flagella locomotion (Spermatozoa, *C. Reinhardtii*)



Cilia locomotion (Ciliate)



Plasmodium sporozites,
Nat. Phys. 18, 586–594 (2022)

Active Polymeric Matter

Microswimmer locomotion

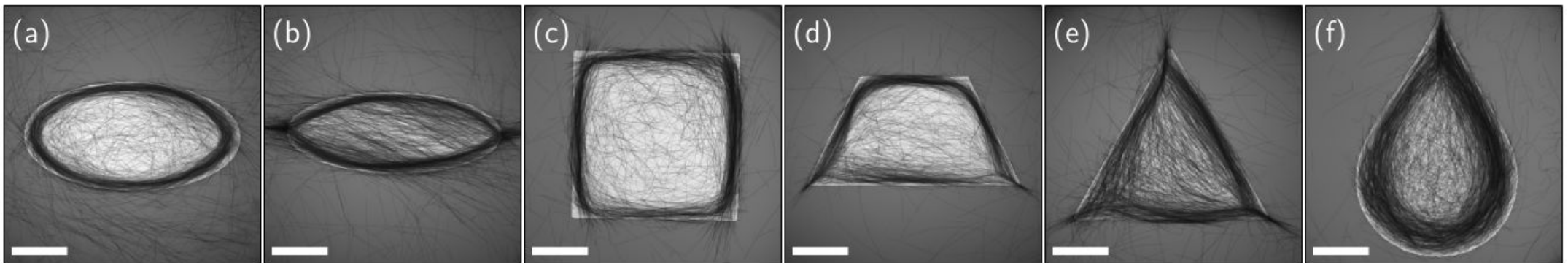
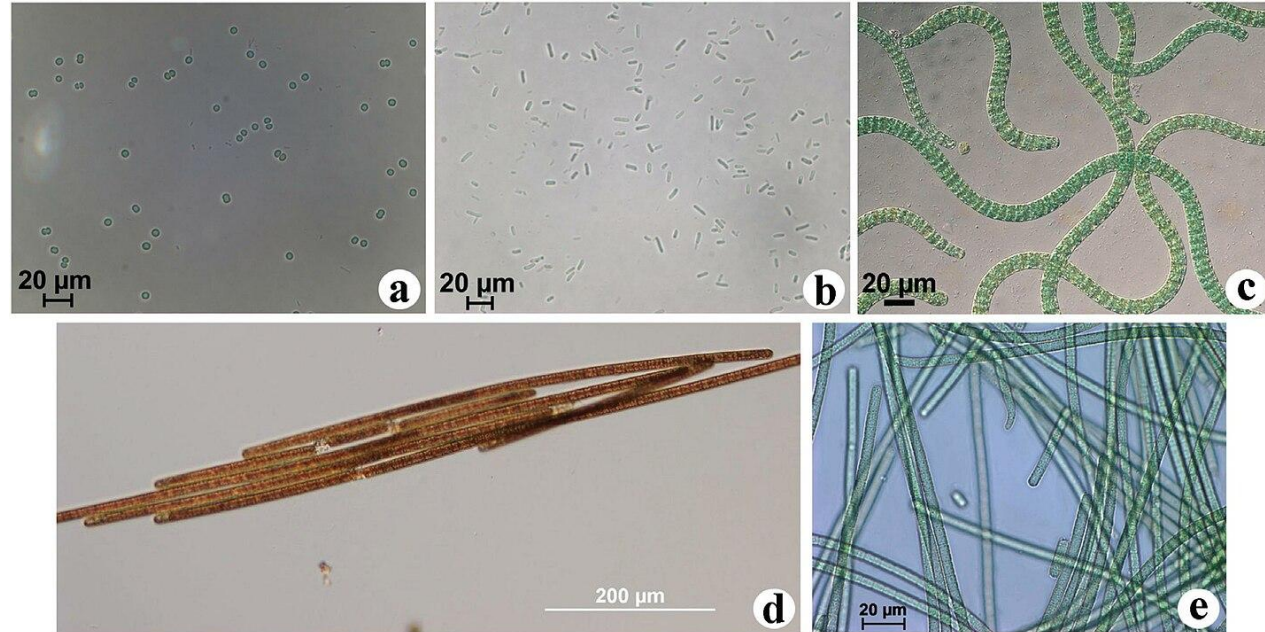
Cyanobacteria

Scientific American 270.1 (1994): 78–86.

eLife 12 (2023)

Mol. Microbiol. 98, 1021 (2015)

arXiv:2403.03093



Active Polymeric Matter

Macroswimmers: *Tubifex Tubifex* & other worms



Organization

Nat. Comm. 10.683 (2019)| arXiv:2301.11667 (2023)

Dynamics & Rheology

Soft Matter, 18, 1174 (2022)| Nat. Phys., 17, 275-283 (2021)| PNAS 116, 51, 25569-25574 (2019)| arXiv:2303.00647| PRL 124, 188002 (2020)| Sci. Adv., 8, eabj7918 (2022)

Phase separation

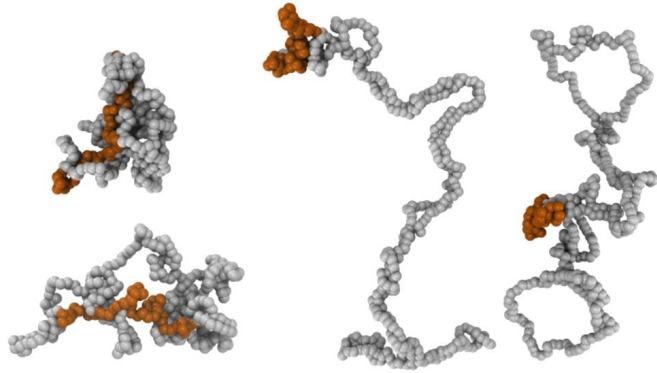
PRL 124, 208006 (2020)| PNAS, 118, 2010542 (2021)| Front. Phys., 9, 734499 (2021)

Entanglement & collective motion

Int. & Comp. Biol., 62, 890-896 (2022)| Science 380.6643 (2023)| Soft Matter 19,10 (2023)

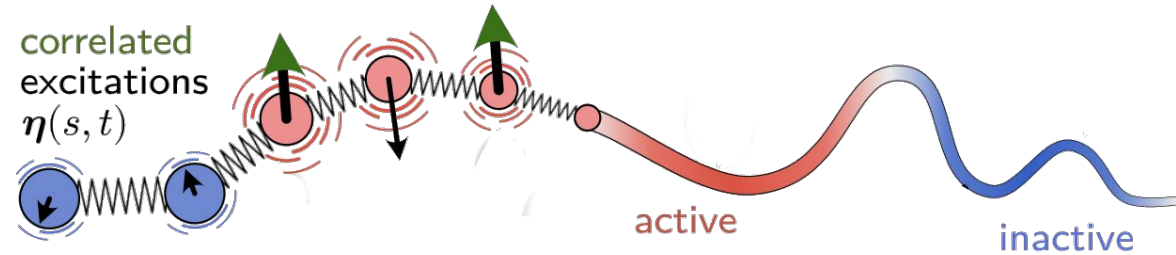
Active Polymer Models

2-temperature model



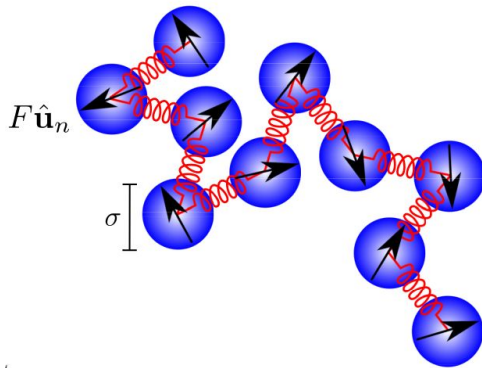
From: Nat. Comm. 11, 26 (2020)

Correlated noise



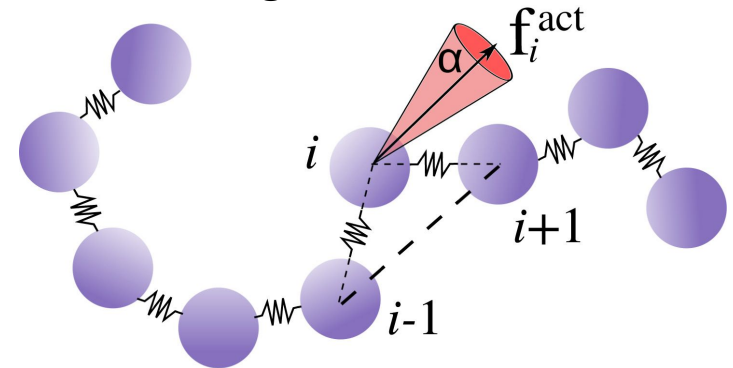
From: PNAS 120 e2221726120 (2023)

Active Brownian Polymers



From: J.Chem.Phys. 142, 124905 (2015)

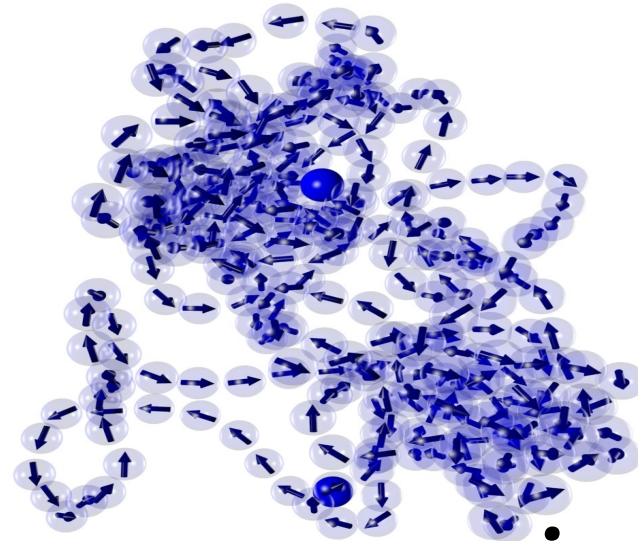
Active Polar (Tangential) Polymers



From: PRL 121, 217802 (2018)

Outline

Polar Active Polymers

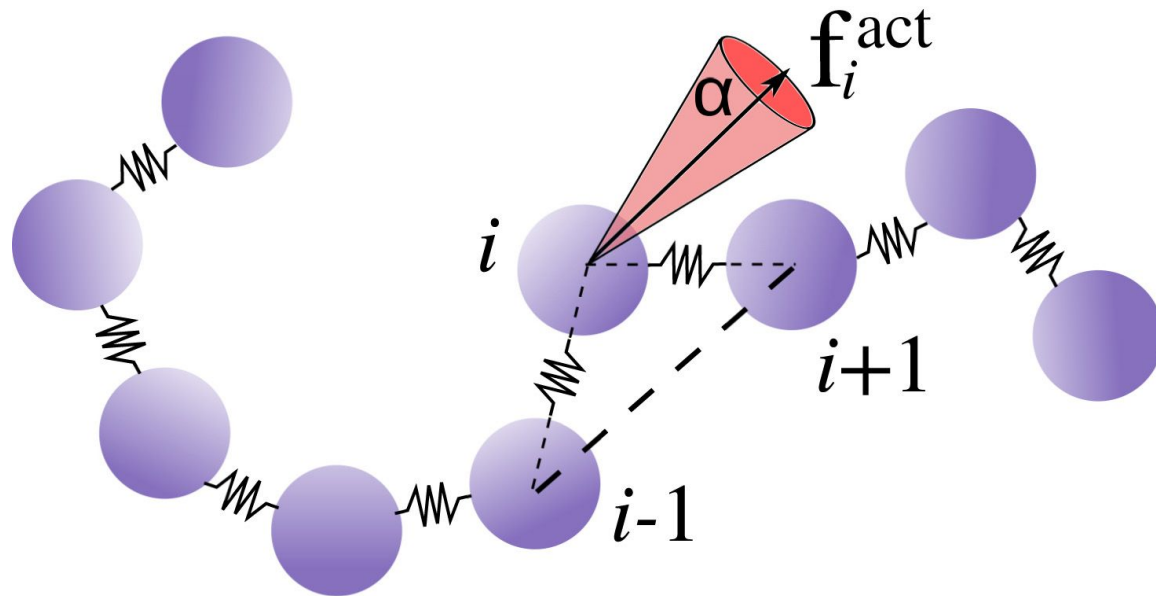


Multi-scale analysis of dense suspensions

Towards modeling biophysical systems

Polar Active Polymer

Polar (tangential) self-propulsion



Decorrelation/relaxation time

$$\tau \frac{D_0}{\sigma^2} \sim \frac{N}{Pe}$$

Total active force

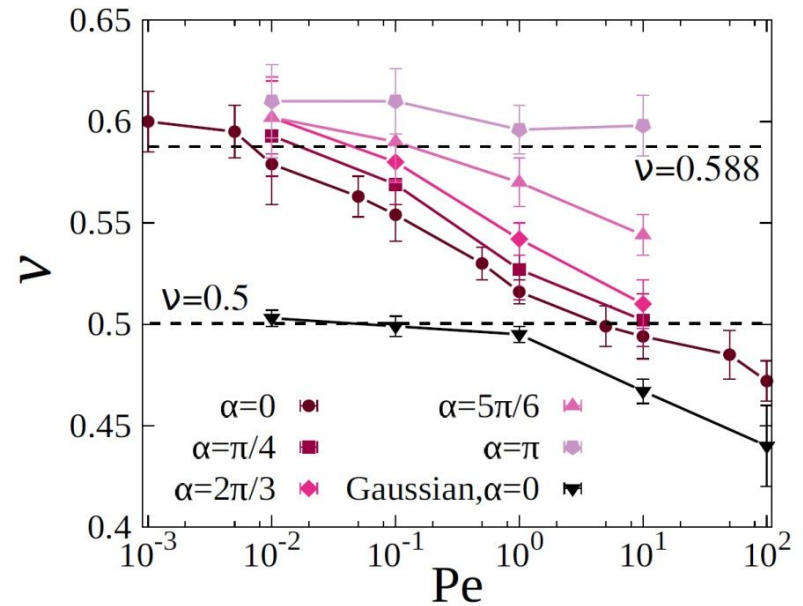
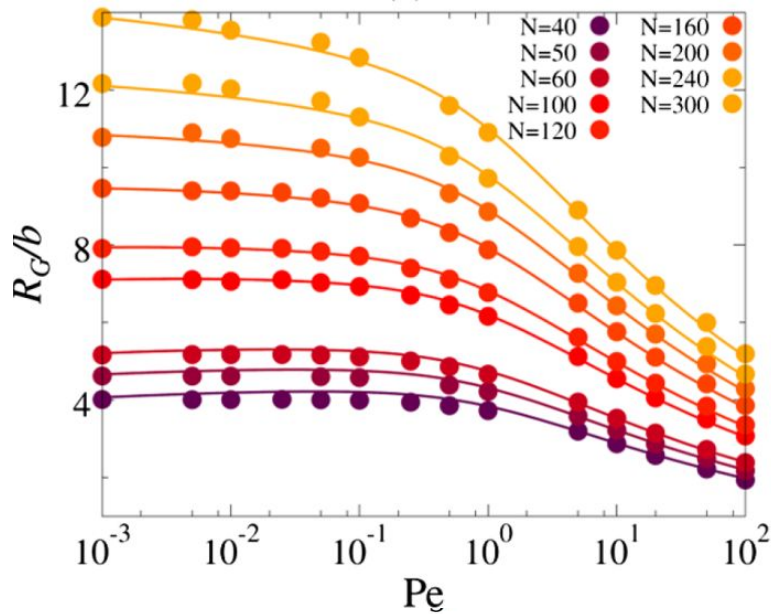
$$\mathbf{F}^{\text{act}} = \sum_i^N \mathbf{f}_i^{\text{act}} \approx \mathbf{R}_E$$

$$Pe \equiv \frac{v^{\text{act}} b}{D_0} = \frac{f^{\text{act}} b}{k_B T}$$

Polar Active Polymer

Radius of gyration $R_g^2 = \frac{1}{N} \sum_i (\mathbf{r}_i - \mathbf{r}_{cm})^2$

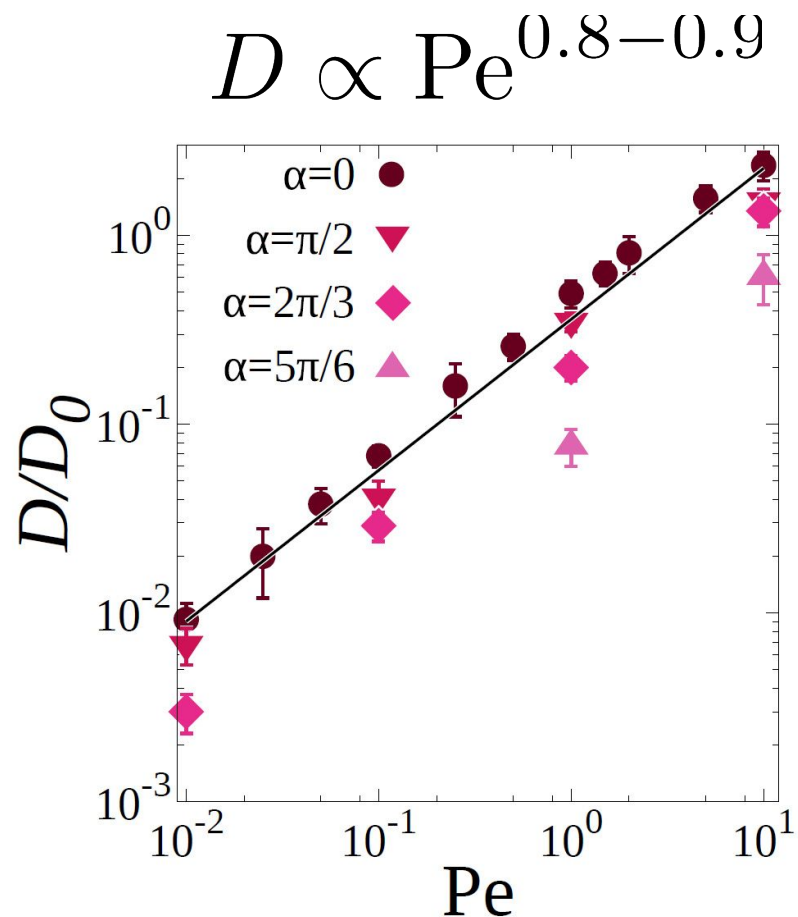
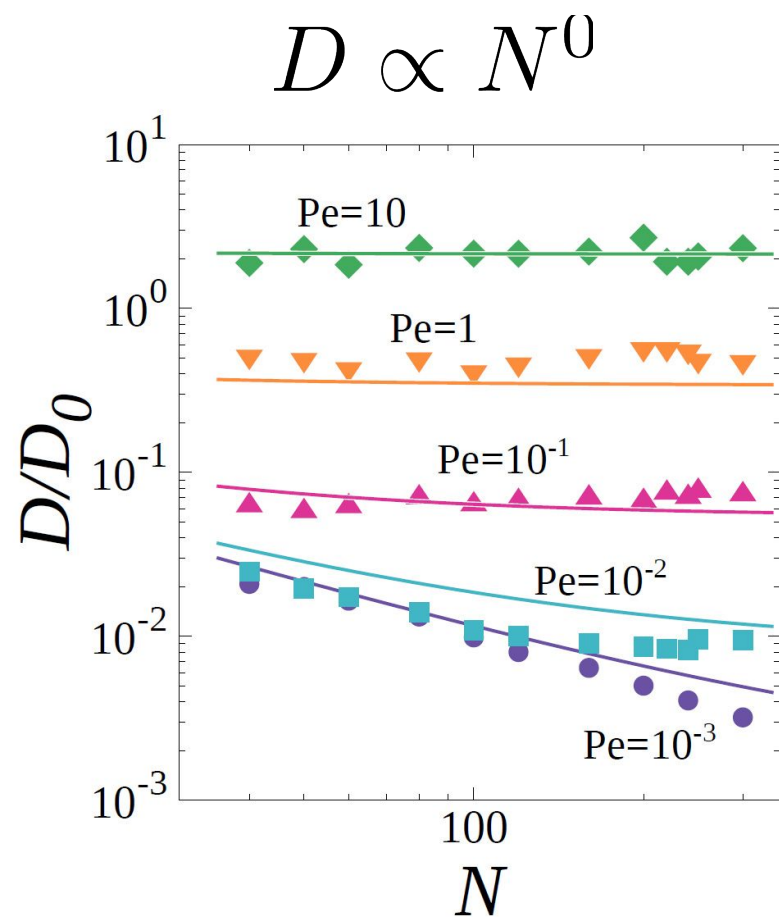
Scaling exponent $\langle R_g \rangle \propto N^\nu$



Coil-to-globule-like transition: the polymer becomes more compact (bundled) for higher values of Pe

Due to follower-forces instability (akin to buckling instability)

Polar Active Polymer



Polar Active Polymer

Minimal stochastic model for the dynamics of the center of mass

$$\dot{\mathbf{r}}_{CM} \equiv \frac{1}{N} \sum_i^N \dot{\mathbf{r}}_i = \frac{1}{N} (\boldsymbol{\xi} + \boldsymbol{\eta})$$
$$\boldsymbol{\xi} \equiv \frac{1}{\zeta} \mathbf{F}^{\text{act}} = \frac{1}{\zeta} \sum_i^N \mathbf{f}_i^{\text{act}}$$
$$\boldsymbol{\eta} \equiv \frac{1}{\zeta} \sum_i^N \boldsymbol{\eta}_i$$

As \mathbf{F}^{act} is almost parallel to the end-to-end vector, we model it as a random force with the same decorrelation properties as $\mathbf{C}(t)$

$$\langle \boldsymbol{\xi}(t) \boldsymbol{\xi}(t') \rangle \propto C(t) \equiv \langle \mathbf{r}_E(t) \cdot \mathbf{r}_E(0) \rangle$$



$$D = 2a N^{2\nu} (\text{Pe})^{-1} + \frac{D_0}{N}$$

Polar Active Polymer

Even simpler approach:

$$D = D_t + \frac{\tau_r v_a^2}{2d} = D_t + \frac{\tau_r (f_a/\gamma)^2}{2d}$$

Active Brownian Particle

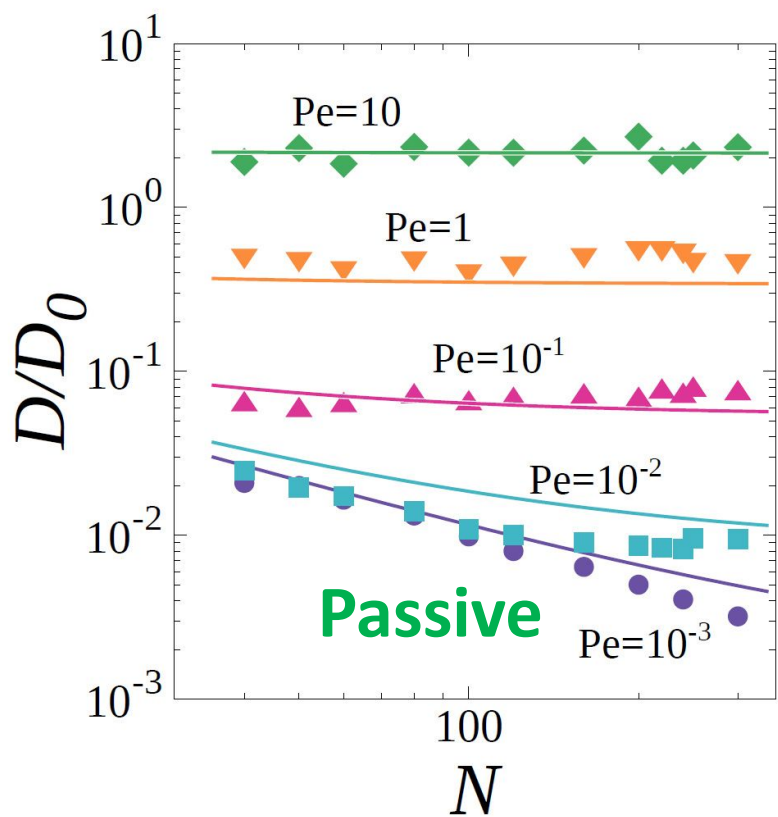


$$\frac{D_a - D_t^P}{D_t^P} = \frac{\tau_r^P D_t}{\sigma^2} \frac{R_e^2}{2N\sigma^2} \text{Pe}^2$$

$$\tau_r^P \propto \frac{N}{\text{Pe}}$$

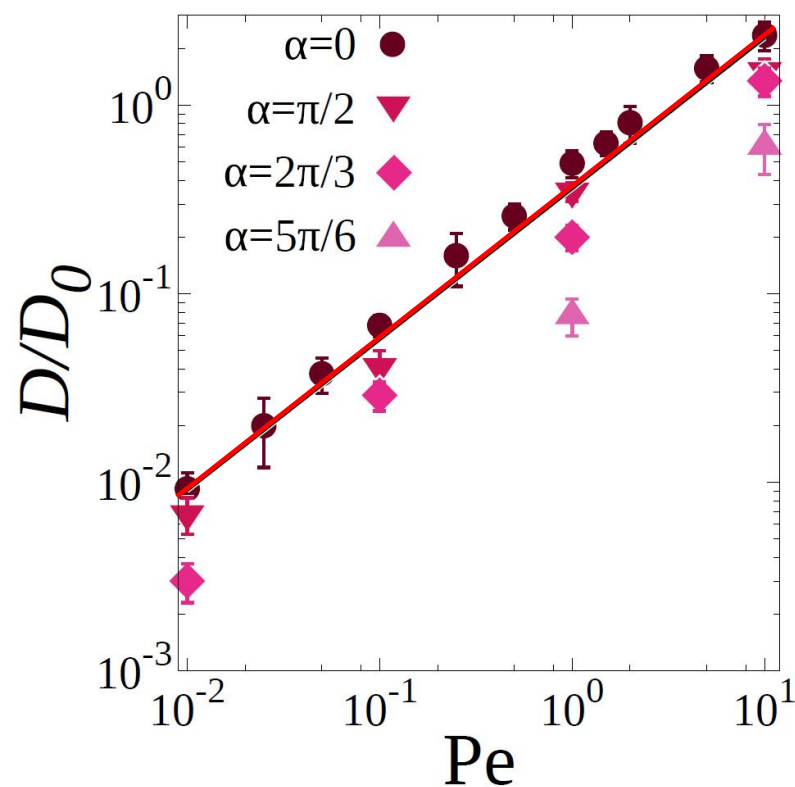
Polar Active Polymer

$$D = 2aN^{2\nu(\text{Pe})-1} + \frac{D_0}{N}$$



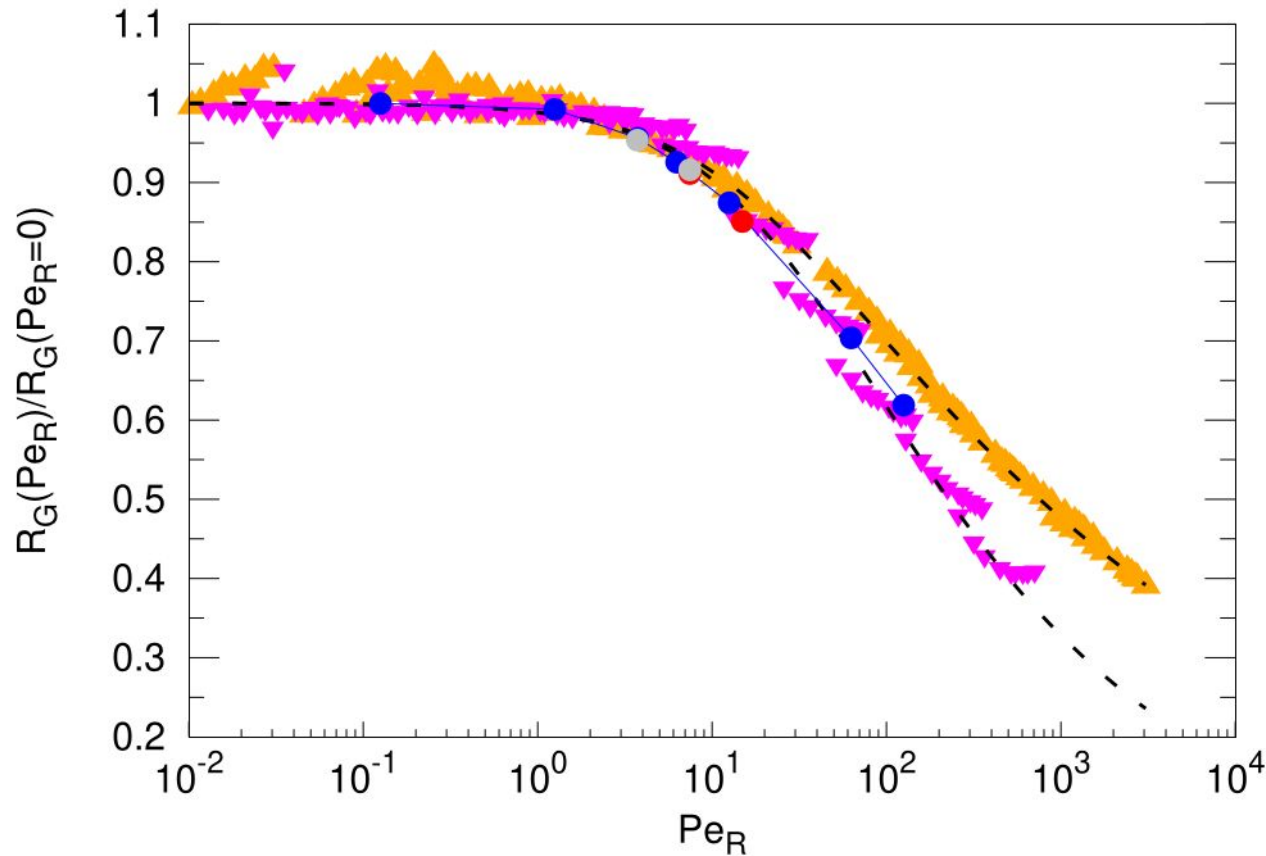
Active

Passive



Polar Active Polymer

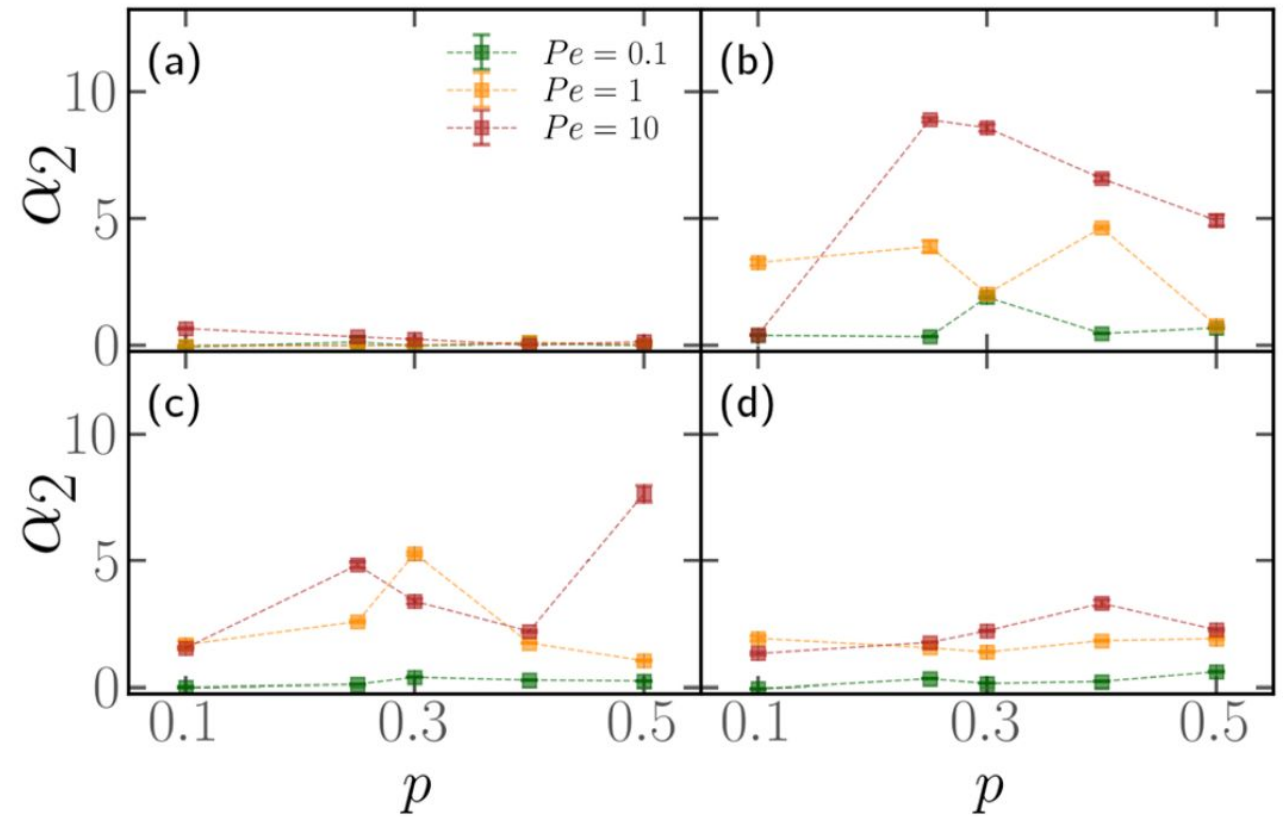
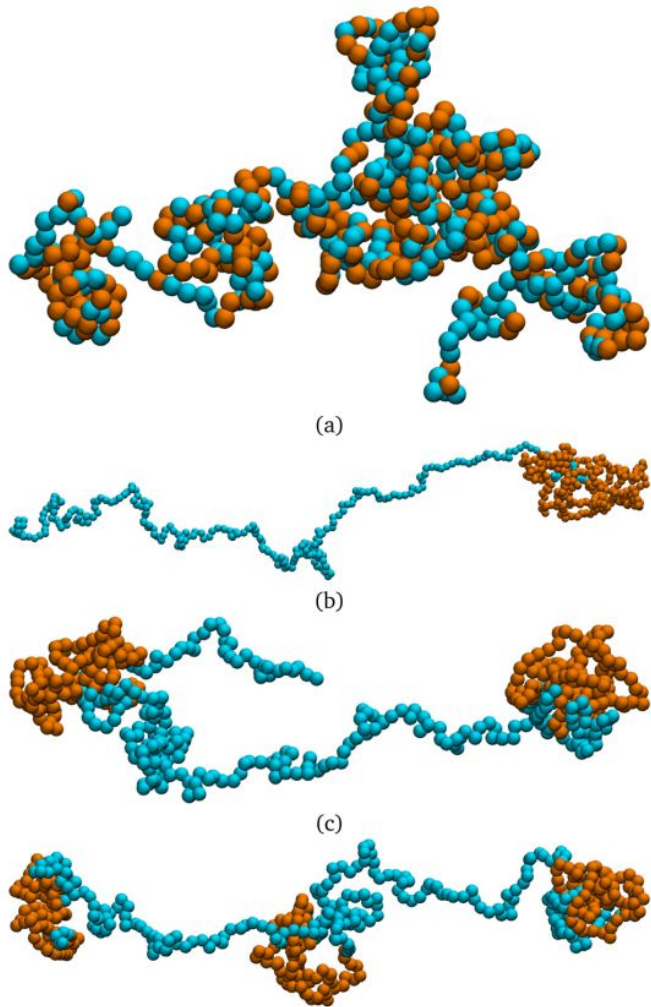
Active Rouse Model (with semi-flexibility)



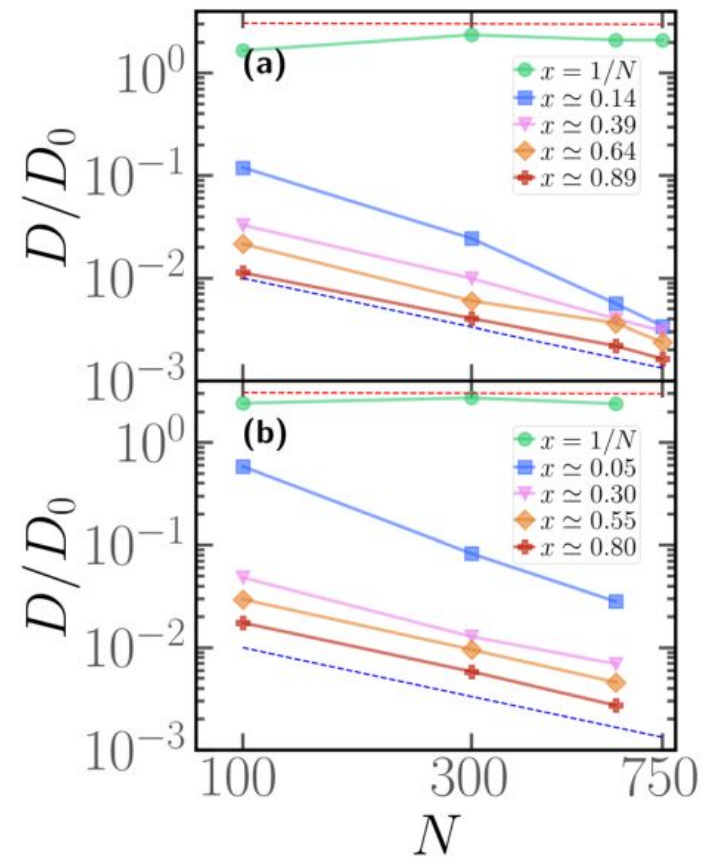
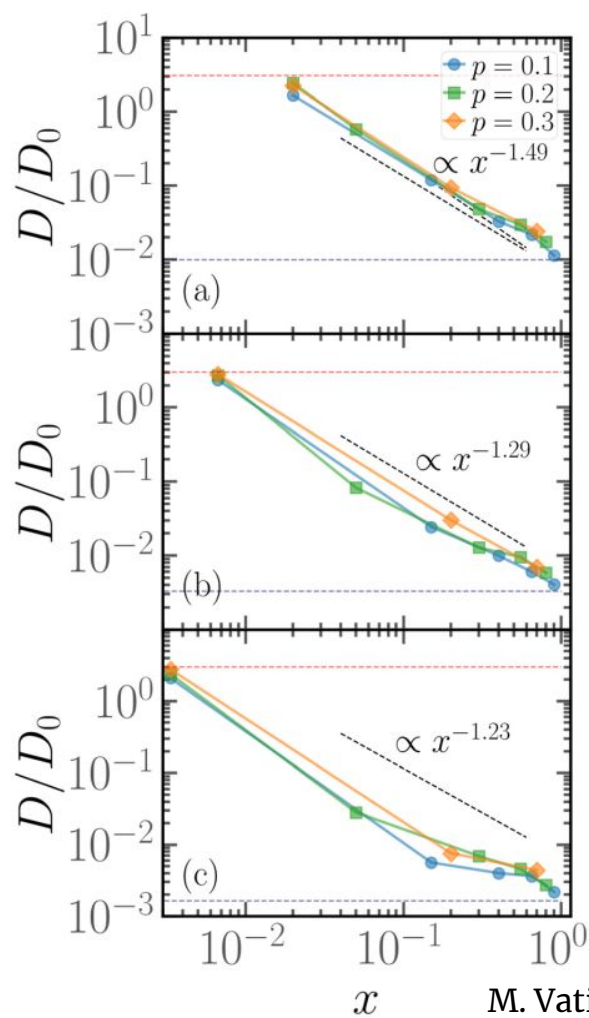
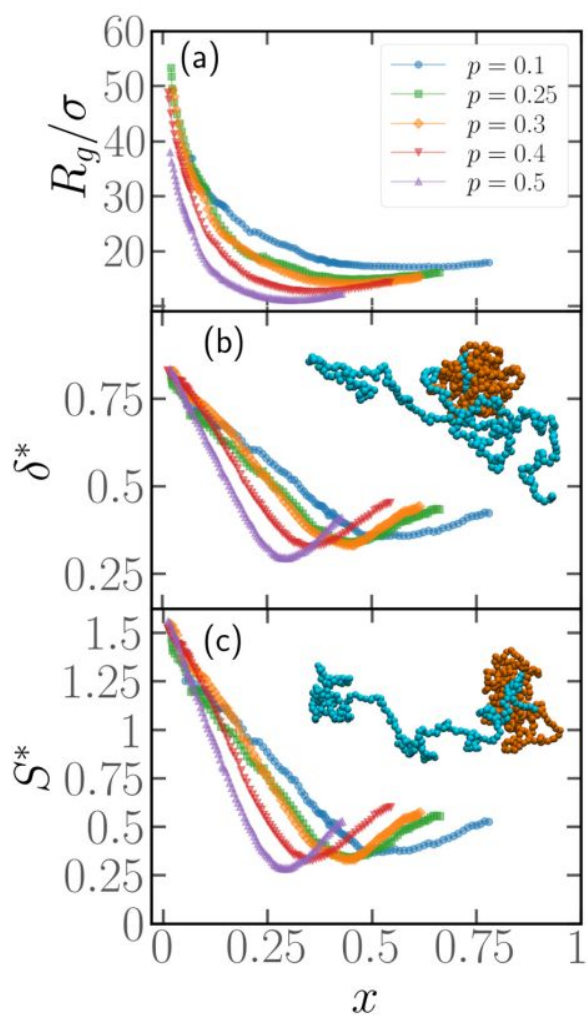
R_g is the relevant length scale of the system

$$\text{Pe}_R = \text{Pe} R_g(N, \text{Pe}=0)$$

Active Diblock Copolymer



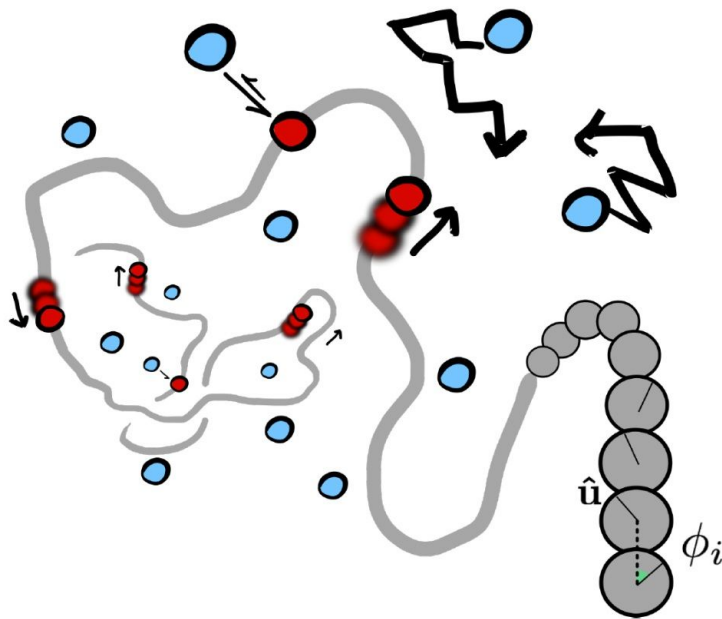
Active Diblock Copolymer



Motor-powered Polymer

Semi-flexible polymer substrate of length L

N Freely diffusing motors of size σ

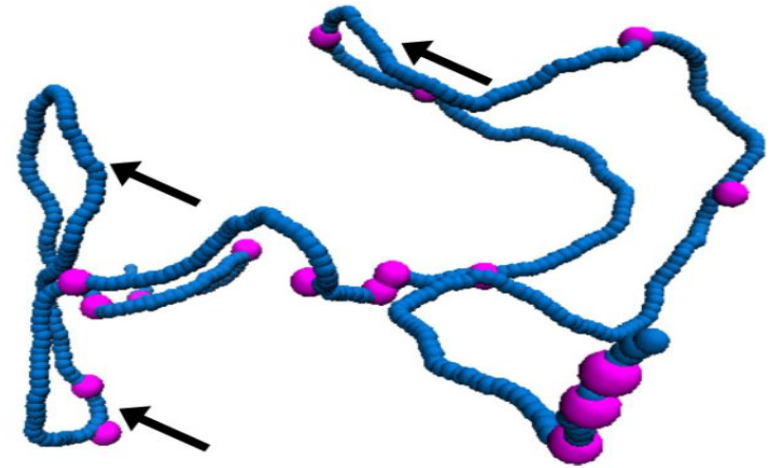
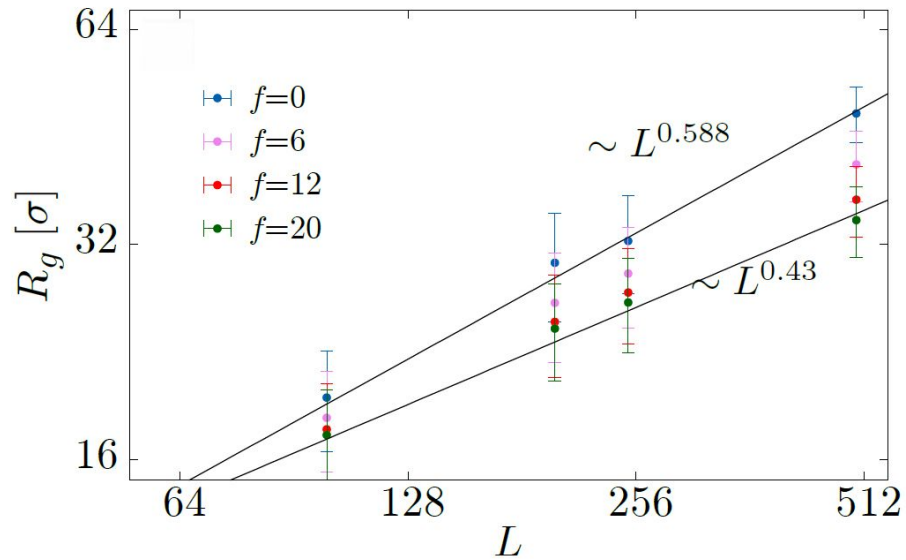


- Average number of bound motors $\langle n \rangle$
- Motor force f
- Binding energy K_{MOT}
- Substrate rigidity K_{BEND}

Freely diffusing motors bind to the substrate and propel themselves in the direction of the local bond vector, pushing the polymer in the opposite direction

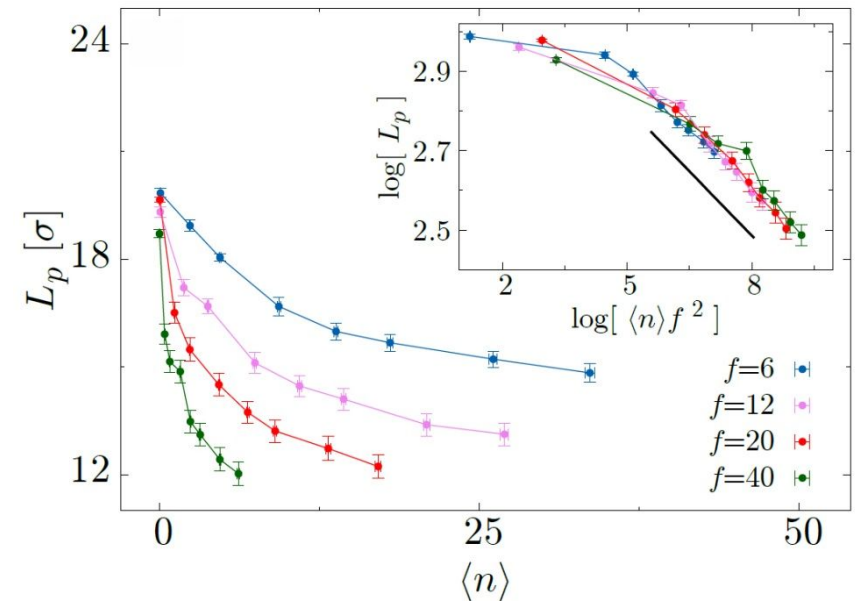
M. Foglino, et.al, *Soft Matter*, 2019

Motor-powered Polymer

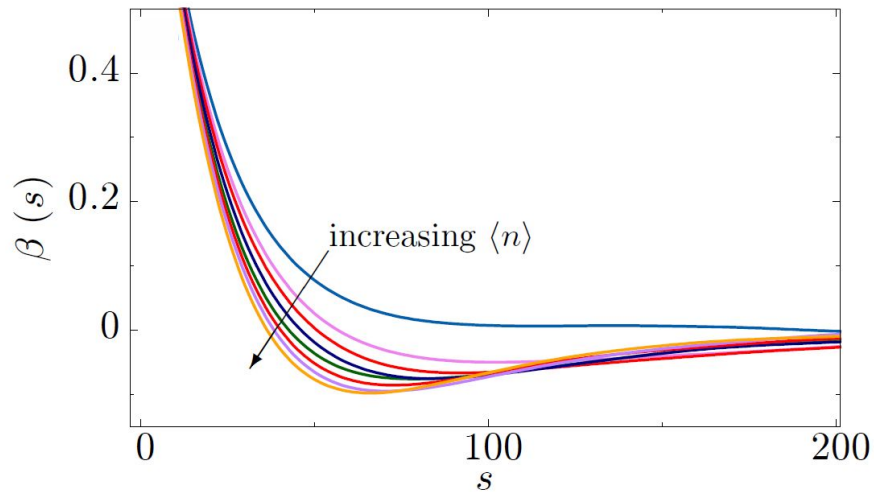


The scaling exponent decreases here too

The buckling instability manifests by the formation of U-shaped tight turns (hairpins) and the “softening” of the substrate

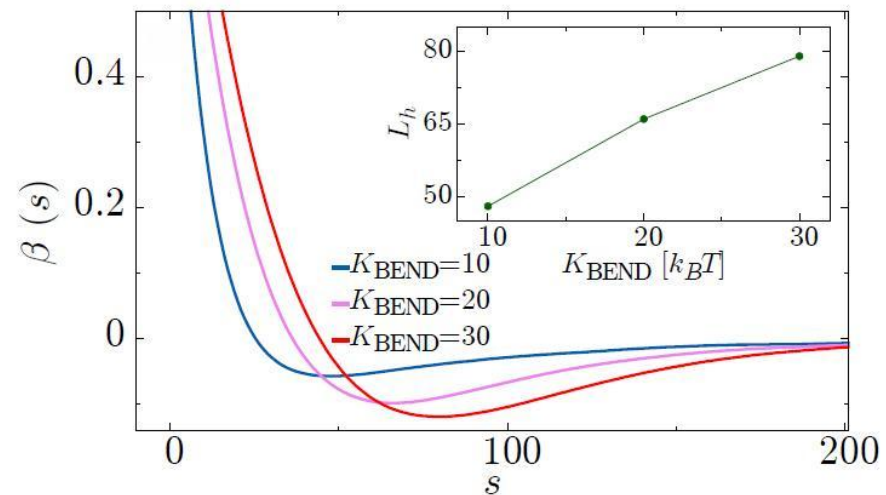


Motor-powered Polymer



$$\beta(s) = \langle \mathbf{b}_i \mathbf{b}_{i+s} \rangle$$

bond autocorrelation function



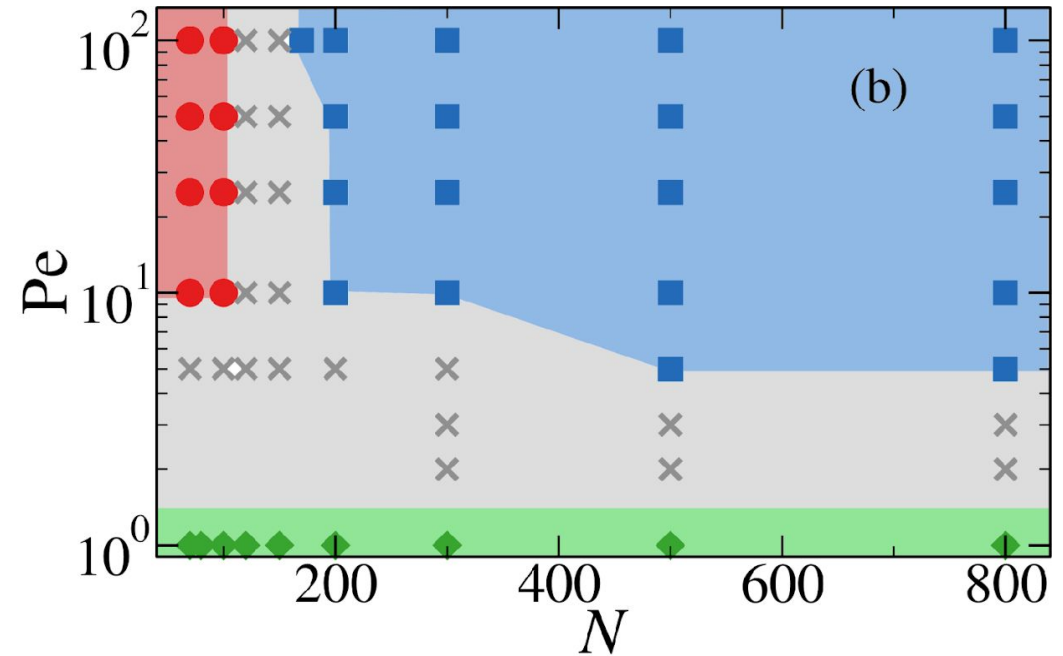
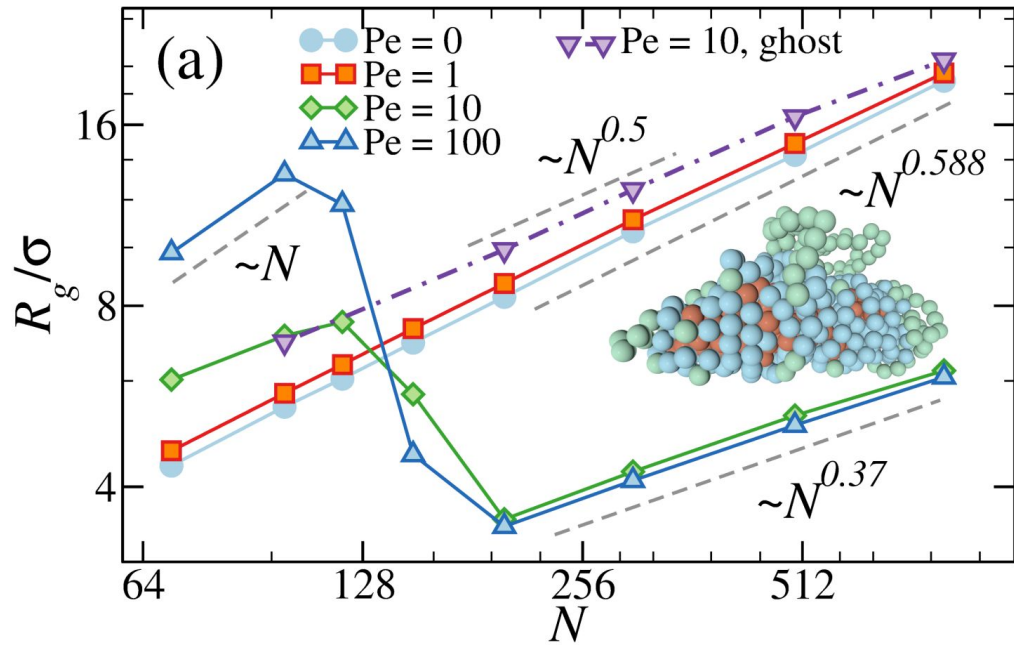
The bending instability manifests here by the formation of U-shaped rather tight turns (hairpins)

Ring Polymers

Why rings?

- Bacterial DNA is ring shaped; other unicellular parasites organize their DNA into a network of interlocked rings (kDNA)
- Ring polymers (solutions and melts) show distinctive properties, very different from their linear counterpart, despite there is a difference of a single bond
- The effects of circularization are of topological origin (topological repulsion)
- Chromosomes can be modeled as ring polymers

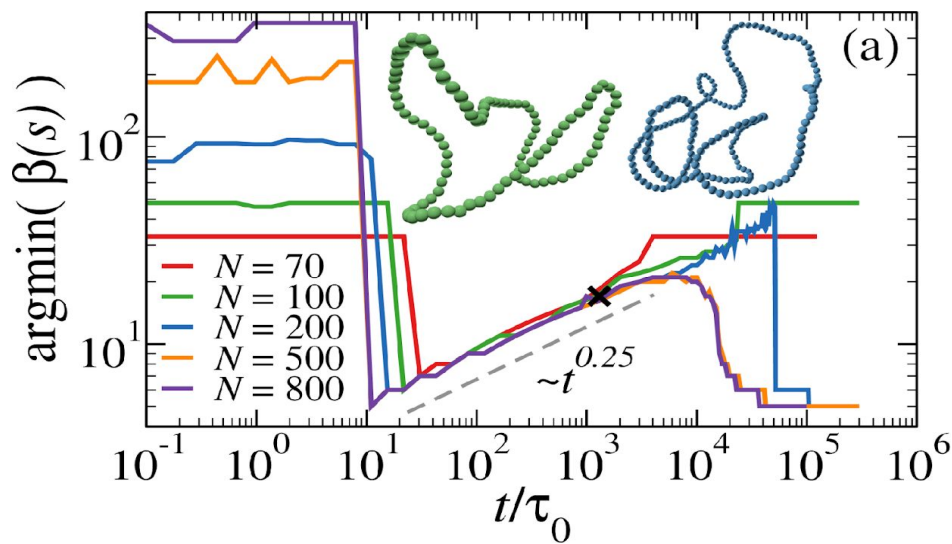
Active Ring Polymers



Swelling – collapse transition that becomes sharper at increasing Pe

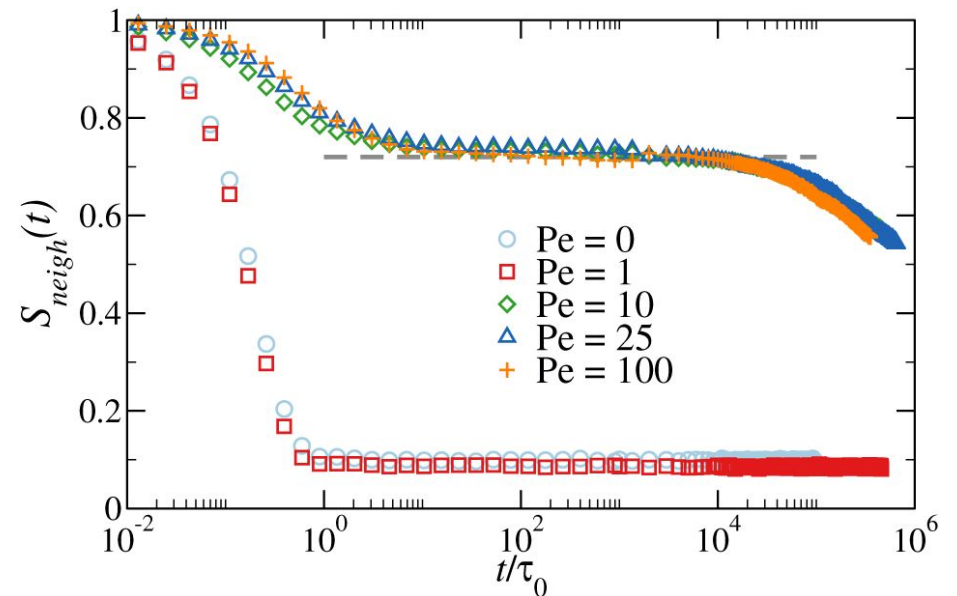
Active Ring Polymers

Kinetics



Non-equilibrium route from passive conformation to steady state is independent of N

Dynamics



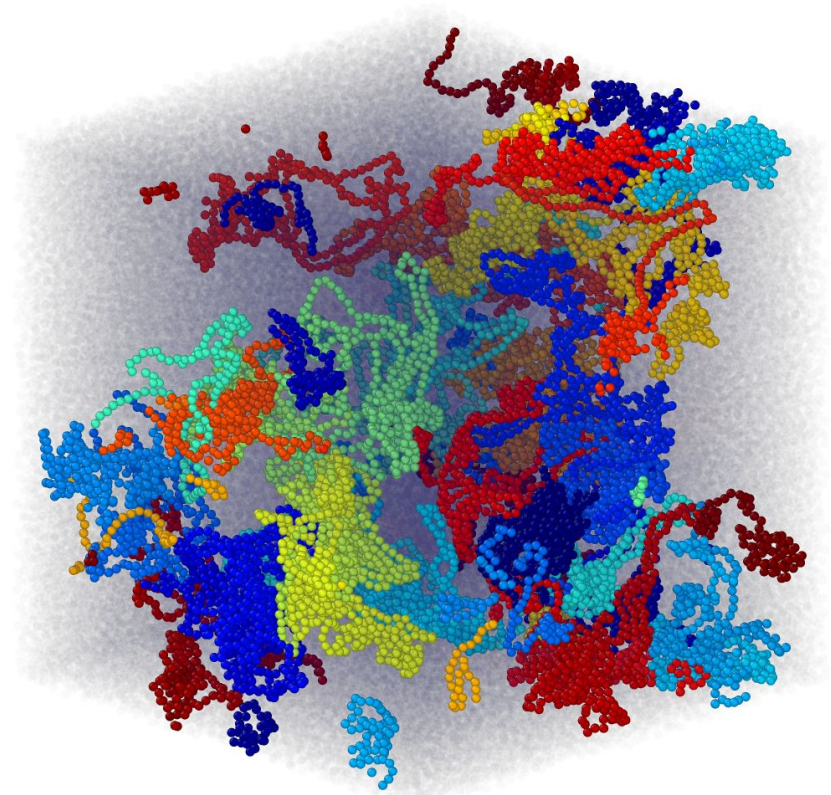
In the steady state, collapsed rings show signs of dynamical arrest in many observables

Outline

Polar Active Polymers

**Multi-scale analysis of
dense suspensions**

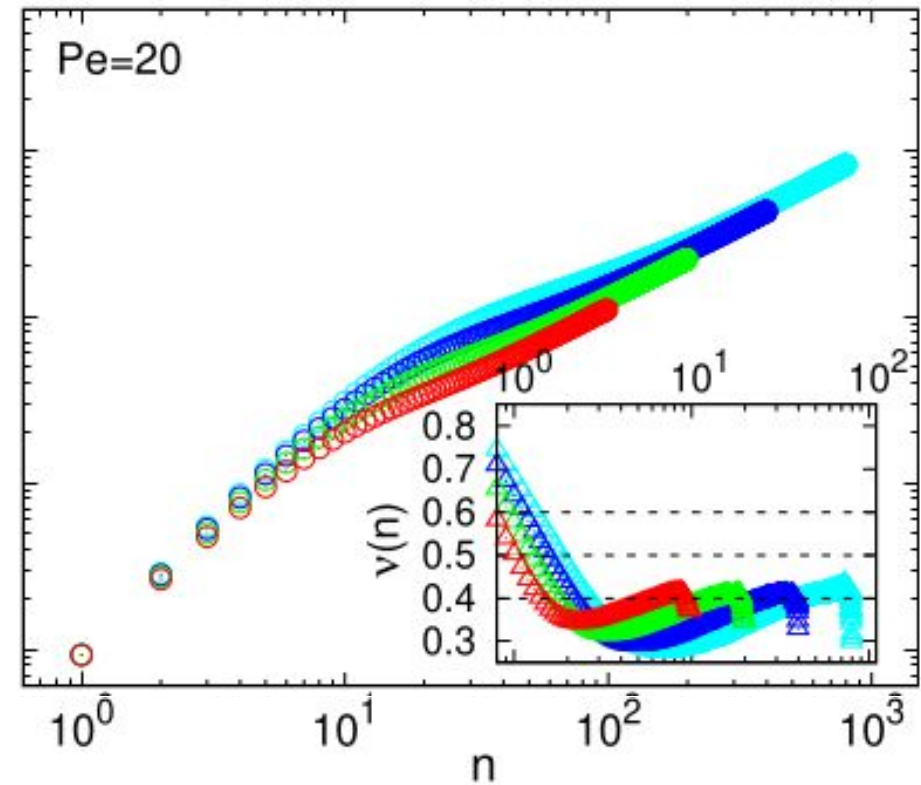
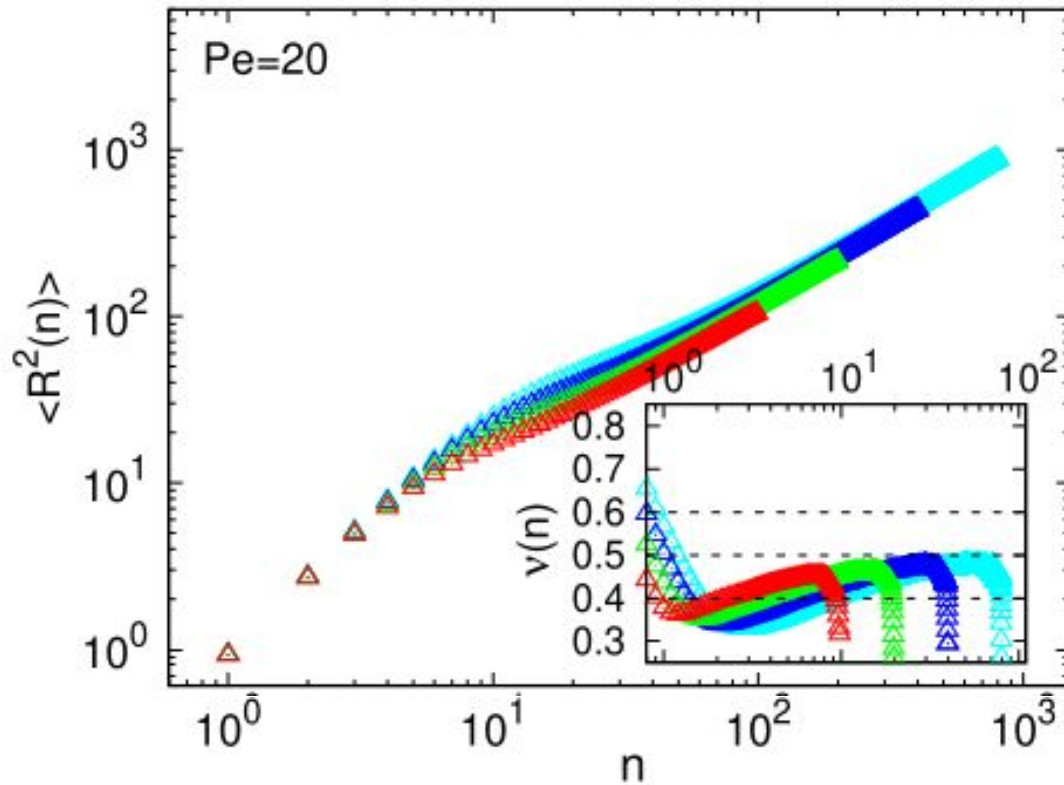
Towards modeling biophysical systems



Melts of Active Linear Chains

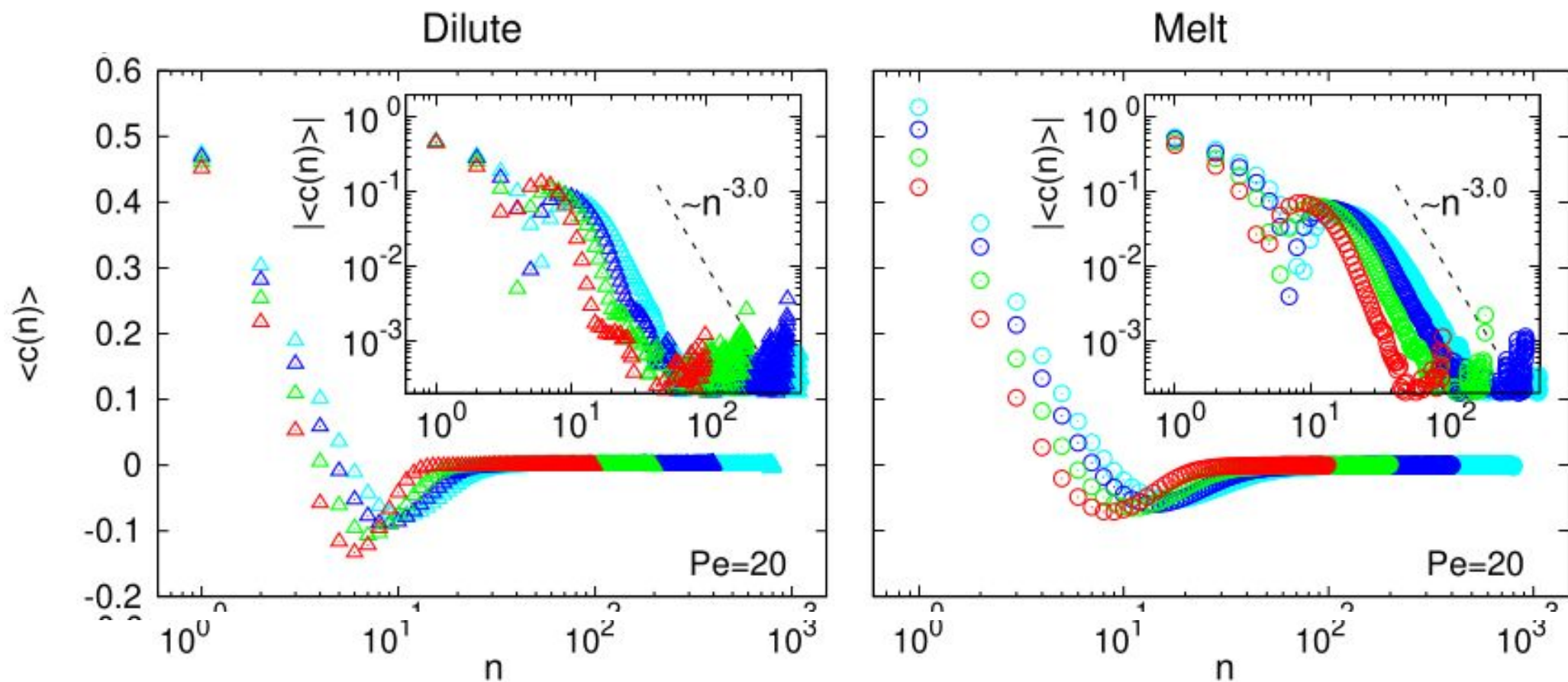
Dilute

Melt



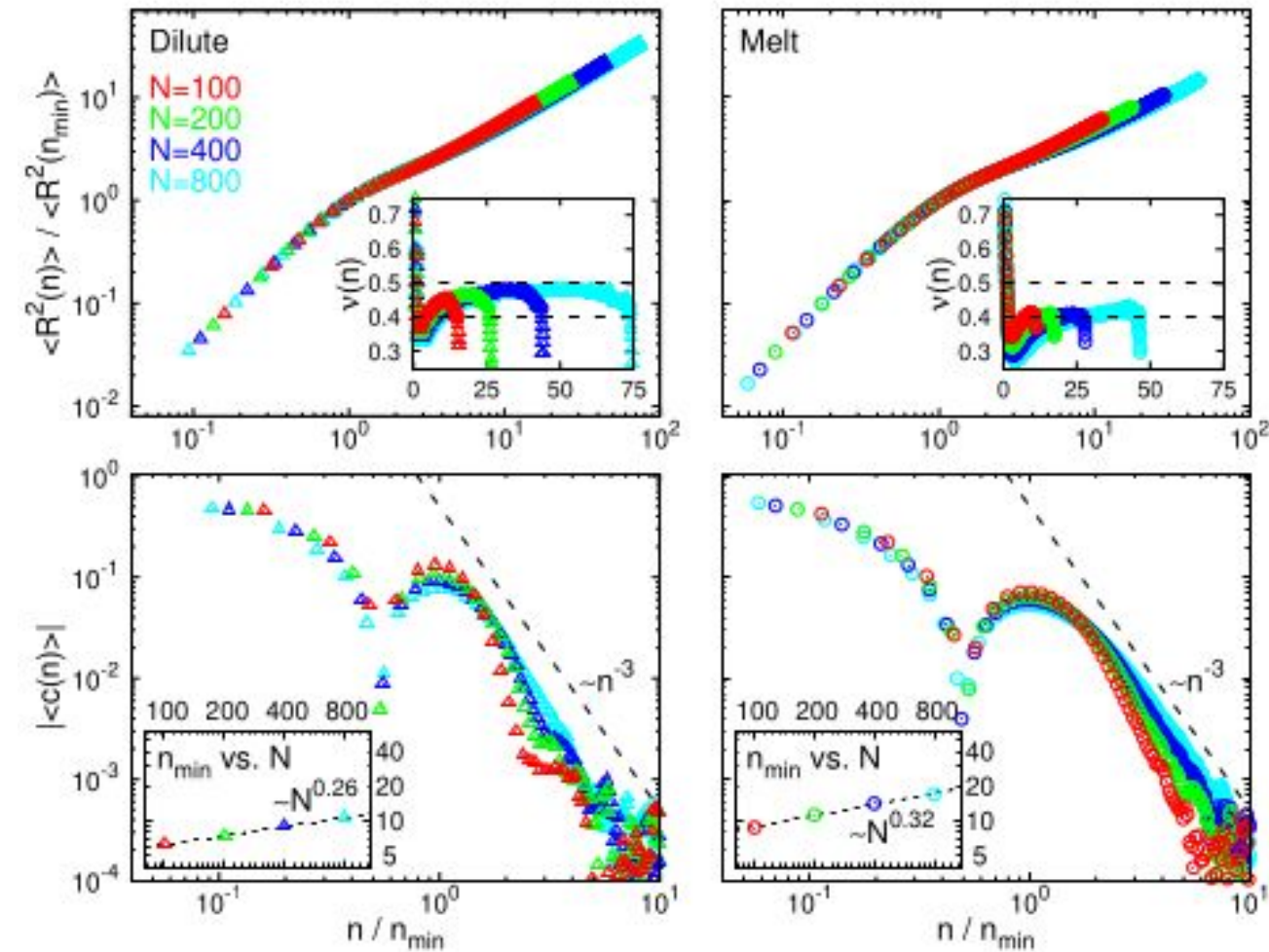
$$\nu(n) \equiv \frac{1}{2} \frac{\log(\langle R^2(n+1) \rangle) - \log(\langle R^2(n-1) \rangle)}{\log(n+1) - \log(n-1)}$$

Melts of Active Linear Chains



$$\langle c(n) \rangle \equiv \frac{\langle \vec{t}_{n'+n} \cdot \vec{t}_{n'} \rangle}{\langle \vec{t}_{n'}^2 \rangle}, \quad n_{min}(Pe, N)$$

Melts of Active Linear Chains

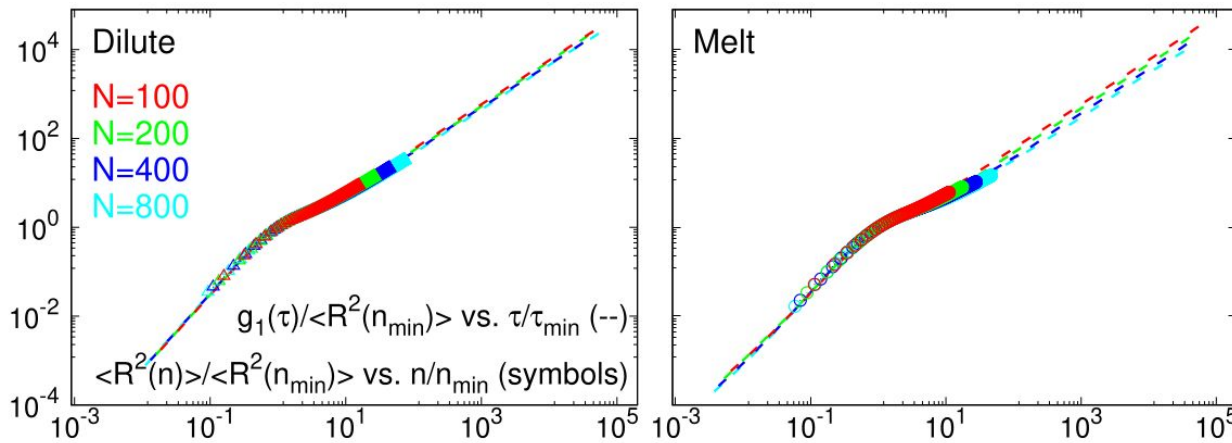


$n_{min}(Pe, N)$ is the length that marks the minimum of the tangent correlation function.

It is a characteristic length scale of the system both at infinite dilution and in the melt.

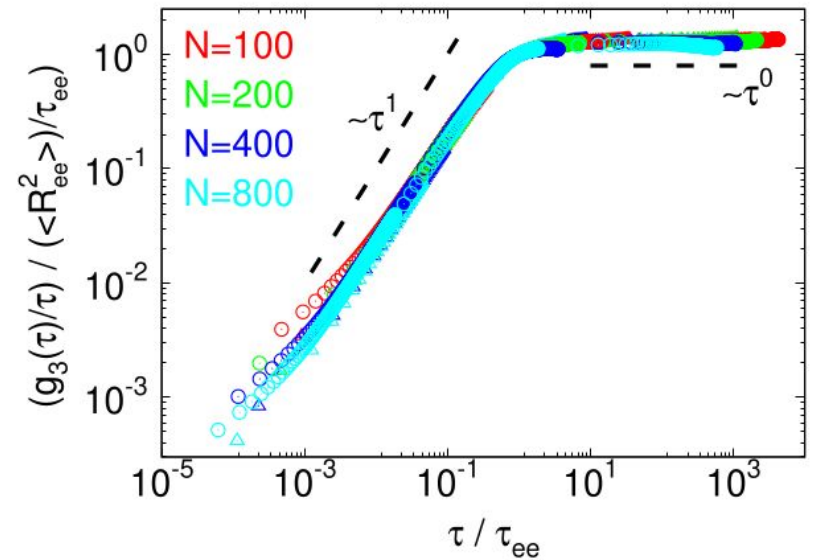
Melts of Active Linear Chains

Dynamics: two different length scales

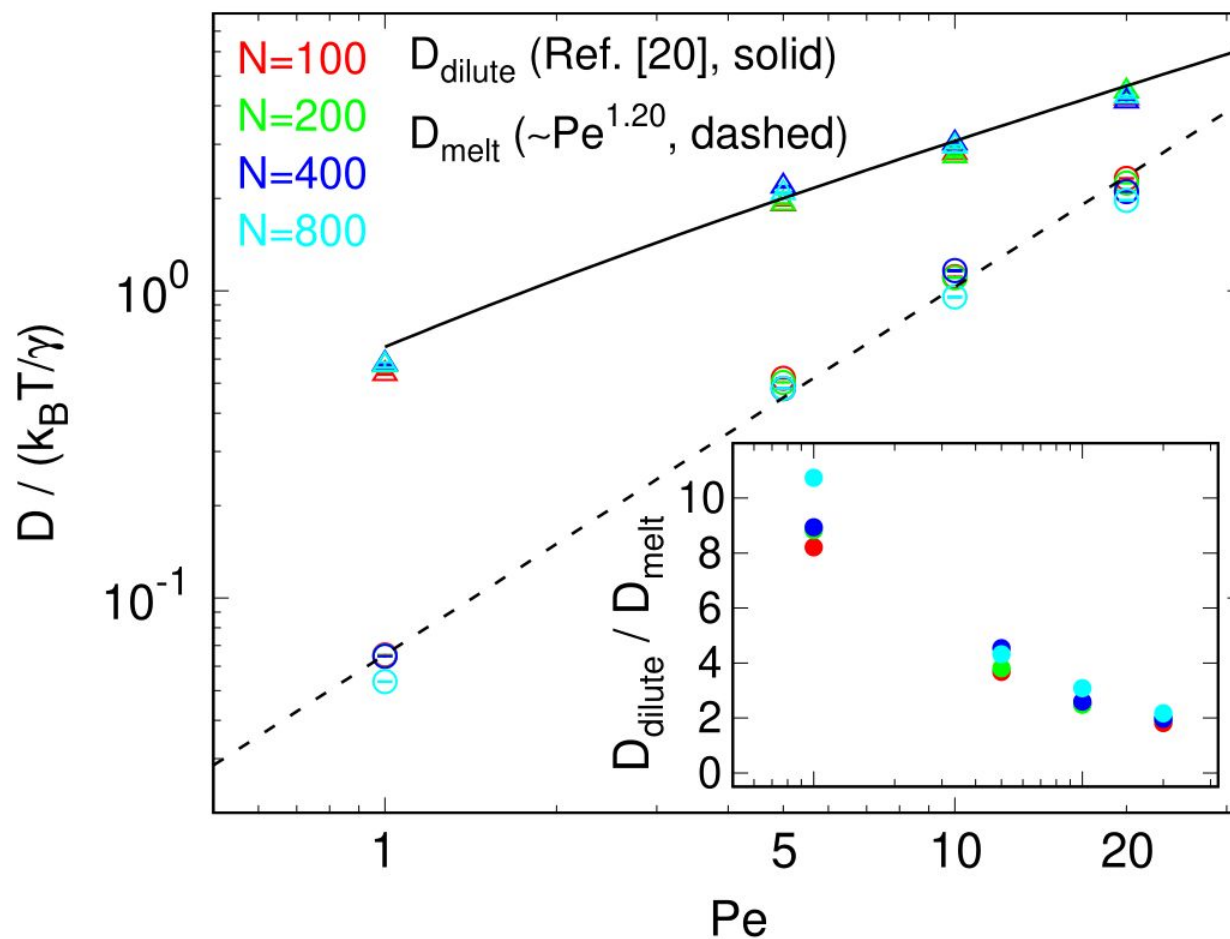
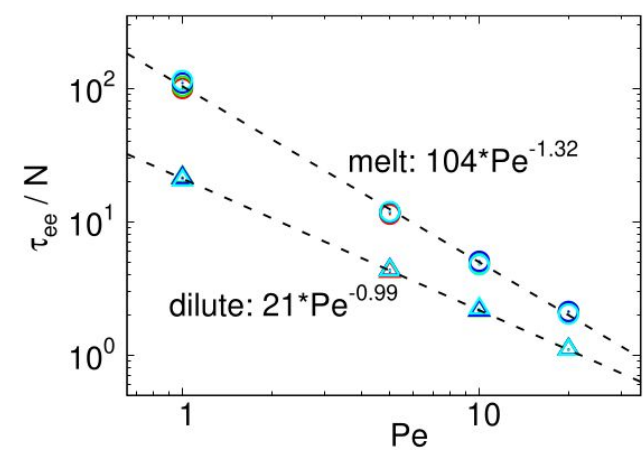
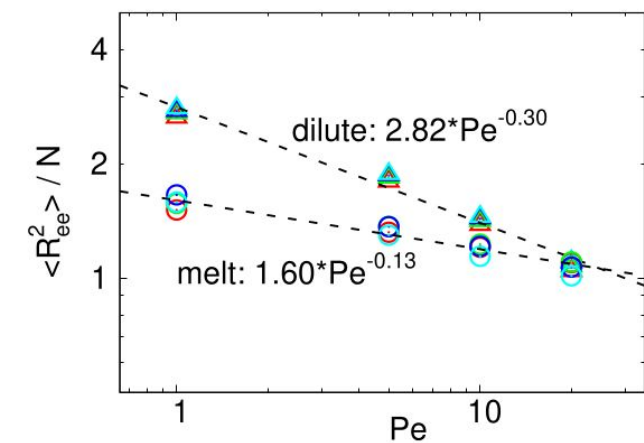


Instead, R_{ee}^2 is the characteristic scale of the centre of mass dynamics, with a characteristic time scale τ_{ee}

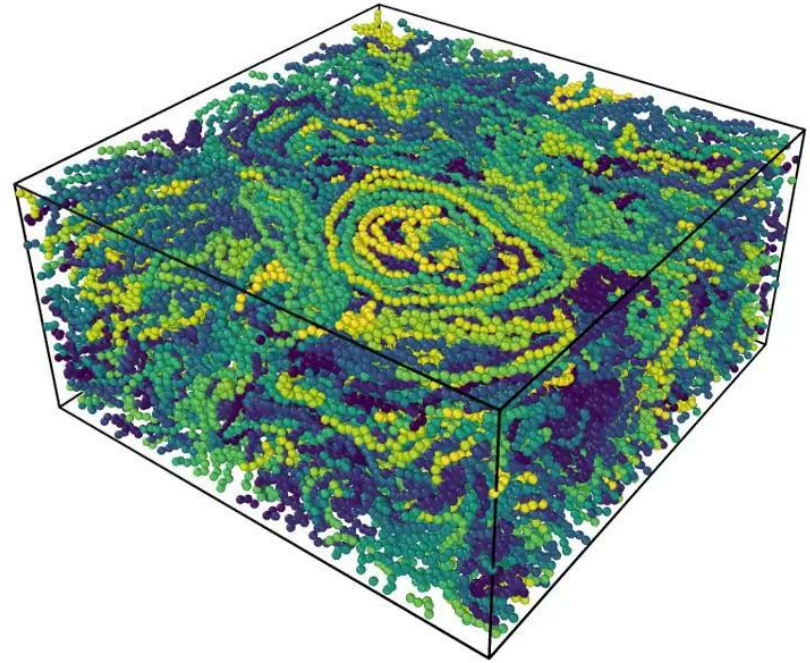
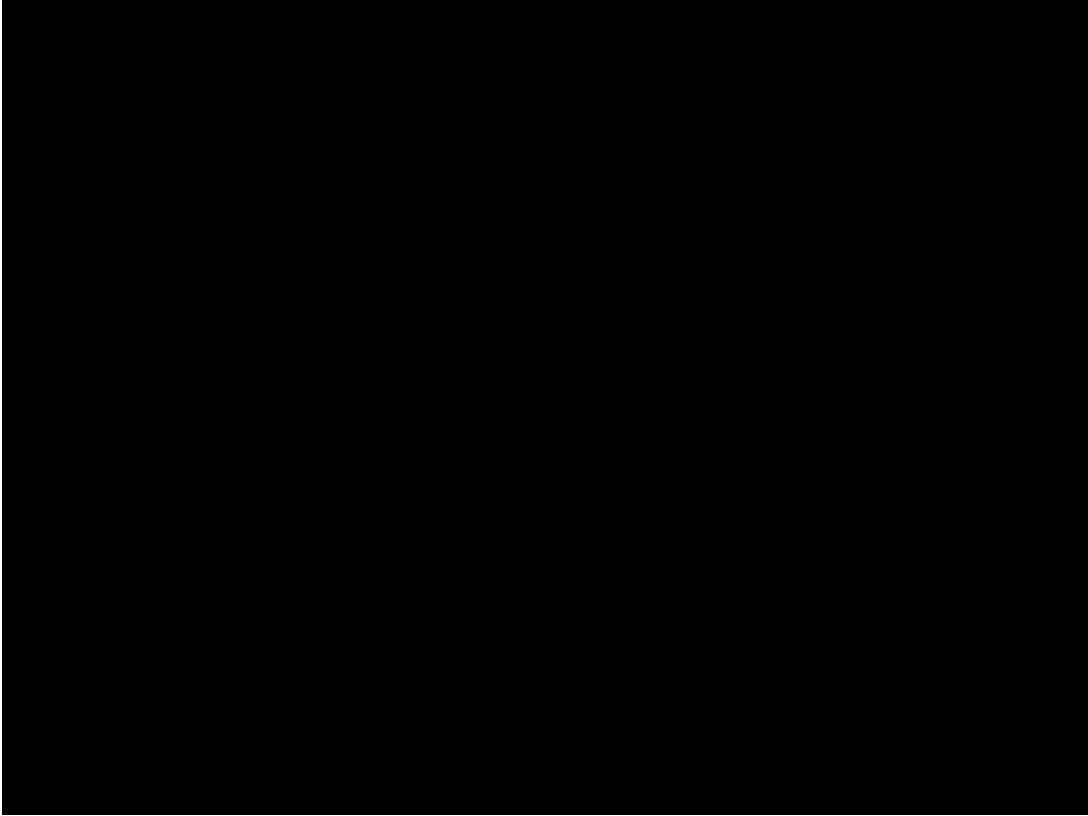
$n_{min}(Pe, N)$ is the characteristic scale of the monomer dynamics



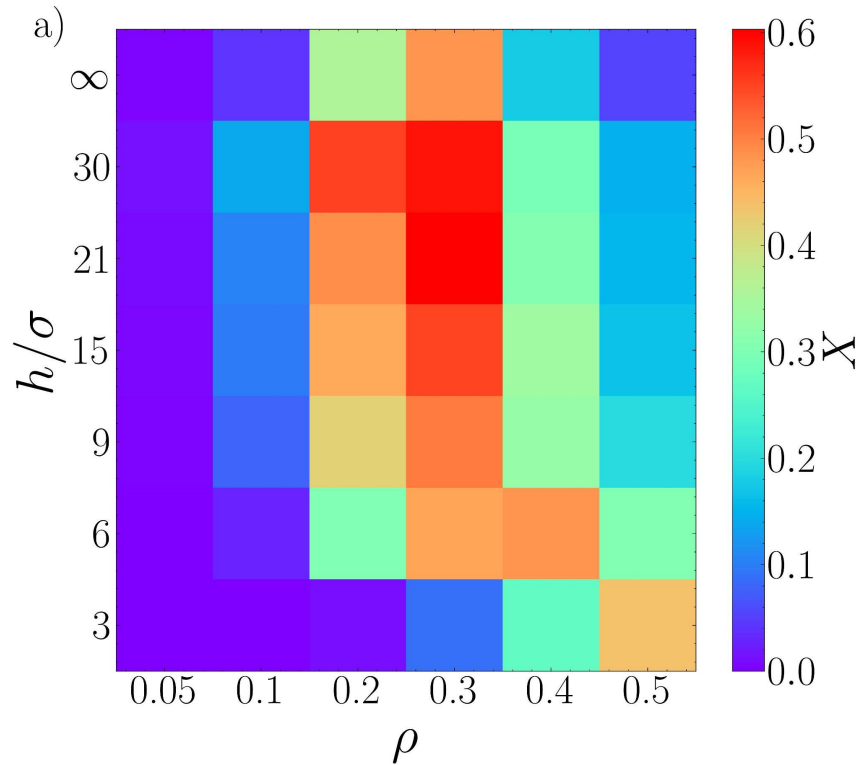
Melts of Active Linear Chains



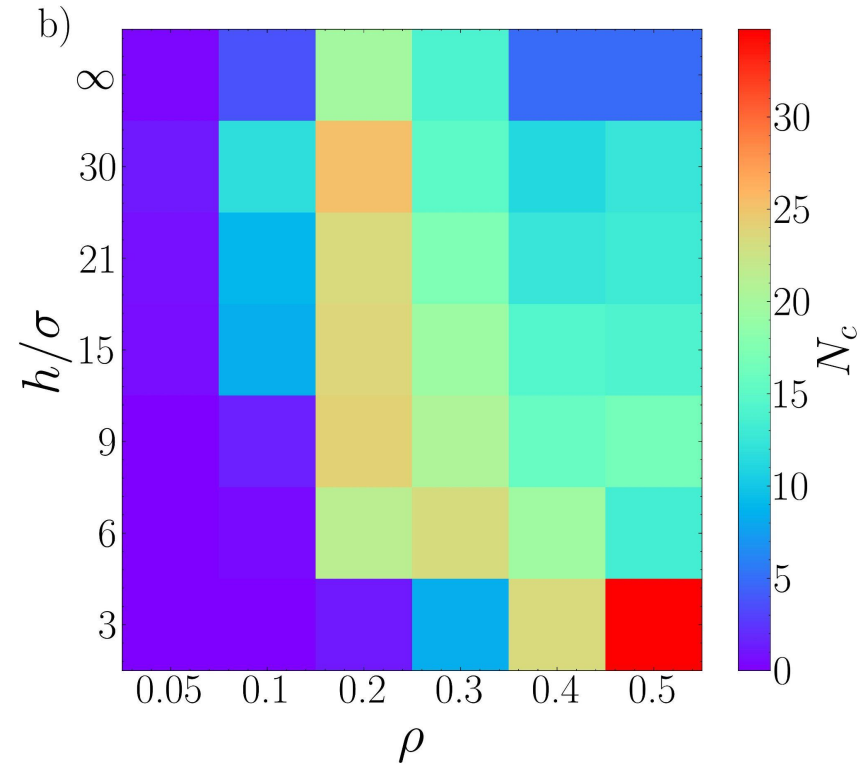
Active Rings under confinement



Active Rings under confinement



Fraction of clustered rings X



Number of clusters N_c

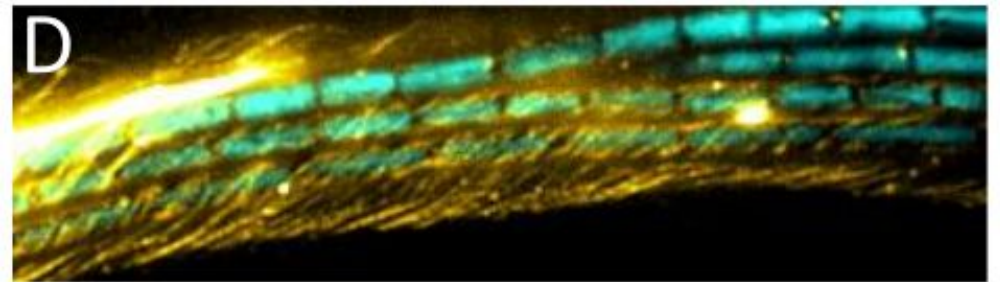
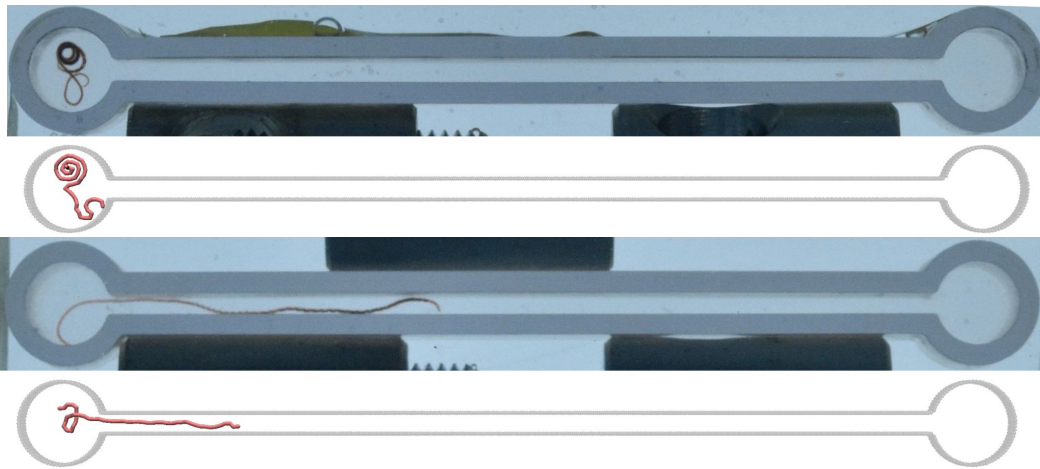
There is a rather large area of the ρ - h plane in which these bundles form;
too high density suppresses organization

Outline

Polar Active Polymers

Multi-scale analysis of dense suspensions

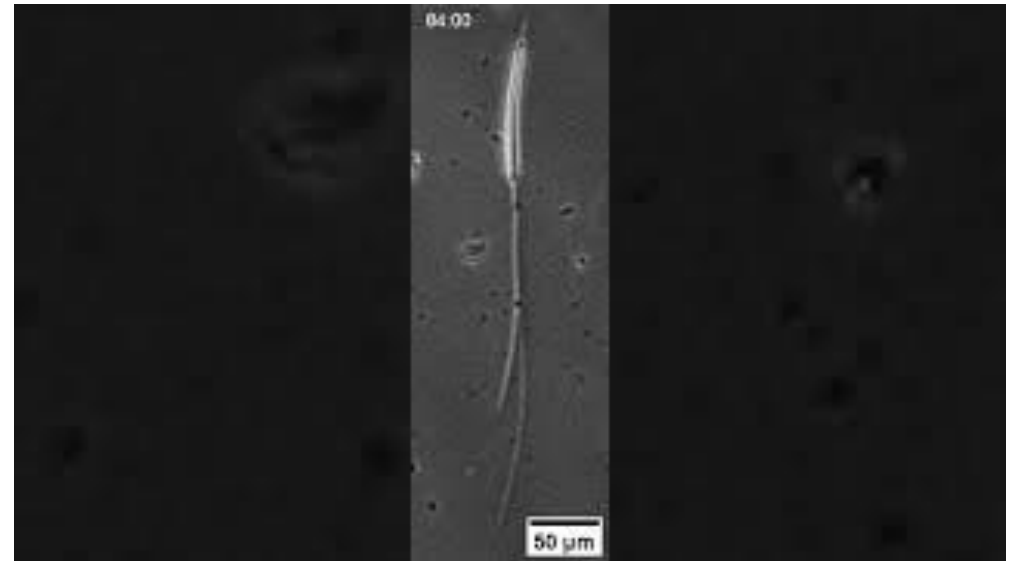
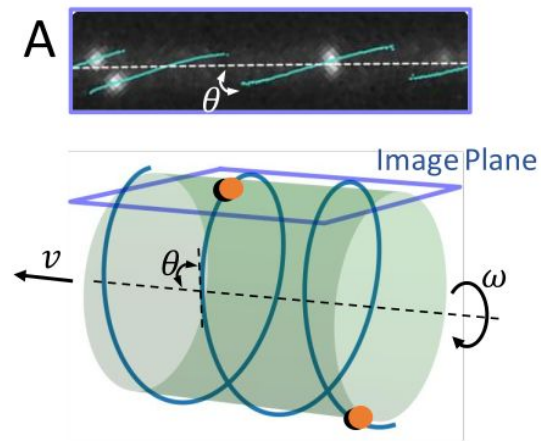
Towards modeling biophysical systems



Cyanobacteria reversals

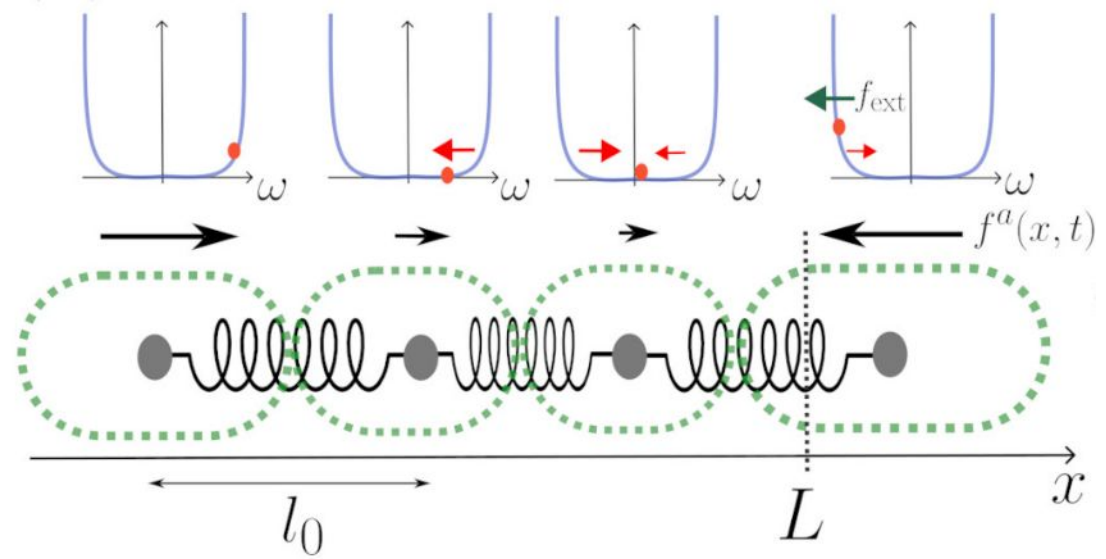
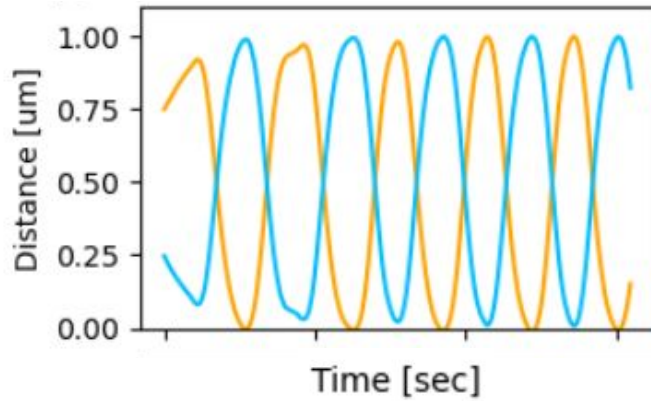


Can we come up with a (local) model for the mechanism of reversals?



J. Rosko, K. Cremin, E. Locatelli, et al, bioRxiv, 2024.02.06.579126

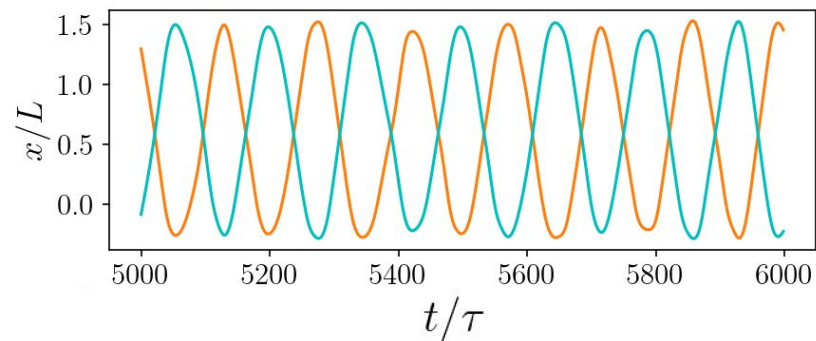
Cyanobacteria coordination model



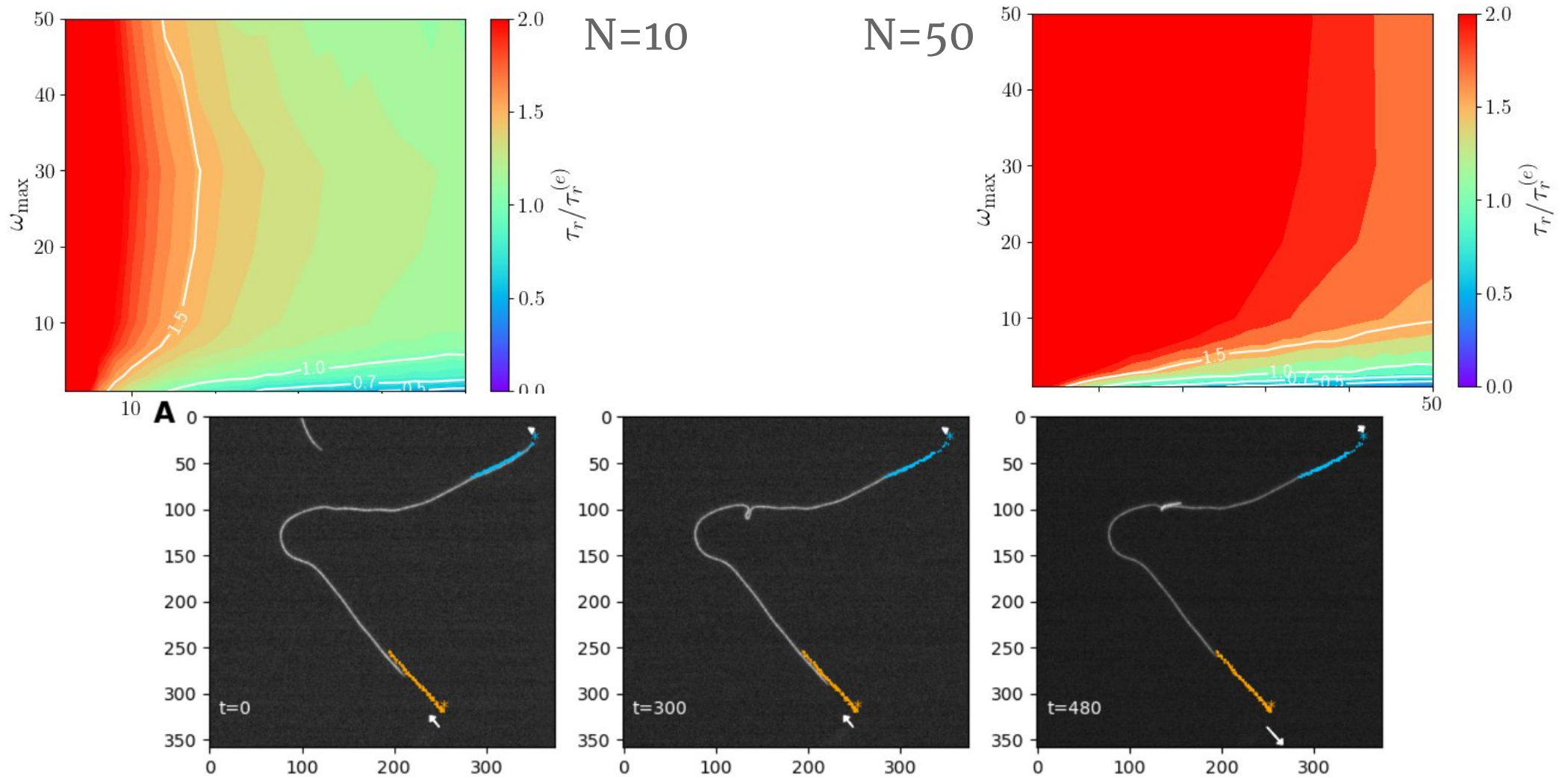
Local (mechanical) sensing mechanism drives coordination

$$\dot{x}_i = \frac{1}{\gamma} \left(-\frac{dV}{dx_i} + f_i^a \cdot s_i(x, t) \right) + \xi_i(t)$$

$$\dot{\omega}_i(x, t) = \frac{1}{\gamma_\omega} \left(-\frac{dU_c}{d\omega_i} - \frac{dV_\omega}{dx_i} + f_{\text{ext}}(x, t) \right) + \eta_i(t)$$

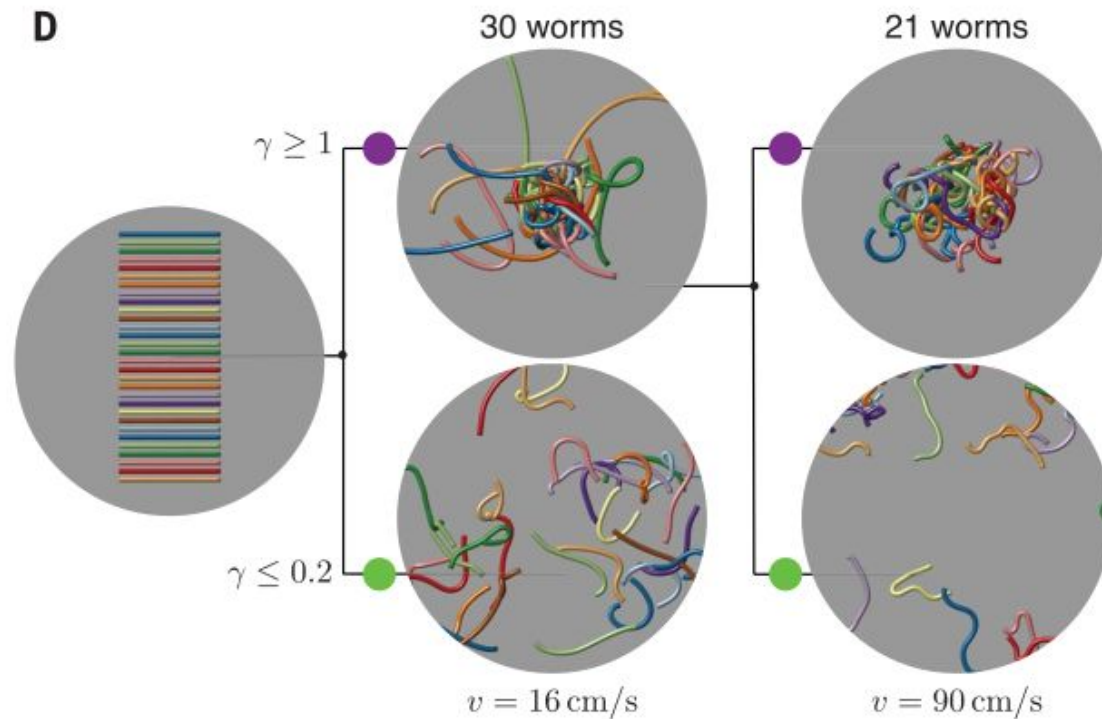


Cyanobacteria coordination model



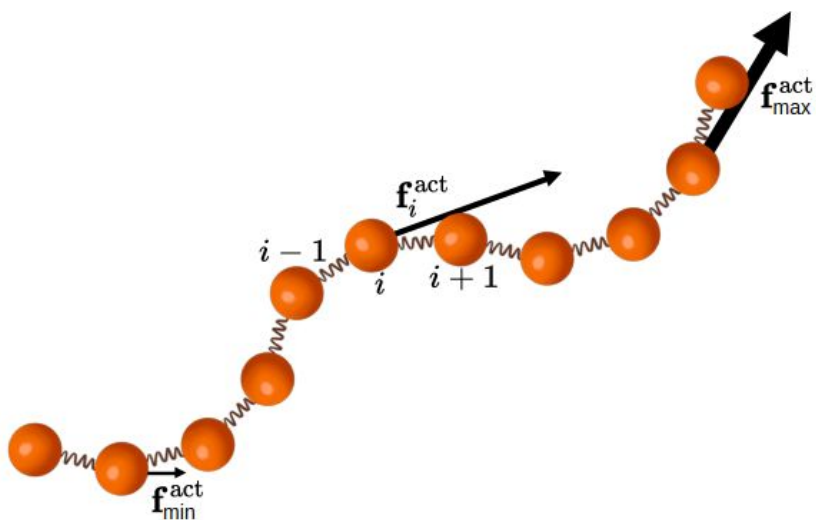
Modeling worms under confinement

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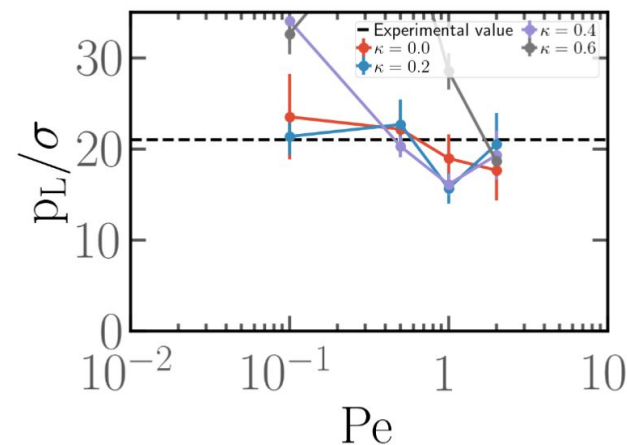
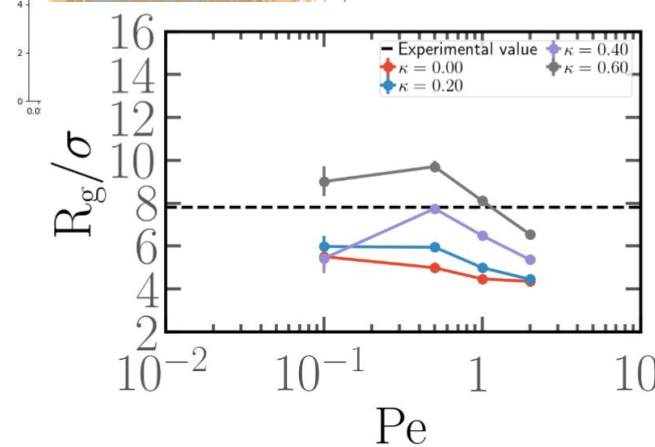
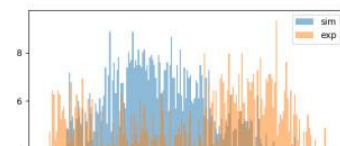
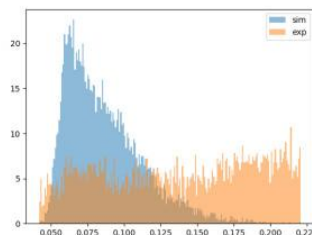
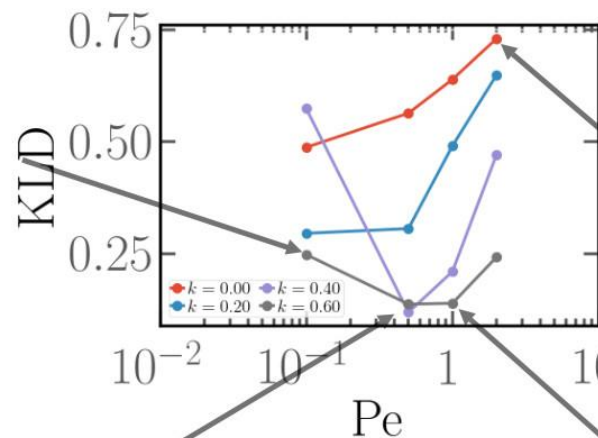
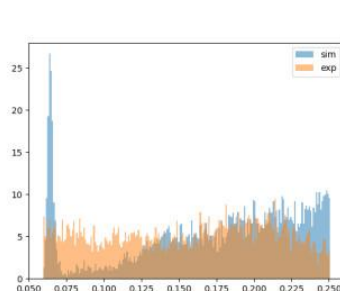
Does a simpler (and more transferable) model exist?

Modeling worms under confinement



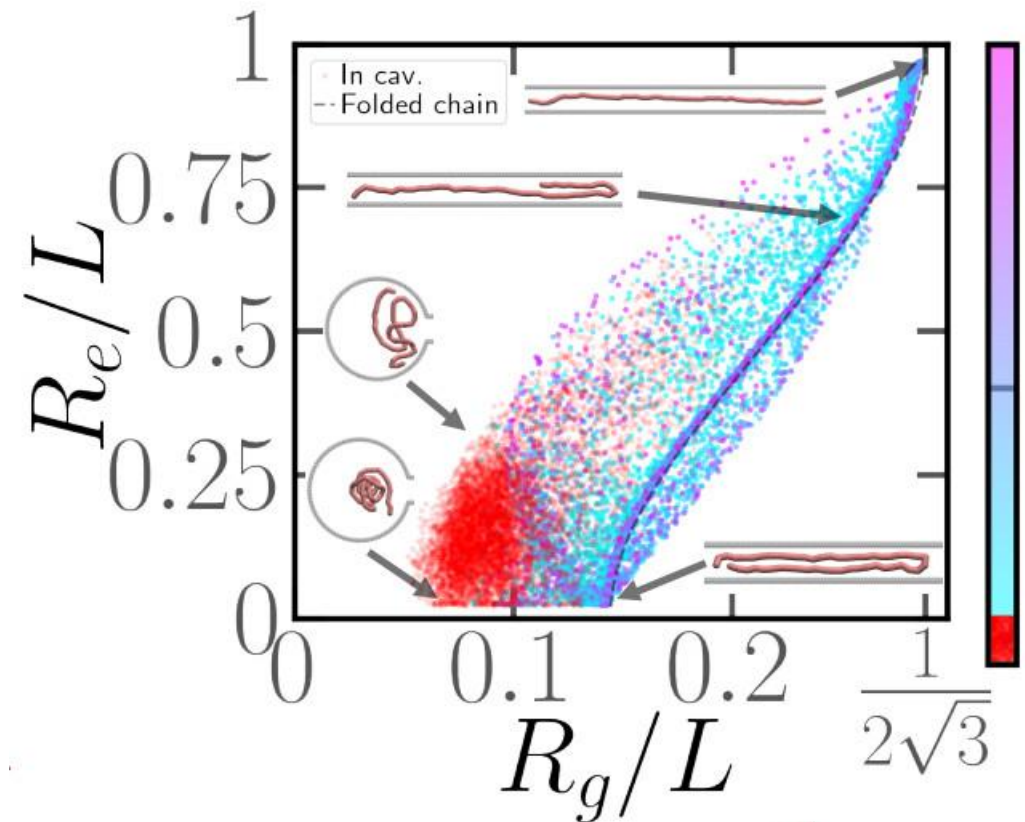
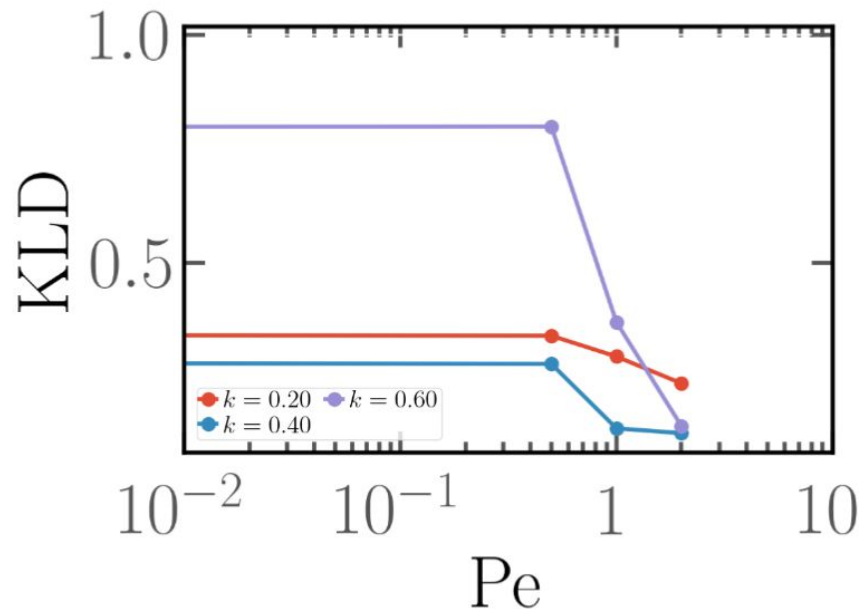
Semi-flexible polymer
 Non-homogeneous activity
 profile
 3D+gravity

Bulk



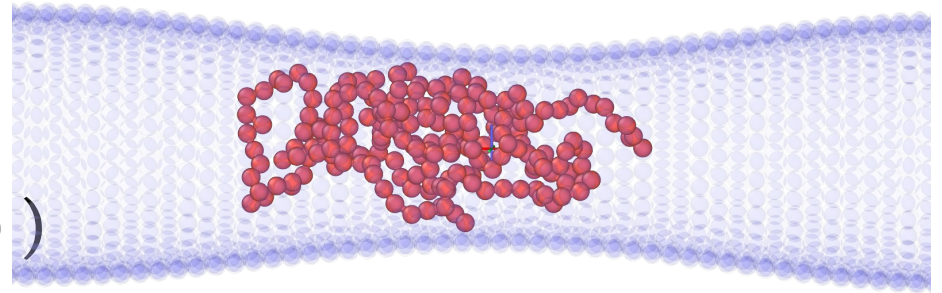
Modeling worms under confinement

Confinement



Perspectives

- Linear polymers in cylindric and corrugated channels
(Phys. Rev. Lett. 131, 048101 (2023))
- Non-homogeneous activity
- Knots & entanglements
- Multi-scale modeling micro/macro-organisms
 - 3D model of reversing bacteria (plectoneme formation)
 - assembly of bacteria (effect of external bias)
 - learning the worm model



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P. Margaretti (Helmholtz Institute Erlangen)



C. Valeriani, J. P. Miranda, J. Martin Roca (UCM)



TECHNISCHE
UNIVERSITÄT
WIEN
Vienna University of Technology

C. N. Likos (Univie)

D. Michieletto, C. Brackley, D. Marenduzzo (Edinburgh)



M. Vatin, E. Orlandini (Padova)

S. Kundu, M. Ubertini, A. Rosa (SISSA)

O. Soyer (Warwick), M. Polin (IMEDEA)



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