# Dynamical phase transitions in complex fluids and nonequilibrium systems

## WP1 Spoke 2 b5 Physics of Complex Systems (UniBa)

- Ordering properties in complex fluids and non-equilibrium systems Active fluids (dense suspensions of bacteria, cytoskeletal extracts, Microtubules bundles)
- Hydrodynamic and Statistical mechanics modelling
- High Performance Computing

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### Phase Field Theories of complex and active fluids



#### Topological Phase transitions of Choleric Shells Confined in Shells

#### Self-Propulsions of active droplets in 3D

Active Turbulence

**Lattice Boltzmann Methods** MPI



#### Non-equilibrium statistical models for self propelled particles and DNA transcription

Non equilibrium Phase transitions in Active Brownian Particles

Polymer models to study transcription and dynamics in Human Chromosomes

> **Molecular dynamics simulations** with implicit and explicit solvents (LAMMPS+ CustomMulti Particle collision Dyanamics Package)

# PHASE FIELD THEORIES





Oriental Order

Smooth Deformations

**Described by a second rank** traceless symmetric Tensor **Order Parameter** 

$$Q_{\alpha\beta} = S \langle \mathbf{n}_{\alpha} \mathbf{n}_{\beta} -$$

## liquid crystals









## Landau-De Gennes Theory

**Tensor order parameter First order transition for orientational order** 

$$\mathscr{F}\left[\phi, Q_{\alpha\beta}\right] = \int dV \left[A_0\left[\frac{1}{2}\left(1 - \frac{\chi(\phi)}{3}\right)\mathbf{Q}^2 - \frac{\chi(\phi)}{3}\mathbf{Q}^3 + \frac{\chi(\phi)}{4}\mathbf{Q}^4\right] + \frac{L}{2}\left[(\nabla \cdot \mathbf{Q})^2 + (\nabla \times \mathbf{Q} + 2q_0\mathbf{Q})^2\right]\right]$$





3 Control Parameters

Chirality Elastic energy

Activity

# **DYNAMICAL EQUATIONS**

### **Dynamical fields**

• Nematic tensor  $Q_{\alpha\beta}$  • Veloc

Advection relaxation for the tensor order parameter  $(\partial_t + \mathbf{v} \cdot \nabla)\mathbf{Q} - \mathbf{S}(\mathbf{W}, \mathbf{Q}) = -\frac{1}{\Gamma}\mathbf{H}$   $\mathbf{H} = -\frac{\delta\mathcal{F}}{\delta\mathbf{Q}} + \frac{\mathbf{I}}{3}Tr\left(\frac{\delta\mathcal{F}}{\delta\mathbf{Q}}\right)$ 

**Advection diffusion for concentration field** 

$$\partial_t \phi + \nabla \cdot (\phi \mathbf{v}) = \nabla \cdot \left( M \nabla \frac{\delta \mathscr{F}}{\delta \phi} \right)$$

### Numerically solved via a hybrid Lattice Boltzmann Scheme

• Velocity field v • Concentration field  $\phi$ 



# LATTICE BOLTZMANN MODELS

Based on phase-space discretisation form of the Boltzmann equation

$$\partial_t f + \xi_{\alpha} \partial_{\alpha} f + \frac{F_{\alpha}}{\rho} \partial_{\xi_{\alpha}} f = \Omega(f)$$

**Discretised both in real and velocity space** 

$$f_{i}(\overrightarrow{r} + \overrightarrow{\xi}_{i}\Delta t, t + \Delta t) - f_{i}(\overrightarrow{r}, t) = \mathscr{C}(f_{i}, t)$$
$$\mathscr{C}(f_{i}, t) = -\frac{1}{\tau}(f_{i} - f_{i}^{eq})$$



Mass and momentum density are defined as

$$\sum_{i} f_{i}^{eq} \xi_{i\alpha} = \rho u_{\alpha} \qquad \sum_{i} f_{i}^{eq} = \rho$$
$$\sum_{i} f_{i}^{eq} \xi_{i\alpha} \xi_{i\beta} = \rho u_{\alpha} u_{\beta} - \sigma_{\alpha\beta}$$

- The Equilibrium distribution functions are expanded up to a given order in the fluid velocity
- The Navier Stokes equations can be formally derived from the Lattice Boltzmann equation in the long wave length limit through Chapman-Enskog expansion



**Review Letters,** 128 (2022)

## **ACTIVE TURBULENCE**



Jörn Dunkel, Sebastian Heidenreich, Knut Drescher, Henricus H. Wensink, Markus Bär, and Raymond E. Goldstein

Phys. Rev. Lett. **110**, 228102 – Published 28 May 2013



Cascade or not cascade? Energy transfer and elastic effects in active nematics LN Carenza, L Biferale, G Gonnella Europhysics Letters 132 (4), 44003 (2021)

Multiscale control of active emulsion dynamics

LN Carenza, L Biferale, G Gonnella Physical Review Fluids 5 (1), 011302 (2020)



# **Planned Activity**

#### **Multi-Phase models**



Typical Runs: 128/256 cores for 48 hours 50 exploratory runs 307200 core hours





# Non-equilibrium statistical models for self propelled particles and DNA transcription



• 
$$U(r) = \begin{array}{c} U_{\text{Mie}}(r) - U_{\text{Mie}}(r_{min}) & \text{if } r < r_{min} \\ 0 & \text{if } r \ge r_{min} \end{array}$$

• 
$$\mathbf{F}_{FENE} = -\frac{k(\mathbf{r}_i - \mathbf{r}_j)}{1 - r_{ij}^2/r_0^2}$$

$$\begin{split} m\ddot{\mathbf{r}}_{i} &= -\gamma\dot{\mathbf{r}}_{i} - \frac{\partial U_{FENE}}{\partial r_{i,i+1}} \,\,\hat{\mathbf{r}}_{i,i+1} - \sum_{j\neq i}^{2N} \frac{\partial U}{\partial r_{ij}} \,\,\hat{\mathbf{r}}_{ij} + F_{\mathrm{act}}\hat{\mathbf{n}}_{i} + \sqrt{2D_{0}} \,\,\eta_{i} \,\,, \\ m\ddot{\mathbf{r}}_{i+1} &= -\gamma\dot{\mathbf{r}}_{i+1} + \frac{\partial U_{FENE}}{\partial r_{i,i+1}} \,\,\hat{\mathbf{r}}_{i,i+1} - \sum_{j\neq i+1}^{2N} \frac{\partial U}{\partial r_{i+1,j}} \,\,\hat{\mathbf{r}}_{i+1,j} + F_{\mathrm{act}}\hat{\mathbf{n}}_{i+1} + \sqrt{2D_{0}} \,\,\eta_{i} \,\,, \end{split}$$



 $\mathbf{F}_{act}$ 

Fact

## ACTIVE PARTICLES

 $\sqrt{2D_0} \eta_{i+1}$ 

## **Molecular Dynamics Methods**



It can be used with multiple CPUs and with a GPU



LAMMPS is a classical molecular dynamics code with a focus on materials modeling.

It's an acronym for Large-scale Atomic/Molecular Massively **Parallel Simulator**.

Bead-spring polymer melt with 100-mer chains and FENE bonds: •32,000 atoms for 100 timesteps reduced density 0.8442 (liquid) •force cutoff of 2^(1/6) sigma •neighbor skin = 0.4 sigma •neighbors/atom = 5 (within force cutoff) •NVE time integration

### **Molecular Dynamics Methods**

## Active self-propelled particles

### **Active (with repulsion) Passive (with attraction)**



- Micro and macro phase separation of active systems
- Self-organization and dynamics of active clusters
- Phase diagram



Cugliandolo et al. PRL 119 (26), 268002 2017 Digregorio et al. PRL 121, 098003 2018 Caporusso et al. PRL 125 (17), 178004 2020

# Planned Activity

Three control parameters Attraction





No Attraction, Any value of density and activity

Attraction, low activity, low density: Gel Phase

> Typical Runs: 256 cores for 48 hours 30 exploratory runs 368640 core hours

#### Activity Density

Attraction, higher activity, low density: Arrested Phase Separation

Attraction, high activity, high density: Percolating Network







## **POLYMER MODEL FOR GENE ORGANIZATION AND TRANSCRIPTION**



Neutral



Binding sites (TU)

3Mbp-> 1000 beads (each 30nm in diameter->3Kbp)



Binders (TF) Weakly with all others

TFs SWITCH BETWEEN ACTIVE AND INACTIVE STATES Rapid exchange of factors and polymerases between bound and free states seen in photobleaching experiments

C. A. Brackley et al. Nat. Comm. (2021)



TU in a cluster

TU+TF





MD SIMULATION





Ranke

## **Active Stress Tensor**

**Dissipative/reactive term** 

 $\sigma_{\alpha\beta}^{act} = -\zeta\phi Q_{\alpha\beta} - \bar{\zeta}\epsilon_{\alpha\mu\nu}\partial_{\mu}(\phi Q_{\nu\beta})$ 





**Non-Equilibrium terms** 



## **Active Cholesteric Droplet:** A novel Motility mode



### 150000



## Molecular Dynamics Methods

Typical Runs: 128 cores for 48 hours 45 exploratory runs 276480 core hours

### **USE CASES (UNIBA)**

### Model for lipid bilayers and ion channels

**Coarse-grained** 



