### WP1@Spoke 2

# b) Theoretical research projects in domains already using HPC solutions 5) Physics of complex systems

#### **HPC Methods:**

-Lattice Boltzmann

-Molecular dynamics

-Monte Carlo

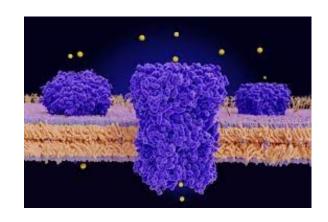
#### **General topics of Use Cases:**

-complex fluid dynamics

-non-equilibrium statistical models

-biological models







#### **UNIBA**:

- -Gonnella Giuseppe
- -Suma Antonio
- -Lamura Antonio
- -Negro Giuseppe



#### **UNINA:**

- -Milano Giuseppe
- -Constantinos Siettos



#### **UNIFE:**

- -Pareschi Lorenzo
- -Dimarco Giacomo
- -Bertaglia Giulia
- -Boscheri Walter



### Lattice Boltzmann Methods for complex fluids

#### Based on phase-space discretized form of the Boltzmann equation

$$\partial_t \, f + \xi_lpha \partial_lpha \, f + rac{F_lpha}{
ho} \partial_{\xi_lpha} \, f = \Omega(\, f \,)$$

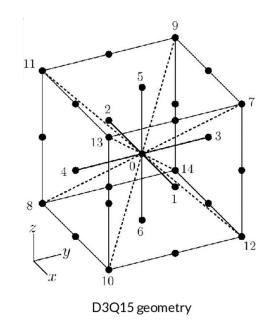
#### Discretizing both tin real and velocity space

$$f_i(ec{r}+ec{\xi}_i\Delta t,t+\Delta t)-f_i(ec{r},t)=\mathcal{C}(\{f_i\},t)$$
  $\mathcal{C}(\{f_i\},t)=-rac{1}{ au}(f_i-f_i^{ ext{eq}})$  BGK approximation

#### Mass and momentum density are defined as

$$\sum_i f_i^{eq} \xi_{ilpha} = 
ho u_lpha \qquad \sum_i f_i^{eq} = 
ho$$

$$\sum_{i}f_{i}^{eq}\xi_{ilpha}\xi_{ieta}=
ho u_{lpha}u_{eta}-\sigma_{lphaeta}$$



- The equilibrium distribution functions are expanded up to a given order in the fluid velocity u.
- The expansion coefficients are determined imposing the above constraints.
- The Navier-Stokes equations can be formally derived from the lattice Boltzmann equation in the long wave length limit through Chapman-Enskog expansion

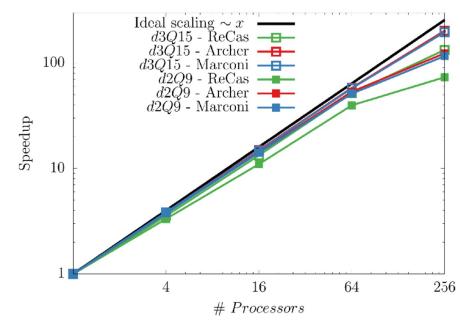
#### **Lattice Boltzmann Method**

Numerical implementation of LBM must face some computational issues

- Memory resources
- Long processing times

"Be wise..." parallelize

- Mpi protocol
- Distributed memory system



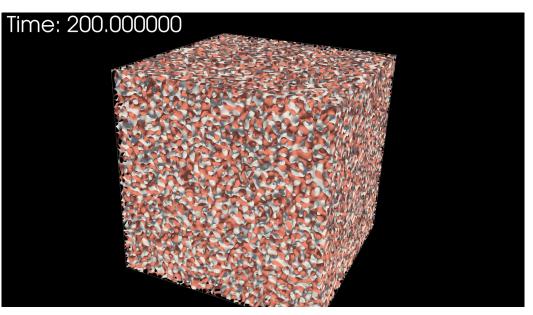
HPC facilities in Bologna, Bari and Edinburgh

Ghost cell method can be implemented to solve the problem of stencils computation on the boundaries between computational subdomains

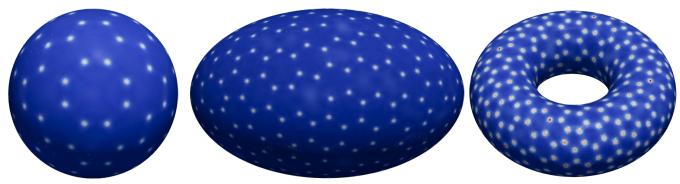
The European Physical Journal E 42 (6), 181

### Complex (liquid crystals, mixtures, etc) fluids

Dynamics of multi-phase fluid mixtures



Topological properties of confined chiral liquid crystals

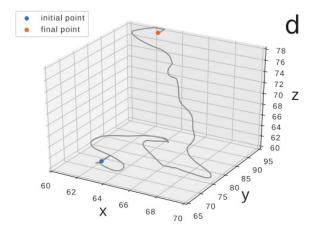


Cholesteric shells: two-dimensional blue fog and finite quasicrystals LN Carenza, G Gonnella, D Marenduzzo, G Negro, E Orlandini Physical Review Letters 128 (2), 027801 (2022)

#### **Active fluids**

#### Self-motility of active droplets





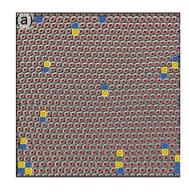
Rotation and propulsion in 3D active chiral droplets LN Carenza, G Gonnella, D Marenduzzo, G Negro

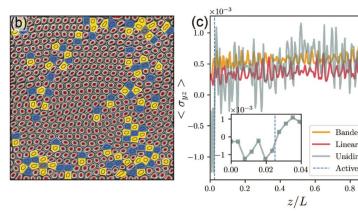
Proceedings of the National Academy of Sciences 116 (44), 22065-22070, (2020)

<u>Chaotic and periodical dynamics of active chiral droplets</u>LN Carenza, G Gonnella, D Marenduzzo, G Negro

Physica A: Statistical Mechanics and its Applications 559, (2020)

Morphology and rheology of active fluids





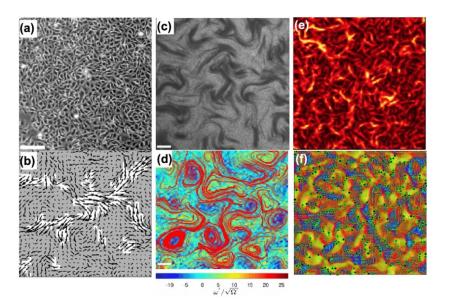
Emergence of negative viscosity states in active polar emulsions

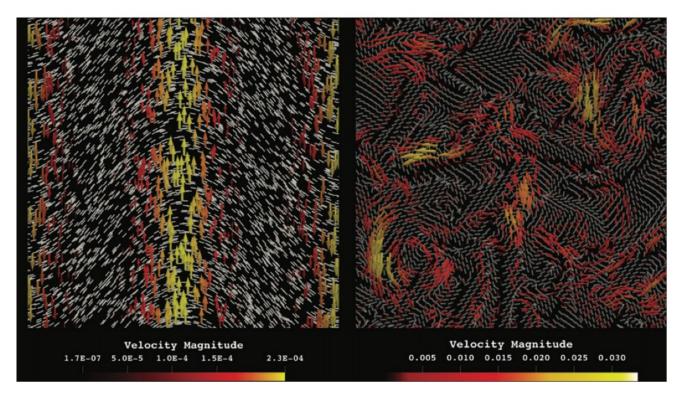
**Lattice Boltzmann Methods** 

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

#### **Active fluids**

#### **Active Turbulence**





No signal of non-linear hydrodynamical transfer and turbulent cascades in active nematic gels

<u>Cascade or not cascade? Energy transfer and elastic effects in active nematics</u>

LN Carenza, L Biferale, G Gonnella Europhysics Letters 132 (4), 44003 (2021)

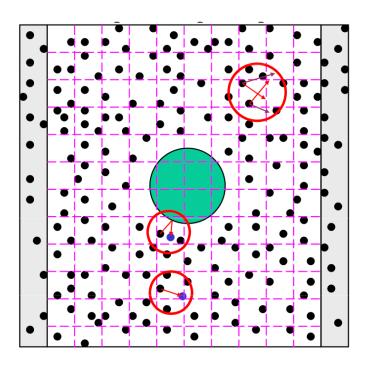
Multiscale control of active emulsion dynamics

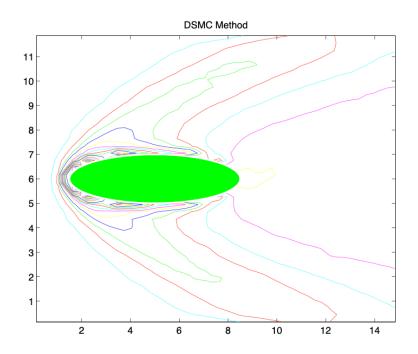
LN Carenza, L Biferale, G Gonnella Physical Review Fluids 5 (1), 011302 (2020)

### General complex systems methods

Introducing an intermediate step between particle systems and fluid models: the so-called kinetic level. These models, characterized by Boltzmann equations, deal with a quantity, the distribution function, which is the density of particles in phase-space (say position and velocity).

The essential idea of Monte Carlo or particle simulations for the Boltzmann equation is to return to the particle description with a number of particles small enough to make the situation computationally treatable but "sufficiently close" to the physical situation.



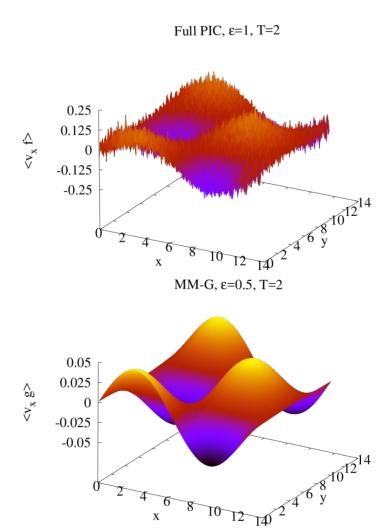


### Fluid dynamics and rarefied flows

Monte Carlo methods for fluid dynamics and rarefied flows. We plan to develop new stochastic based methods for the resolution of flows out of equilibrium. In particular we will address the resolution of the Boltzmann and of the Fokker-Planck equations. The focus is on efficiency and low variance methods by coupling the solution of deterministic equilibrium flows with non equilibrium stochastic solvers.

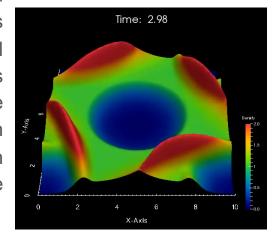
#### **Bibliography**

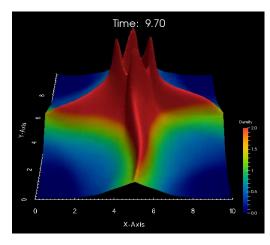
- Asymptotic preserving and time diminishing schemes for rarefied gas dynamic. N. Crouseilles, G. Dimarco and M. Lemou. Kinetic and Related Models, Vol. 10, pp. 643-668, (2017).
- Time diminishing schemes (TDS) for kinetic equations in the diffusive scaling. A Crestetto, G. Dimarco, N. Crouseilles, M. Lemou. Journal of Computational Physics 394, pp. 243-262 (2019).
- A new deviational Asymptotic Preserving Monte Carlo method for the homogeneous Boltzmann equation. A. Crestetto, N. Crouseilles, G. Dimarco, M. Lemou. Communications in Mathematical Sciences 18, pp. 2305-2339 (2020).



#### Emergence behavior in living biological systems

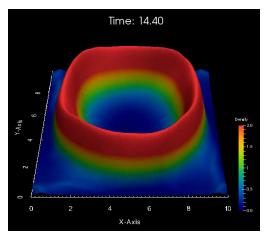
Understanding of the emergence behavior in living biological systems and humanity. The focus here is in the description of complex systems such as organisms, human behaviors and economics. Knowledge, social stratification, opinion dynamics and spread of fake news will considered as prototypes of complex dynamics. The capability of dealing with large amount of data and to infer about these dynamics will be out of the main goal of this WP. We will also consider data-driven approaches in which experimental information measured from real-world systems will be inserted into the model to calibrate the parameters.

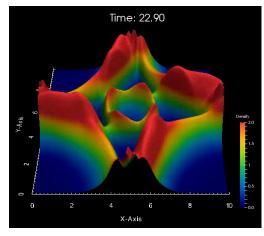




#### **Bibliography**

- Self Alignement driven by jump processes: macroscopic limit and numerical investigation. G. Dimarco and S. Motsch. Mathematical Models and Methods in Applied Sciences, Vol. 26, pp. 1385-1410. (2016).
- Kinetic modeling of alcohol consumption. G. Dimarco, G. Toscani. Journal of Statisical Physics 177, pp. 1022-104 2(2019).
- Social climbing and Amoroso distribution. G. Dimarco, G. Toscani. Mathematical Models and Methods in Applied Sciences 30, pp. 2229-2262 (2020).
- Wealth distribution under the spread of infectious diseases. G. Dimarco, L. Pareschi, G. Toscani, M. Zanella. Physical Review E 102, 022303 (2020).



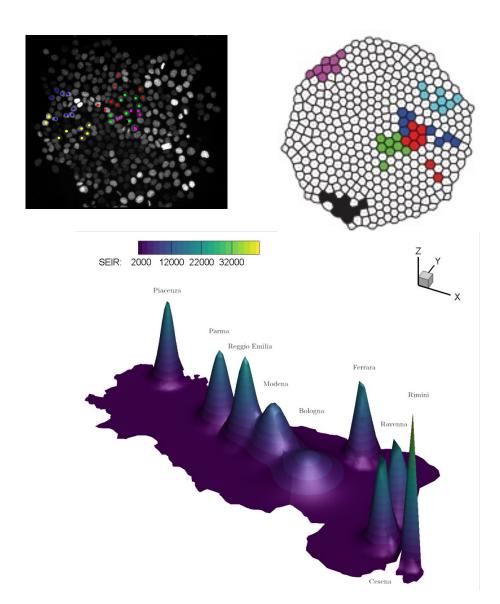


### Medical applications

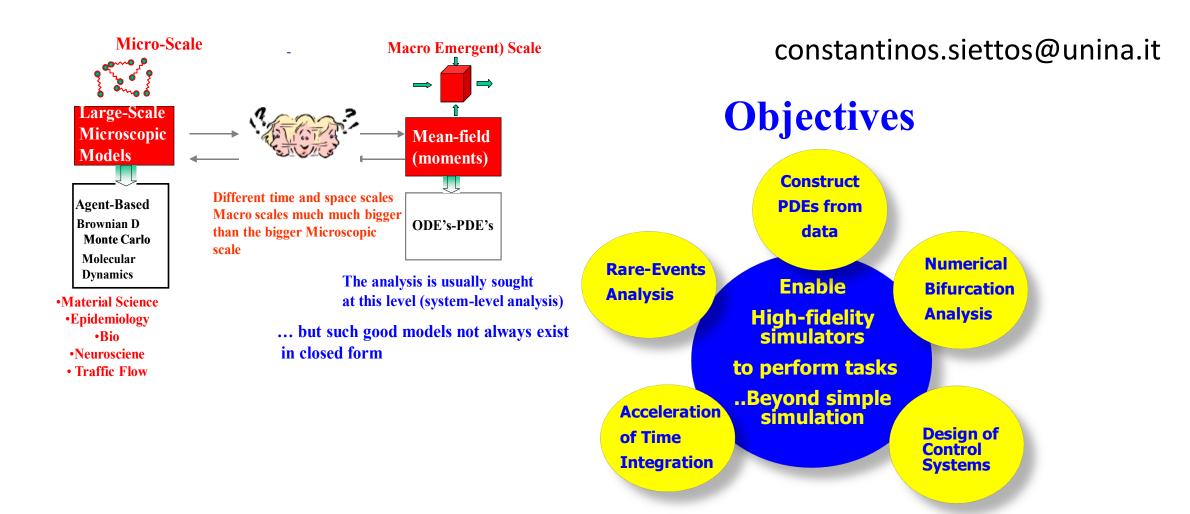
Medical applications such as modeling and simulations of the spread of viruses (Covid-19) through spatially dependent multiscale approaches will be addressed. An important role will be played by the contact dynamics analysis and modeling. The growth, the spread and the control of tumor cells will be studied. For these new models, particle methods and fitting with the data will be also considered. The role of uncertainty in the data will be also discussed and addressed for the construction of reliable models.

#### **Bibliography**

- Are tumor cell lineages solely shaped by mechanical forces? M. Leroy-Leretre, G. Dimarco, M. Cazales, M.L. Boizeau, B. Ducommun, V. Lobjois, P. Degond. Bulletin of Mathematical Biology Vol. 79, pp. 2356-2393, (2017).
- Spatial spread of COVID-19 outbreak in Italy using multiscale kinetic transport equations with uncertainty. G. Bertaglia, W. Boscheri, G. Dimarco, L. Pareschi. Math. Biosci. Eng. 18 (2021) 7028-7059.
- A data-driven epidemic model with social structure for understanding the COVID-19 infection on a heavily affected Italian Province. M. Zanella, C. Bardelli, G. Dimarco, S. Deandrea, P. Perotti, M. Azzi, S. Figini, G. Toscani. Mathematical Models and Methods in Applied Sciences 12, 2533-2570 2021.
- Kinetic models for epidemic dynamics with social heterogeneity. G. Dimarco, B. Perthame, G. Toscani, M. Zanella. Journal of Mathematical Biology 83(1),4 (2021).



#### Bridge the gap between micro and macro scales of Complex Systems



What if physics are known in a more detailed (microscopic) level?

SIMULATION: LIKE EXPERIMENTS

i.e. run atomistic-based models for many units (atoms, agents etc) and for different parameters and for long time to get macroscopic (system-level) information

...but there is no description (in moments of distribution) of the macroscopic behavior;

EQUATION-FREE METHODS

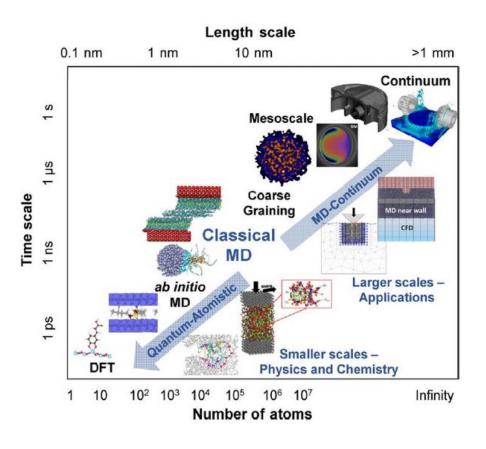
Physics-Informed Machine Learning

**CONTROL** 

**OPTIMIZATION** 

But good macroscopic models do not always exist in closed form ....

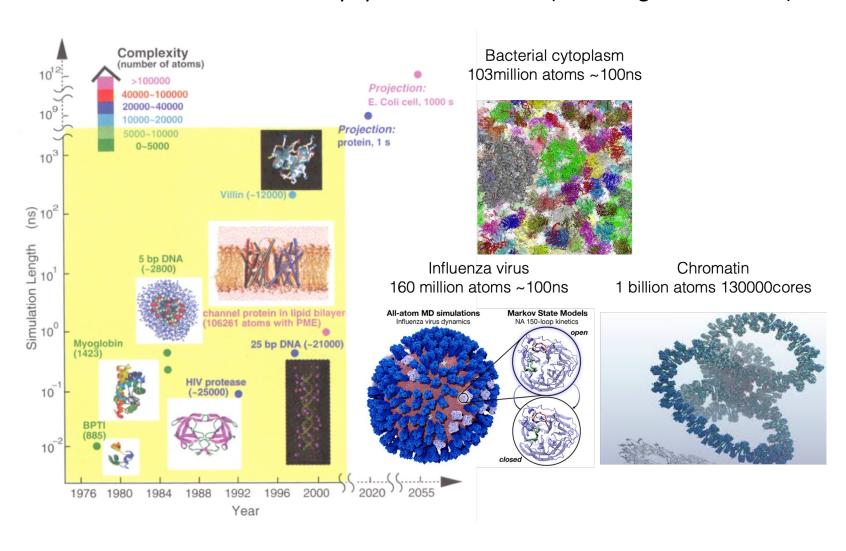
AIM: Systematically bridge the gap between Micro-scopic and Macro-scopic (system-level) World

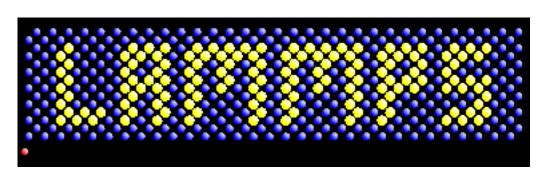


Basic elements are atoms (as a whole, no electrons, protons) or group of atoms in more coarse-grained systems

They are considered as classical objects (no quantum mechanics involved, only Newton's equation of motions.)

General outlook of MD by system's dimension (for biological structures)

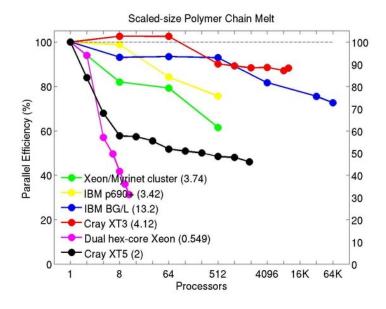




LAMMPS is a classical molecular dynamics code with a focus on materials modeling.

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

It can be used with multiple CPUs and with a GPU



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

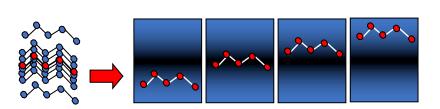
### **OCCAM: Hybrid Particle-Field Parallel MD Simulations**

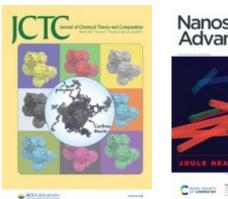
- -Development of coarse-grained models based on Field Theory for soft matter simulations
- -High Performance Large Scale Molecular Dynamics parallel simulations giuseppe.milano@unina.it

Parallel OCCAM for distributed memory (Implementation of MPI interface for CPU, multi GPU in progress)



https://sites.google.com/view/occammd
Complex Molecular Architectures using Field Theory

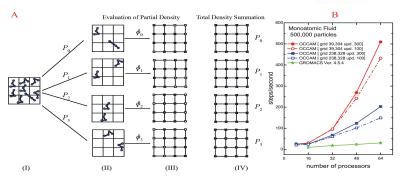


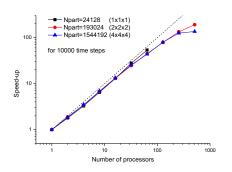






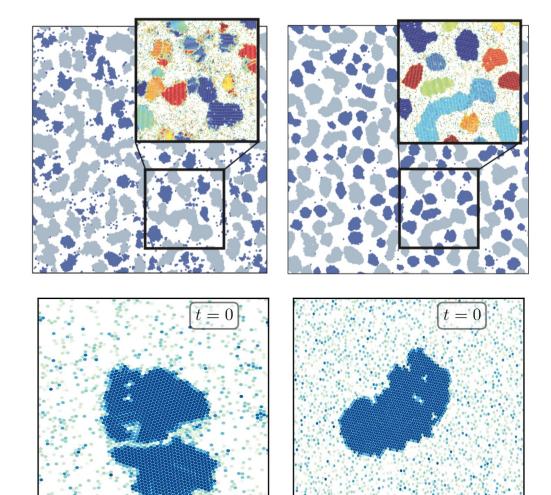




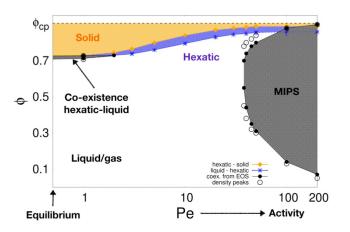


### **Active self-propelled particles**

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



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

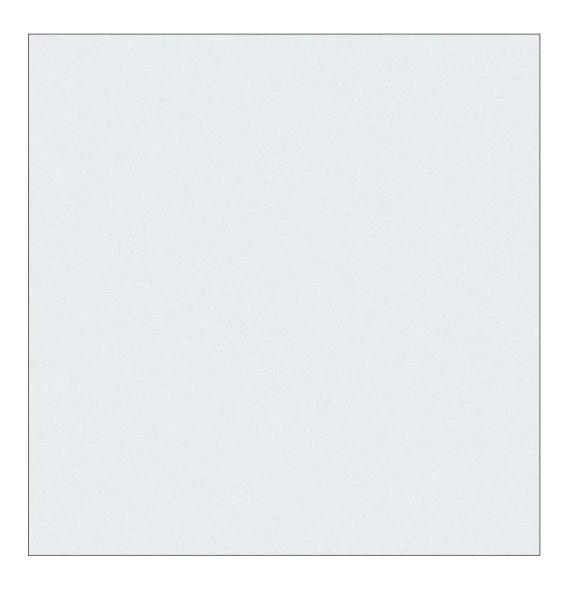


Cugliandolo et al. PRL 119 (26), 268002 2017

Digregorio et al. PRL 121, 098003 2018

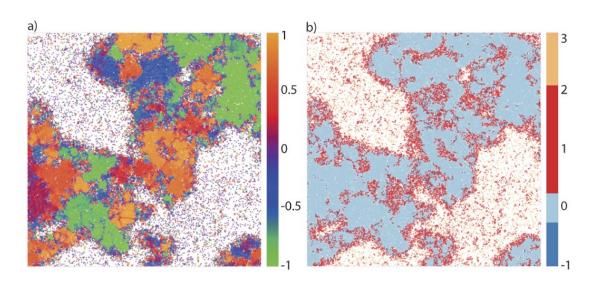
Caporusso et al. PRL 125 (17), 178004 2020

### **Active self-propelled particles**



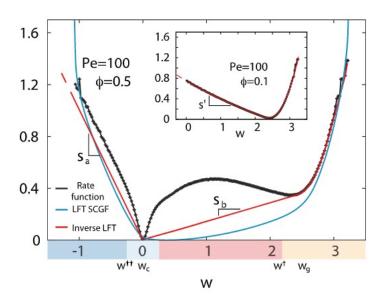
Cluster tracking for 1024<sup>2</sup> particles using clustering algorithms

### Non-equilibrium statistical mechanics

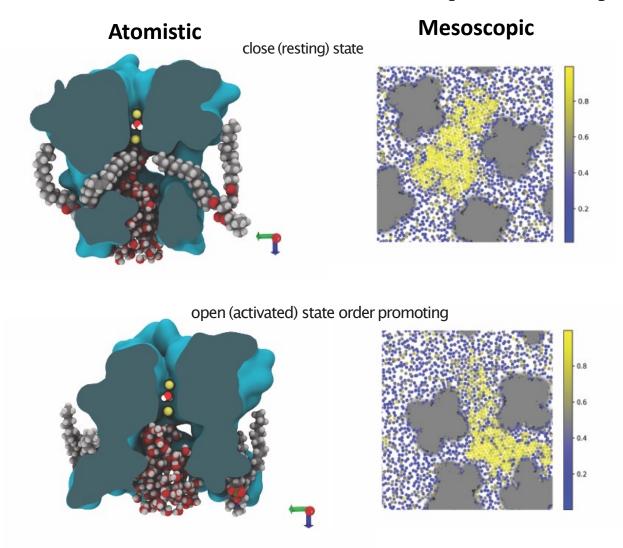


- Analytical and numerical evaluation of the Active Work distributions for active particles
- Study of Fluctuation Relations

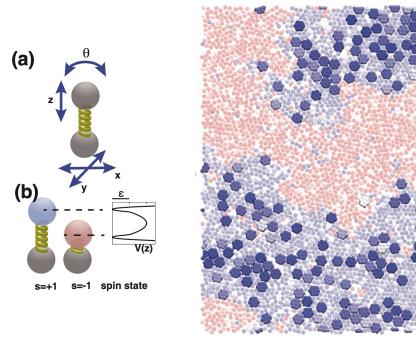
Semeraro et al., J Stat Mech, 12 2021
Cagnetta et al. PRL 119 (15), 158002 2018
Chiarantoni et al. Journal of Physics A 53 (36), 36LT02 2020
Petrelli et al. 102 (1), 012609 2020



### Model for lipid bilayers and ion channels

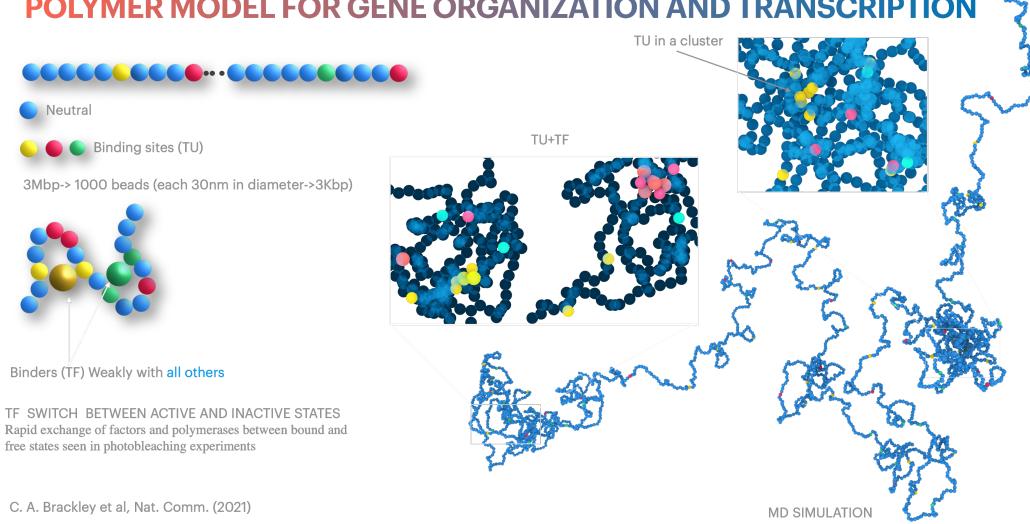


#### **Coarse-grained**

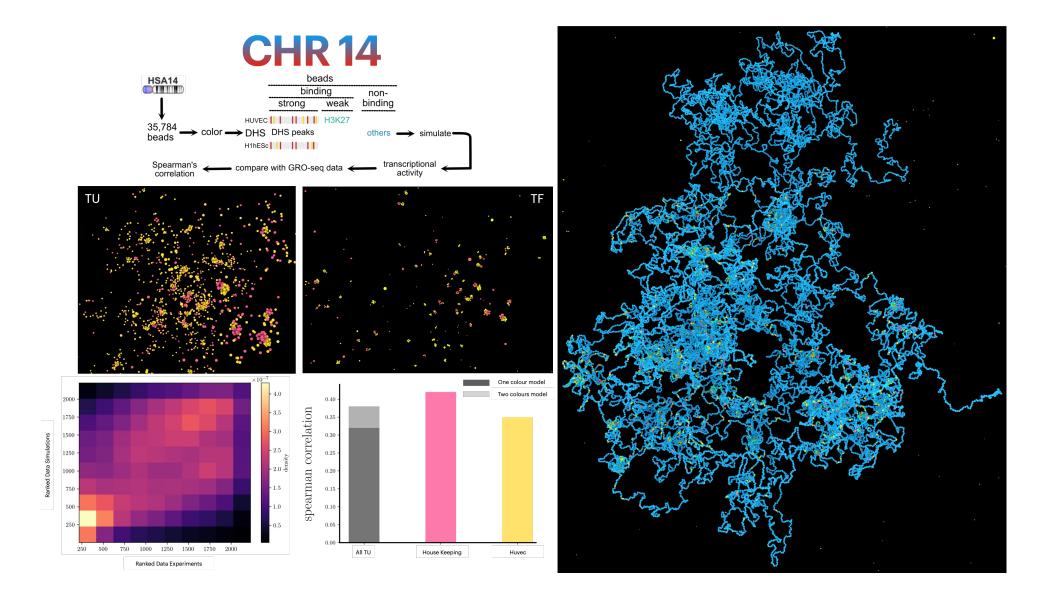


#### **Model for chromatin**

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

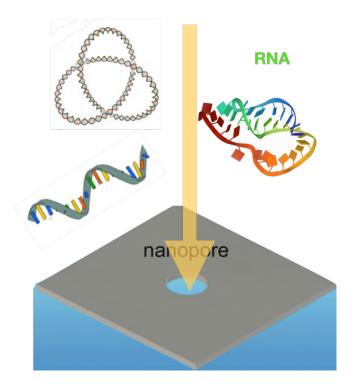


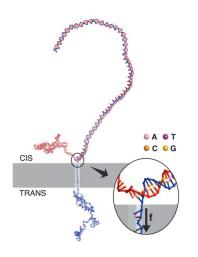
### **Model for chromatin**



### Translocation of biopolymers

### DNA (dsDNA or ssDNA)





Detection of knots and other topological structures Study the impact of different topologies on translocation See how topological structures are exploited by nature

Suma et al. Nature communications 11 (1), 1-9 2020

Suma et al. NAR gkaa080 2020

Suma et al. PNAS 114 (15), E2991-E2997 2017

## Thank You