

FCO. VAZQUEZ

THIS IS THE TITLE OF THE
THESIS

Contents

<i>1</i>	<i>Introduction</i>	<i>5</i>
	<i>1.1 State of art: Hydrogels</i>	<i>5</i>
	<i>1.1.1 Polymeric Structures</i>	<i>5</i>
	<i>1.1.2 Basic Mechanical Properties</i>	<i>5</i>
	<i>1.1.3 What is a hydrogel?</i>	<i>5</i>
	<i>1.2 About computer simulations</i>	<i>5</i>
<i>2</i>	<i>Theoretical framework</i>	<i>7</i>
<i>3</i>	<i>Numerical Experiments</i>	<i>9</i>
	<i>3.1 Simulation tools</i>	<i>9</i>
	<i>3.1.1 LAMMPS</i>	<i>9</i>
	<i>3.2 Simulation methodology</i>	<i>10</i>
	<i>3.2.1 Assembly protocol</i>	<i>11</i>
	<i>3.2.2 Shear protocol</i>	<i>12</i>
	<i>3.3 Results</i>	<i>12</i>
	<i>3.3.1 Network analysis</i>	<i>12</i>
	<i>3.3.2 Mechanical response</i>	<i>12</i>
<i>4</i>	<i>Conclusion</i>	<i>13</i>
	<i>4.1 Discussion</i>	<i>13</i>
	<i>4.2 Future work</i>	<i>13</i>
<i>5</i>	<i>Bibliography</i>	<i>15</i>

1

Introduction

1.1 State of art: Hydrogels

1.1.1 Polymeric Structures

1.1.2 Basic Mechanical Properties

1.1.3 What is a hydrogel?

1.2 About computer simulations

2

Theoretical framework

2.1 Soft Colloids

2.2 Mechanical response

2.3 Hydrogels

2.3.1 Gels

2.3.2 Cross-linking mechanisms

2.3.3 Mechanical properties

2.4 Molecular Dynamics

3

Numerical Experiments

3.1 Simulation tools

Description of what is in this chapter Becuase yes

3.1.1 LAMMPS

Talk about LAMMPS LAMMPS is a computational engine for modeling interacting particles at any length scale, so long as the interactions are primarily short-range, and particles densities are moderately bounded[Thompson et al., 2022]. LAMMPS implemented MD algorithms to enable parallelism across CPUs via MPI, and some versions of these algorithms for GPUs.

For test simulations we used the mpi paralelization scheme, meanwhile the final results where computed with the serial version, since we had access to a cluster.

A LAMMPS input script (text file) is simply a series of lines each beginning with a command name followed by one or more whitespace separated arguments. Programming-like commands are included which define variables, perform conditional tests, execute loops, or invoke shell commands, eg to launch a program external to LAMMPS. The input script is parsed and executed one line at a time which means a single script can be used to run a simulation in stages, altering one or more parameters between the stages, or to run a series of independent simulations where the entire system is reinitialized multiple times.

- Pair styles - Atom styles - Fix styles - Computes styles - units definitions

Make a resume of partitioning, communication, neighbr lists

Since we had access to the lavis cluster we...

Talk about the Fix styles Fix styles implement operations performed during a dynamics timestep or an energy minimization iteration[Thompson et al., 2022]. The `initial_integrate()` and `final_integrate()` methods of fix styles can be

used to implement portions of the velocity-Verlet algorithm for various ensembles at the appropriate points in the timestep.

Units Lennar Jones units. distance = σ . time = reduce LJ τ . mass = ratio to unitless 1.0. temperature = reduced LJ temp. pressure = reduced LJ pressure energy = ϵ . velocity = σ/τ . force = reduce LJ force (σ/τ^2). <https://docs.lammps.org/99/units.html>

3.2 Simulation methodology

Overview of the simulation workflow.

The simulation methodology is based on the work presented in[Gnan et al., 2017] and[Sorichetti et al., 2023], with the objective of create a representative polymer structure of a microgel and characterize the mechanical response under shear deformation. This methodology creates the structure by using a mixture of two types of patchy particles. The patchy particles are spheres of identical size and mass decorated by patches to represent interaction sites. One type represent a *Crosslinker* and is define by 1 central particle with 4 patches placed at the vertices of a circumscribed tetrahedron. The other one represent a *Monomer* define by 1 central particle and 2 patches placed at the poles.

The interaction between the central particles is modeled with a Weeks-Chandler-Andersen repulsive potential,

$$U_{WCA}(r_{i,j}) = \begin{cases} 4\epsilon_{i,j} \left[\left(\frac{\sigma}{r_{i,j}} \right)^{12} - \left(\frac{\sigma}{r_{i,j}} \right)^6 \right] + \epsilon_{i,j}, & r_{i,j} \in [0, 2^{1/6}\sigma], \\ 0, & r_{i,j} > 2^{1/6}\sigma \end{cases}, \quad (3.1)$$

where $r_{i,j}$ is the distance between the center of the central particles, σ is the diameter of the particles and $\epsilon_{i,j}$ is the energy of the interaction. The patch-patch interaction is modeled with an attractive potential,

$$U_{\text{patchy}}(r_{\mu\nu}) = \begin{cases} 2\epsilon_{\mu\nu} \left(\frac{\sigma_p^4}{2r_{\mu\nu}^4} - 1 \right) \exp \left[\frac{\sigma_p}{(r_{\mu\nu} - r_c)} + 2 \right], & r_{\mu\nu} \in [0, r_c], \\ 0, & r_{\mu\nu} > r_c, \end{cases} \quad (3.2)$$

where $r_{\mu\nu}$ is the distance between two patches, σ_p is the diameter of the patches, r_c is the cut distance of interaction set to $1.5\sigma_p$ and $\epsilon_{\mu,\nu}$ is the interaction energy between the patches. Moreover, the interaction between patches is complemented by a three-body repulsive potential, defined in terms of (3.2), that provides an efficient bond-swapping mechanism making possible to easily equilibrate the system at extremely low temperatures, while at the same time, retaining the single-bond-per-patch condition[Sciortino, 2017],

$$U_{\text{swap}}(r_{l,m}, r_{l,n}) = w \sum_{l,m,n} \epsilon_{m,n} U_3(r_{l,m}) U_3(r_{l,n}), \quad r_{l,n} \in [0, r_c], \quad (3.3)$$

where

$$U_3(r) = \begin{cases} 1 & r \in [0, r_{\min}], \\ -U_{\text{patchy}}(r)/\epsilon_{m,n}, & r \in [r_{\min}, r_c] \end{cases}. \quad (3.4)$$

The sum in (3.3) runs over all triples of bonded patches (patch l bonded both with m and n). $r_{l,m}$ and $r_{l,n}$ are the distances between the reference patch and the other two patches. The parameter $\epsilon_{m,n}$ is the energy of repulsion and w is used to tune the swapping ($w = 1$) and non-swapping bonds ($w \gg 1$). The cut off distance r_c is the same as in the potential of interaction between patches, meanwhile the minimum distance r_{\min} is the distance at the minimum of (3.2), *i.e.* $\epsilon_{m,n} \equiv |U_{\text{patchy}}(r_{\min})|$. Finally, the energy of interaction between crosslinker patches (ϵ_{μ^i, μ^j}) are set to 0 to allow only crosslinker-monomer and monomer-monomer bonding (figure 3.2).

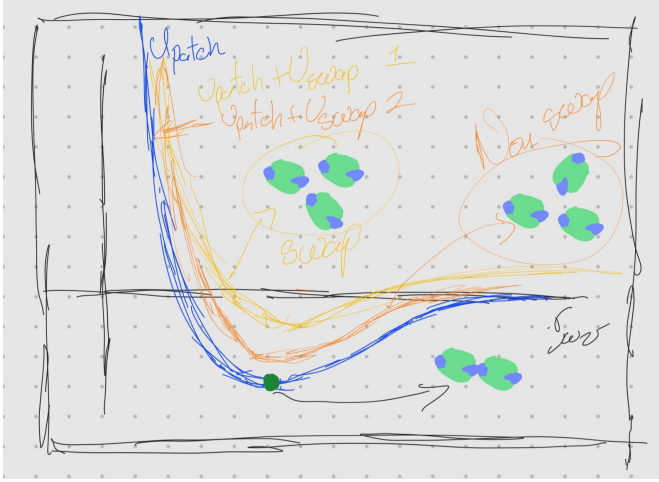


Figure 3.1: La idea de la figura es poner el potencial de interacción entre parches y ver el efecto del potencial de 3 cuerpos cuando $w = 1$ y cuando $w \gg 1$.

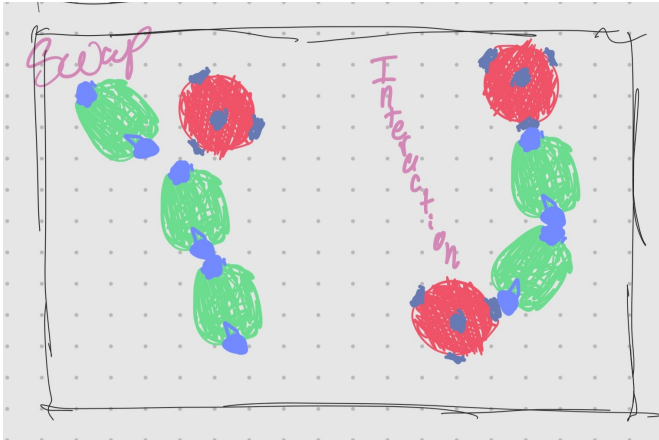


Figure 3.2: La idea de esta es mostrar las posibles configuraciones (monomero-monomero, monomero-crosslinker y un poco de potencial de 3 cuerpos)

3.2.1 Assembly protocol

We perform molecular dynamics (MD) simulations at fixed temperature $T = kBT/\epsilon = 0.05$, where kB is the Boltzmann constant. Thanks to such a low temperature, the system tends to maximize the number of bonds. In addition, owing to the bondswapping mechanism, the system is able to continuously

restructure itself, until the large majority of possible bonds are formed. It is important to said that the main difference between the articles cited and the implementation in this htesis are the absence of FENE bonds and the swelling potential.

3.2.2 *Shear protocol*

Parameters discussion because yes, why shear rate and that stuff and discussion about the damp

Figures about the deformation?

3.3 *Results*

3.3.1 *Network analysis*

After the shear

Before the shear

3.3.2 *Mechanical response*

Varying shear rates and stuff

4

Conclusion

- What we achieve
- Future work

4.1 Discussion

About the mechanical response. If it works or not and so on.

About the network characterization. What it work or order arameters and so on.

4.2 Future work

About different crosslinking mechanisms. To use FENE in some crosslinkers and yes.

About parameters Shear rate, temperature, damp

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