

Kinetic Monte Carlo Simulations for Polymerization in 2 dimensions

Group Lunch - 2024-09-18 - Simon Briesenick

Motivation

- Formation of 2D ordered networks is highly dependent on Monomer + Substrate interactions

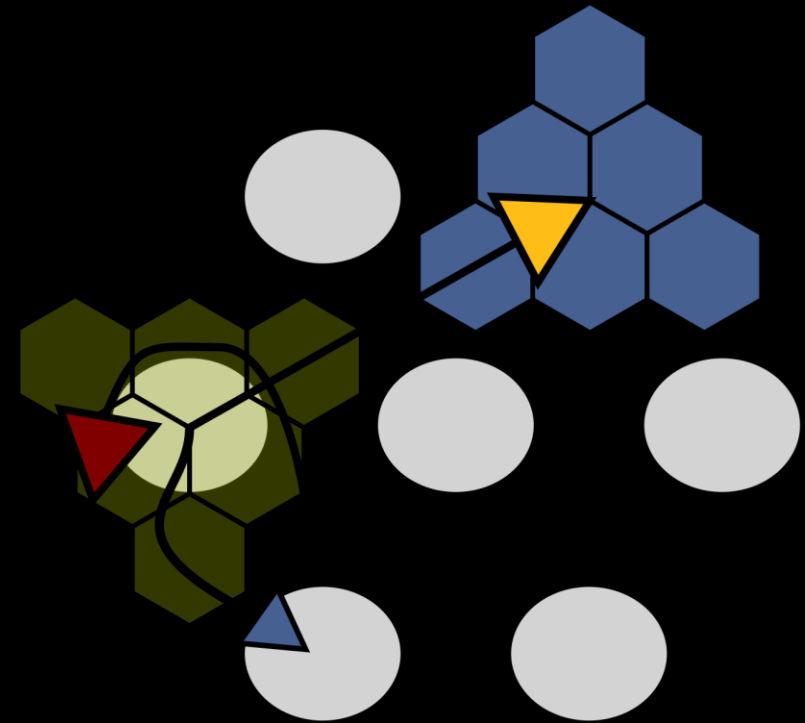
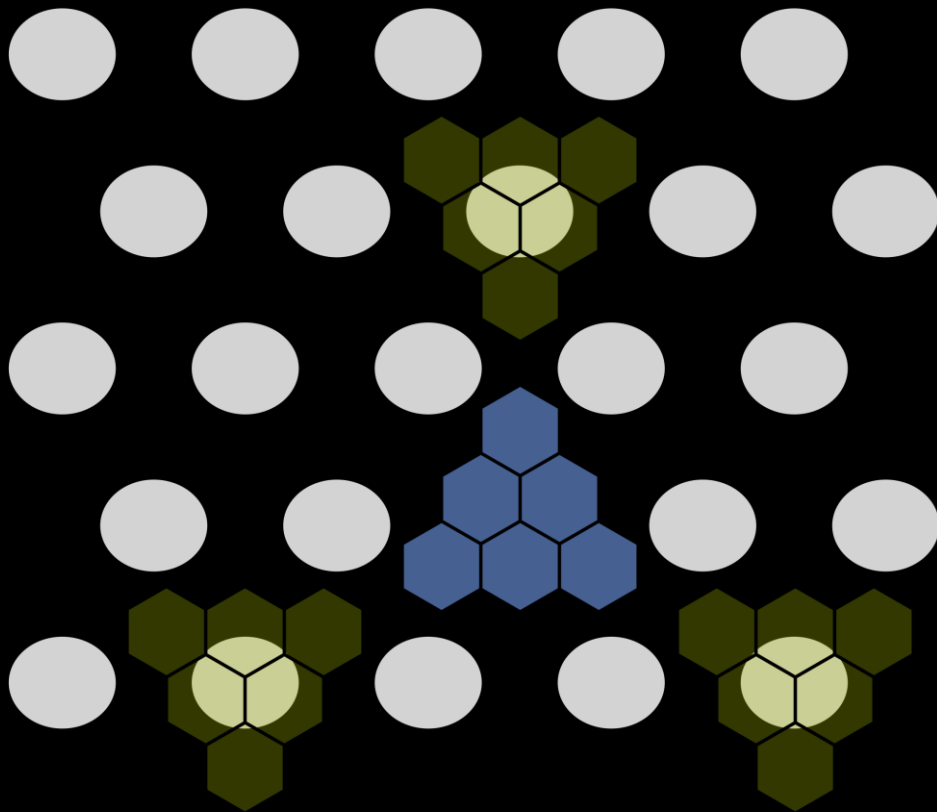
- Diffusion Barrier
- Coupling Barrier
- Rotational Barrier
- Epitaxial Relationship

} Can be tweaked by
material used, surface
orientation, temperature,
adatoms,

➔ Can we use this model to better understand the reaction kinetics on different surfaces, and could we perhaps even extract the Monomer + Substrate interactions

Physical Model

Monomers appear on a hexagonal grid, and can **diffuse**, **rotate**, or **couple** to other monomers (if possible)



Implementation in Python

- Object-oriented approach with **Monomer** and **Lattice** classes
- Initialize lattice with single (fixed) monomer in the center.



- Add one monomer at a time on random position on the lattice

Each timestep, the (free) monomer can diffuse, rotate, and couple. The probability for each of these actions is given by an Arrhenius equation:

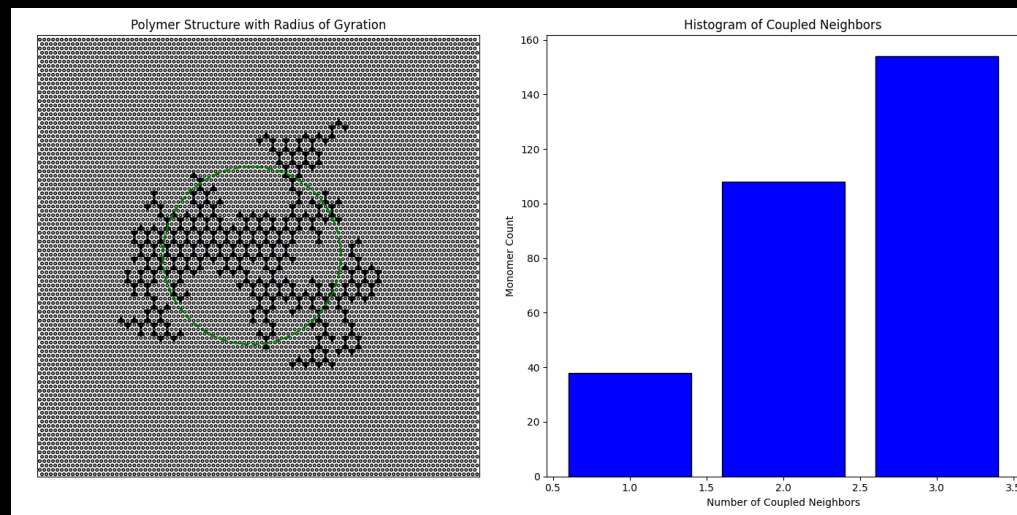
$$p^{d,r,c} = P_0^{d,r,c} \exp\left(-\frac{E^{d,r,c}}{k_B T}\right)$$

Run until new monomer has coupled to growing oligomer, and repeat.

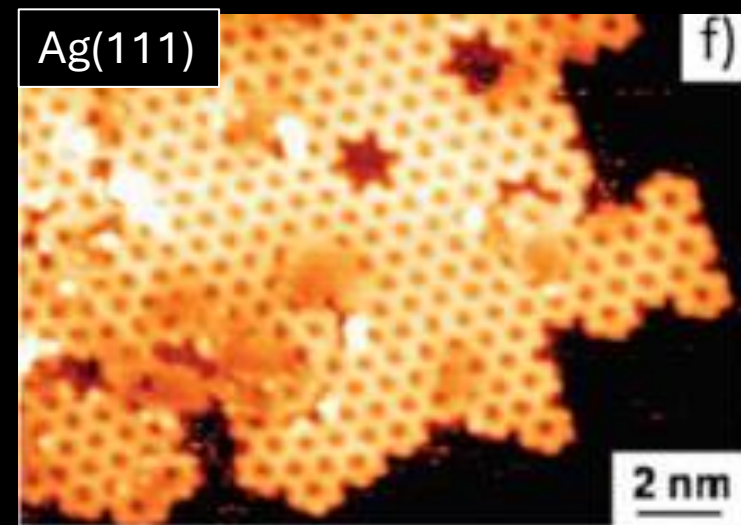
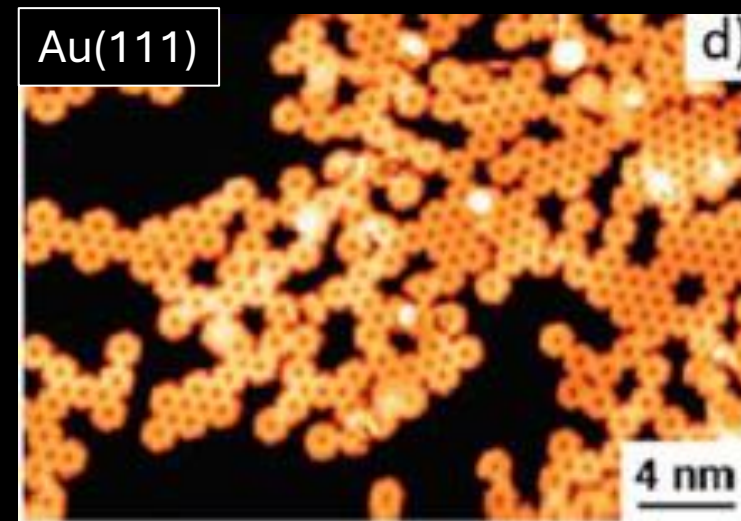
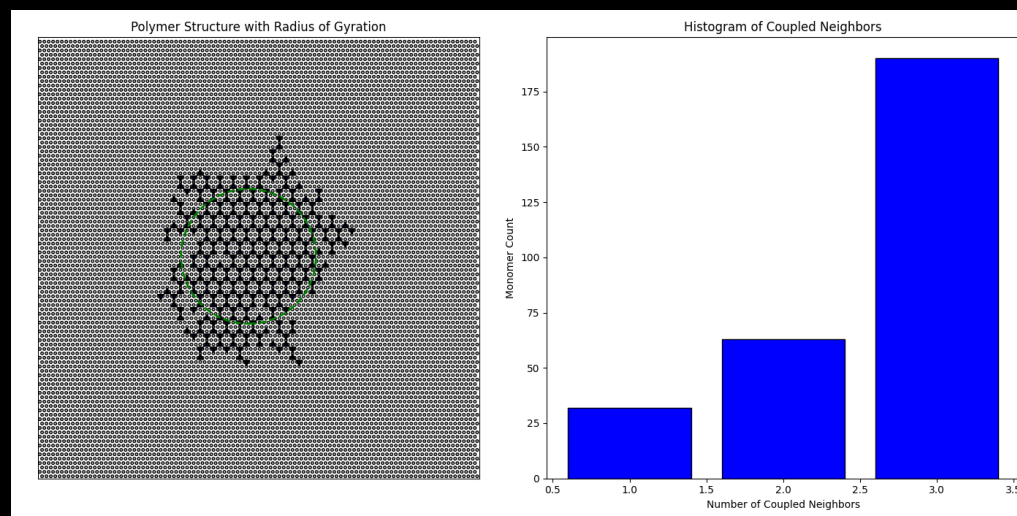
A coupled monomer's probability for all actions is set to 0, i.e., they become immobile

Results

High diffusion
barrier, low coupling
barrier



High coupling
barrier, low diffusion
barrier



Bieri, et al. *JACS* **2010** 132 (46)

Next Steps

- Introduce surface defects, like step edges, point defects, etc.
- Model **copolymerization**, where two different monomer species can polymerize together as **block-**, or as **random-copolymers**
- Scale up the model to analyze how islands merge

