Simulations of homopolymer collapse using forward-flux sampling.

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Third placed in recent QS World University Rankings 'Top 50 Under 50'





Atomic and molecular simulation

• An attempt to visualise the field.

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Motivation - biomineralisation



Columns of calcite



Overview

- Forward-flux sampling on a toy system
 - Freezing of a hard-sphere polymer chain.
 - Kinetics vs thermodynamics.
 - Breakdown of the two state assumption.
 - Improved reaction coordinate?





Attractive hard sphere chain

• Simulate with both Metropolis MC and collision dynamics (event driven molecular dynamics).

Monte-Carlo move set

- Crank shaft
- Pivot
- End-bridging
- Regrowth (Rosenbluth / CBMC)

connectivity altering

Collision dynamics

- Ballistic between events
- Hard core collision
- Well capture and escape events
- Thermal 'jolt' events

connectivity constrained







Properties

 Thermodynamics of system extensively studied previously <u>interviously</u>

[Taylor, Paul and Binder. *Phys. Rev. E.* **79**, 050801 (2009) *J. Chem. Phys.* **131**, 114907 (2009)]

- Single stage "protein-like" collapse for $\chi \lesssim 1.06$
- We study globule-crystal transition for larger χ .
- Brute force sampling feasible for $\chi \ge 1.15$.
- Use Forward Flux Sampling (FFS) for smaller χ .
- FFS simulations use collision dynamics.
- Wang-Landau simulations use MC move set to locate thermodynamic transition temperature.

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Brute force CD trajectory at transition temperature.



Sampling and FFS

- All globule states can be connected by short CD trajectories.
- Different realisations of the crystal state separated by high barriers.
- Breakdown of two-state assumption.



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An improved reaction coordinate?



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Summary

- Studied a very simple model, amenable to extensive MC/MD sampling.
- Brute force rates can be computed in some parameter ranges.
- Computed both forward and backward rates with FFS;
 - as a function of temperature
 - as a function of model parameters
- Discrepancy between 'kinetic' and Wang-Landau transition temperature
 - small c.f. typical error bar in bulk FFS crystallisation studies, but annoying nonetheless.
 - breakdown of two-state assumption due to large barriers between crystalline states?
 - effect of thermalisation or non-statistical dynamics?
 - poor choice of reaction coordinate?
- Proposed alternative reaction coordinate for future study.

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Thank You!

Ruzicka, S.; DQ & Allen, M.P. Folding kinetics of a polymer. *Phys. Chem. Chem. Phys.*, **2012**, 14, 6044-6053.



29/01/2007 Lunchtime Seminar