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Going up in Time and Length Scales in Modeling Polymers

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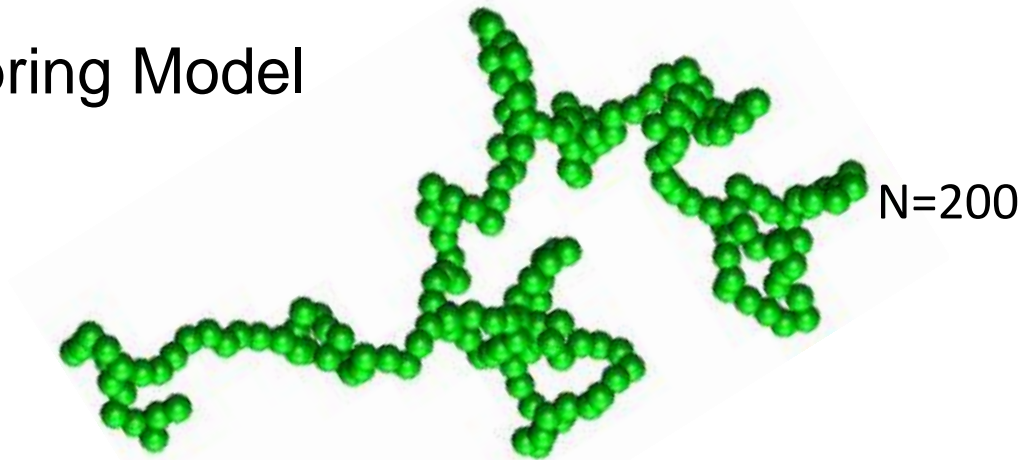
Les Houches, May 16, 2017

Computational Challenges in Polymers

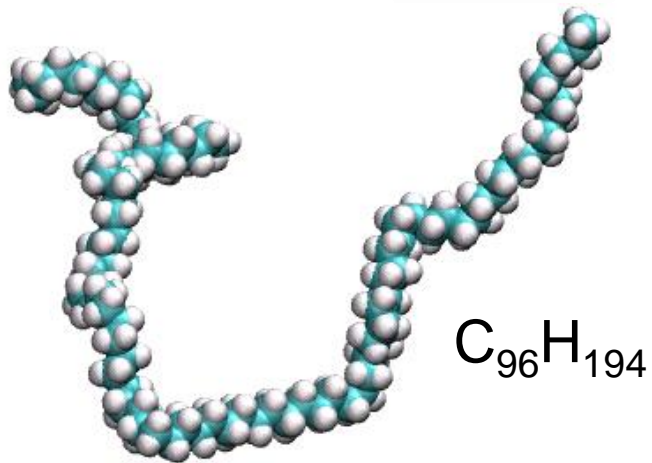
- Longest relaxation time $\tau \sim N^3$
- Chains are Gaussian coils – $R \sim N^{1/2}$
 - Number of chains must increase as $R^3 \sim N^{3/2}$ so polymer chains do not see themselves through periodic boundary conditions
- Double chain length – cpu required increases by at least a factor of $2^{4.5} \sim 23$
 - 1-2 month simulation becomes 2-4 years
- Number of processors limited: ~ 500 -1000 particles/processor

Polymer Simulation Models

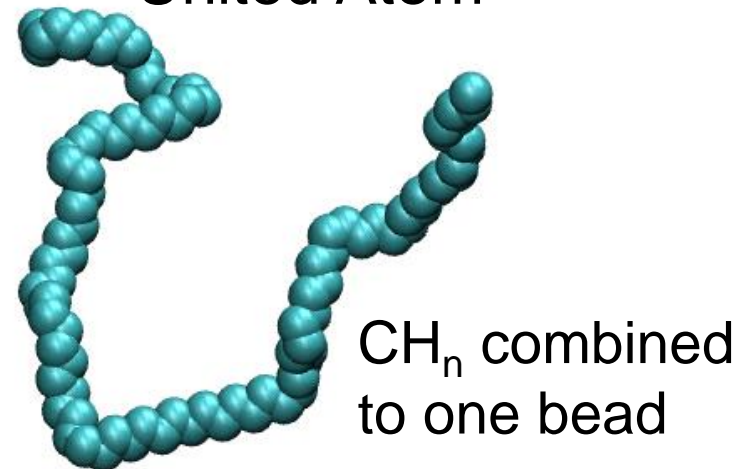
- Bead-Spring Model



- Atomistic: All Atom

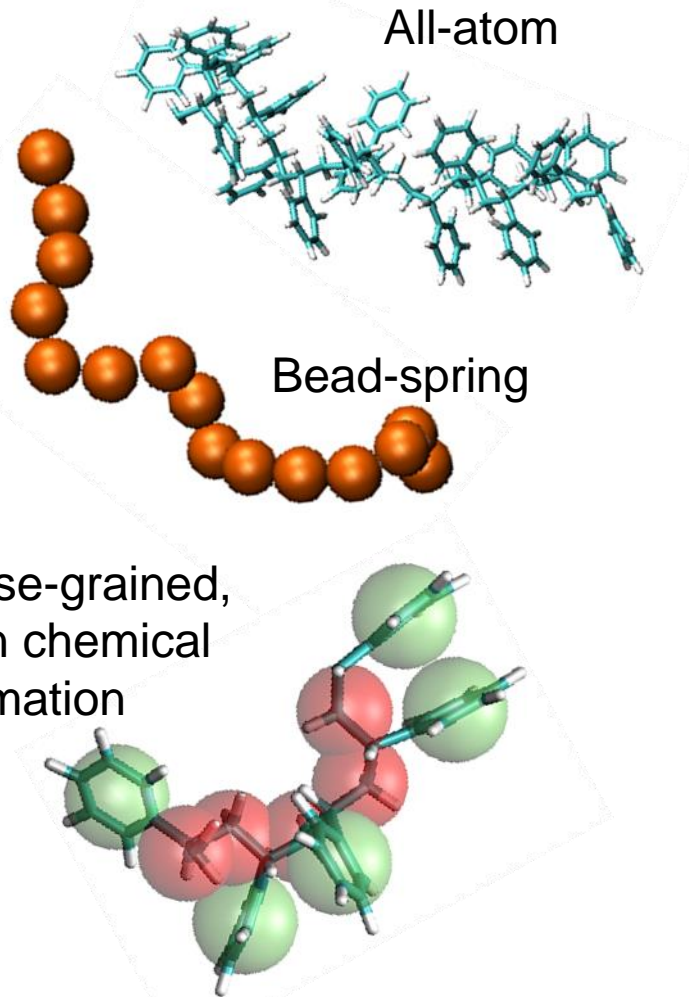


United Atom



Coarse-Graining of Polymers

- To reach larger length/time scales, new coarse graining methods are an active area of research



- Reduced number of degrees of freedom, simpler interaction potentials, reducing the overall computational effort
- Larger time steps (10-20x)
- Reduced effective bead friction due to lower energy barriers and/or a smoother energy landscape
- Back-mapping to fully atomistic model

Bead-Spring Model

- Short range - excluded volume interaction

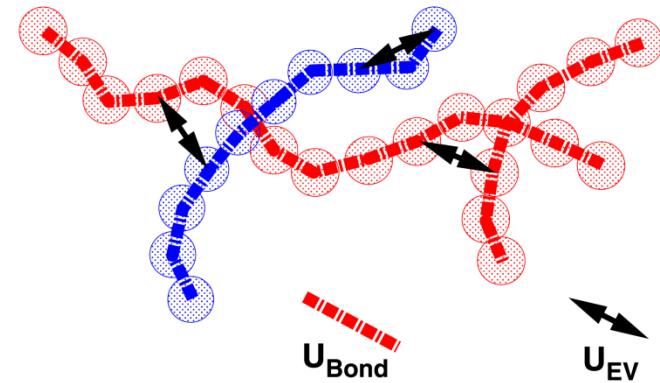
$$U_{\text{LJ}}(r) = \begin{cases} 4\epsilon \left\{ \left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^6 + \frac{1}{4} \right\} & r \leq r_c \\ 0 & r \geq r_c \end{cases}$$

- Bonded interaction - FENE spring

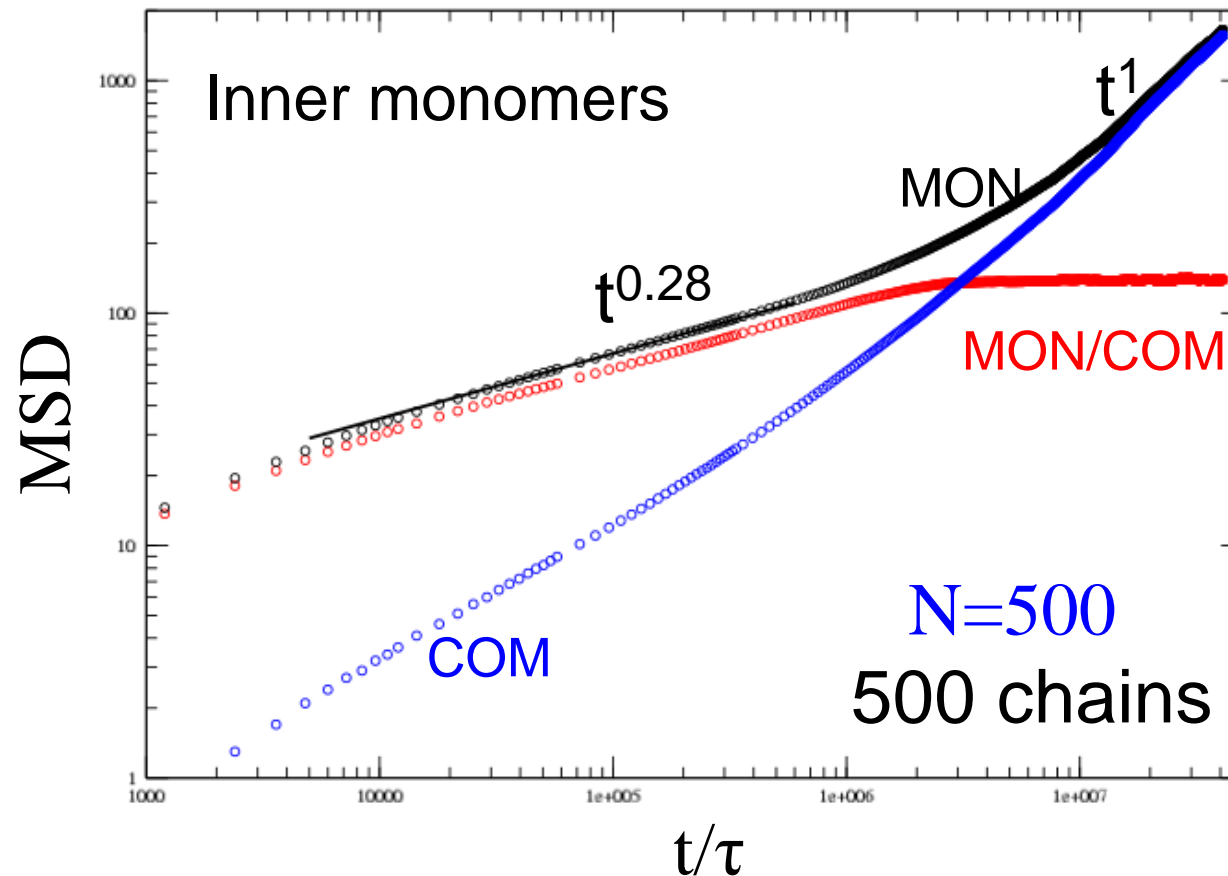
$$U_{\text{FENE}}(r) = \begin{cases} -0.5kR_0^2 \ln(1 - (r/R_0)^2) & r \leq R_0 \\ \infty & r > R_0 \end{cases} \quad k=30\epsilon/\sigma^2, R_0=1.5\sigma$$

- Energy barrier prohibits chains from cutting through each other topology conserved

$$m_i \frac{d^2 \vec{r}_i}{dt^2} = -\vec{\nabla} \cdot U_i - m_i \Gamma \frac{d\vec{r}_i}{dt} + \vec{W}_i(t) \quad \text{Time step } \Delta t \sim 0.01\tau, \tau = \sigma(m/\epsilon)^{1/2}$$



State of the Art: Motion of Entangled Polymer

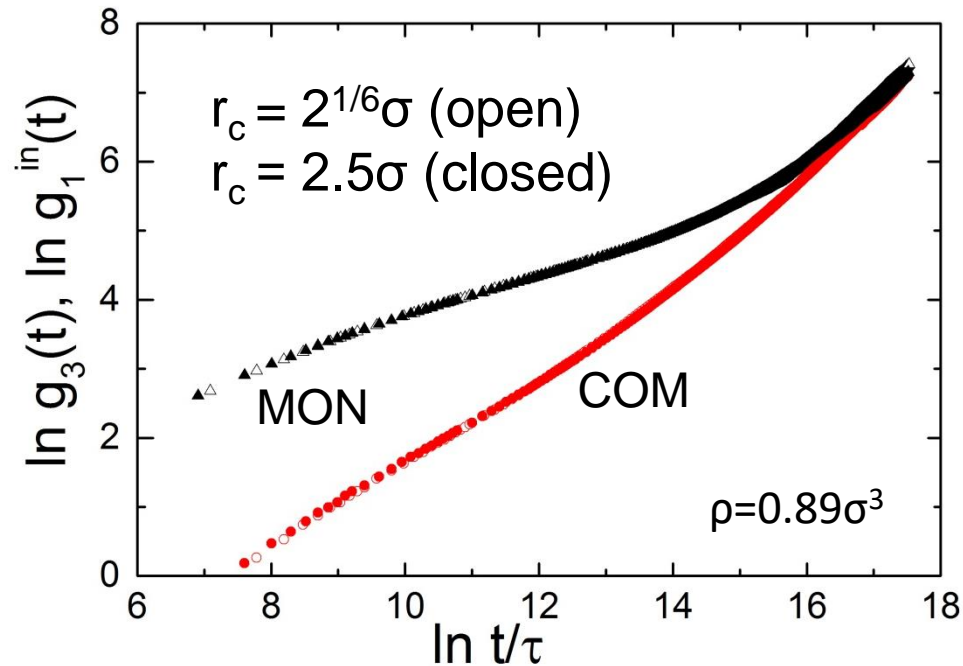
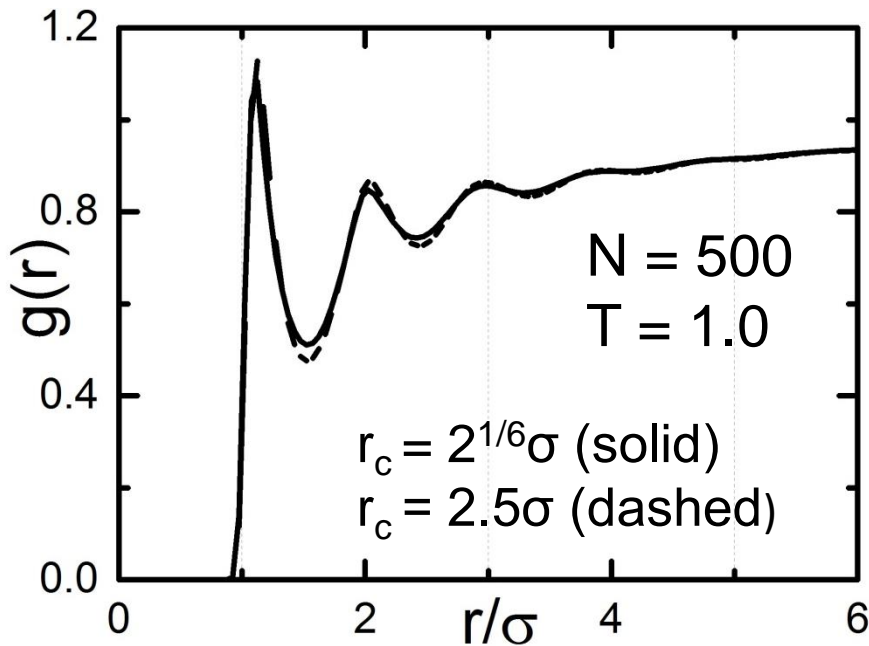


*2 million
core hours*

- $t^{1/4}$ motion is clearly seen for inner monomers
- Theoretically predicated second $t^{1/2}$ region still unresolved

Effect of Attractive Interactions

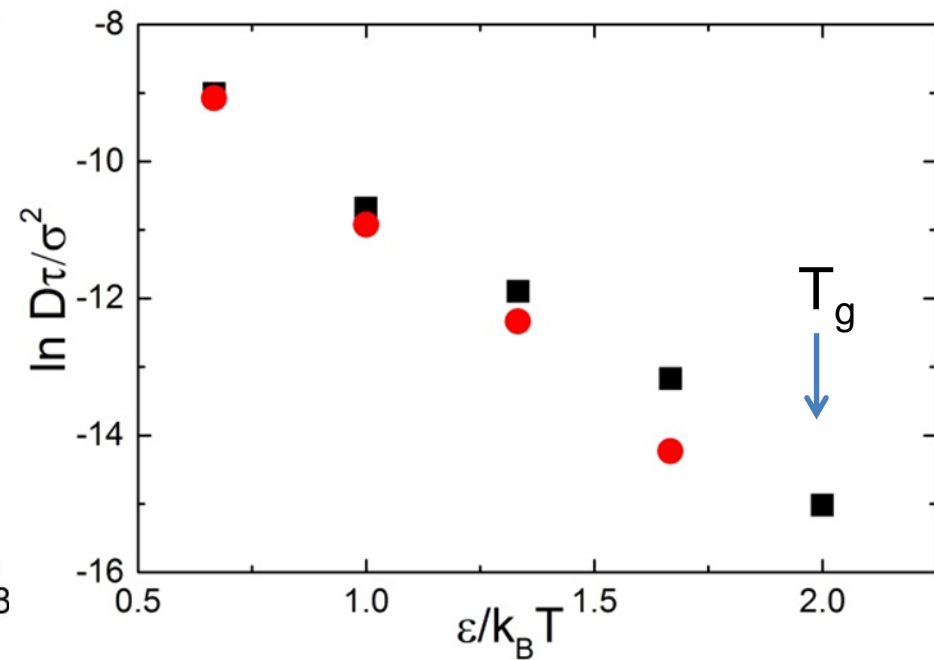
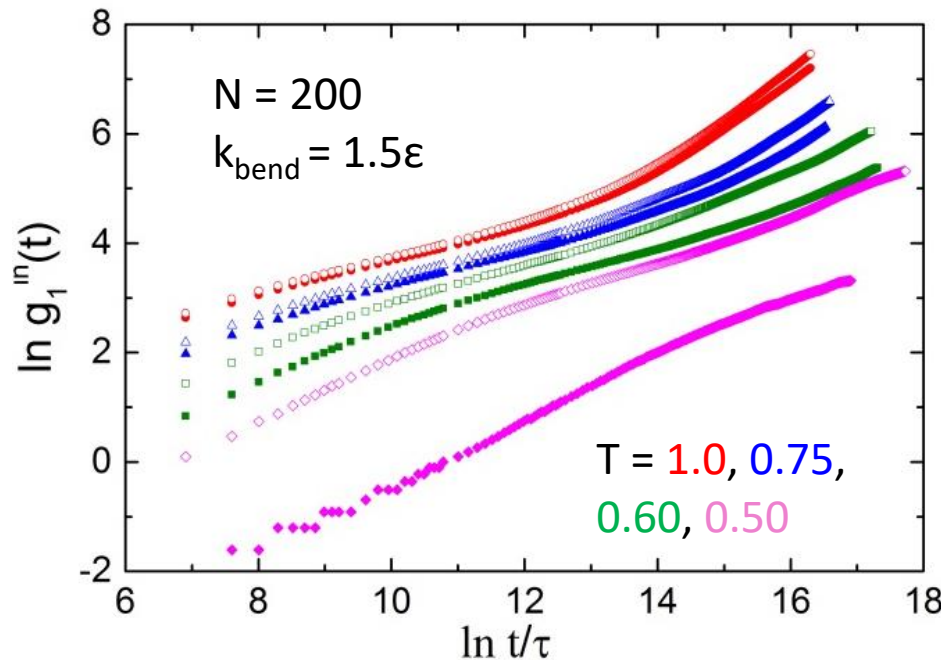
- Purely repulsive interactions widely used for modeling melts
 - computationally the most efficient



- Attractive interactions have essentially no effect on melt properties
- Justify implicit assumption that the dynamics of entangled polymers melts can be studied with short range, repulsive interactions

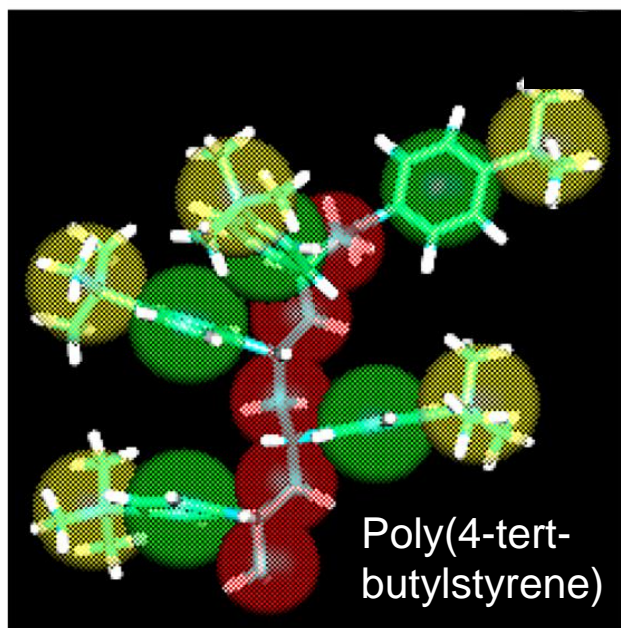
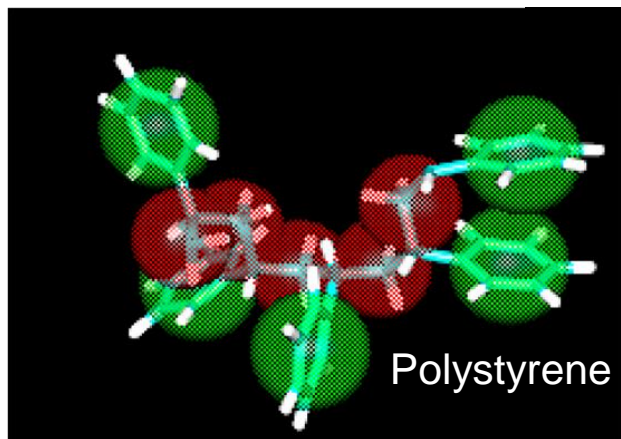
Attractive Interactions: Temperature Effects

- Cool melt slowly at constant pressure
- Compare repulsive and attractive interactions
- Attractive interactions have little effect on the local packing, single chain statics and entanglement length N_e for all T

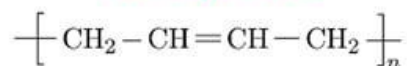


- Attractive interactions reduce chain mobility for $T < 2T_g$

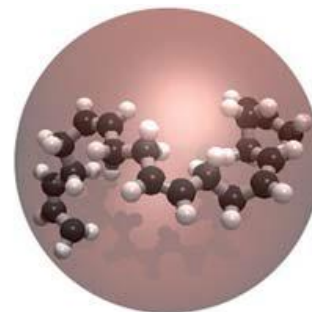
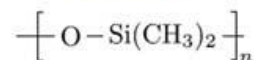
Coarse Graining Polymers



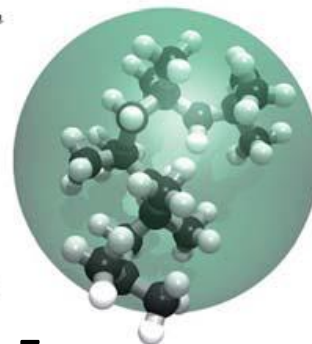
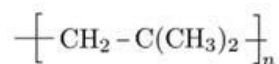
cis-1,4-PB



PDMS



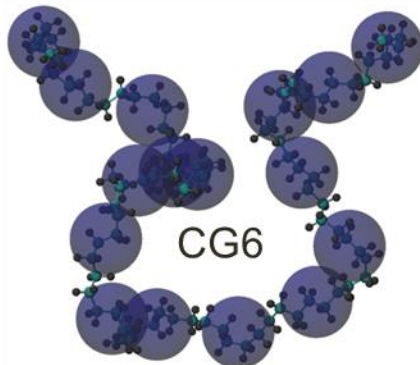
PIB



$n=5$

G. Mauerl et al, RSC Adv., 5, 14065 (2015)

Degree of Coarse Graining Polyethylene



$C_{96}H_{194}$ chain with increasing
degree of coarse graining

- Largest lengths scales of polymer dynamics are controlled by entanglements
- Shortest time and length scales required to resolve dynamic properties not obvious

- **Probe the degree of coarse graining (CGing) required to simultaneously retain significant atomistic detail and access large length and time scales**

Coarse-Graining Methodology

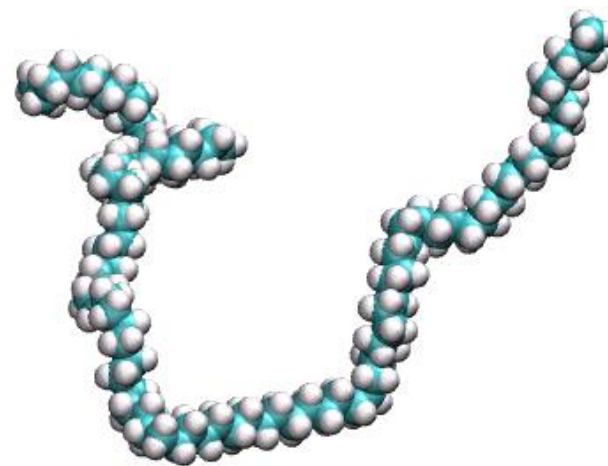
All-atom Melt MD
simulation – LAMMPS*

Define Beads

Bonded Interactions by
Boltzmann Inversion

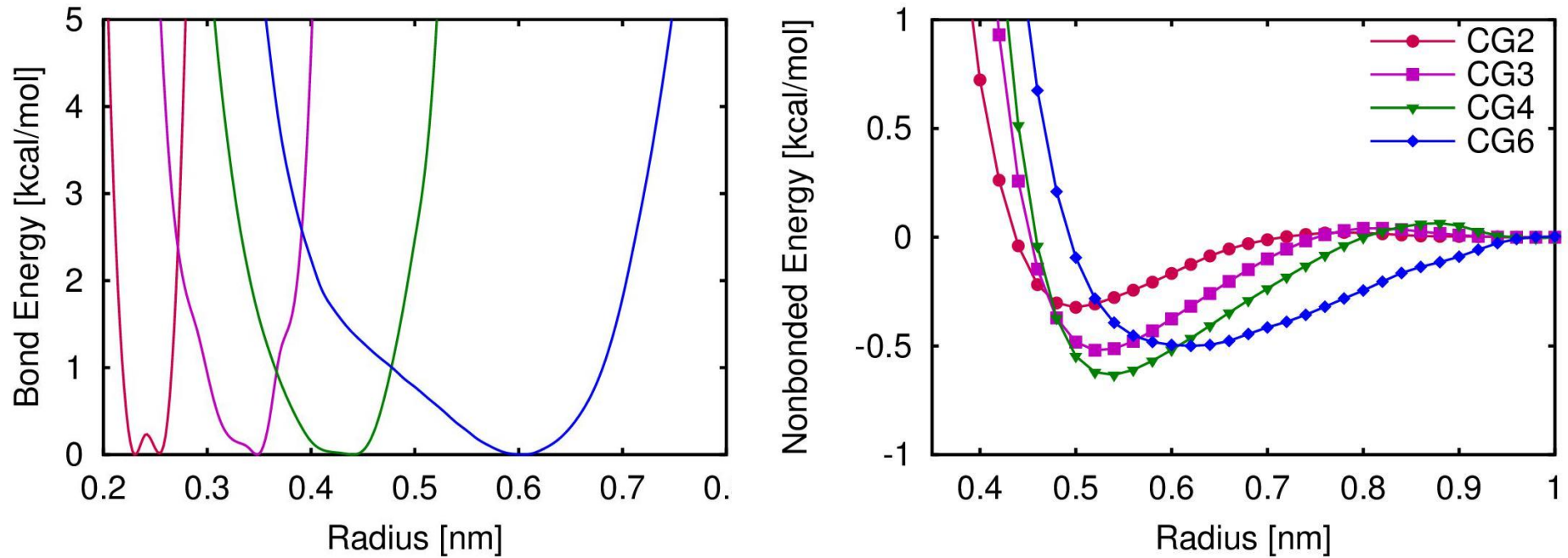
Nonbonded Interactions by
**Iterative Boltzmann
Inversion (IBI)**

Validation



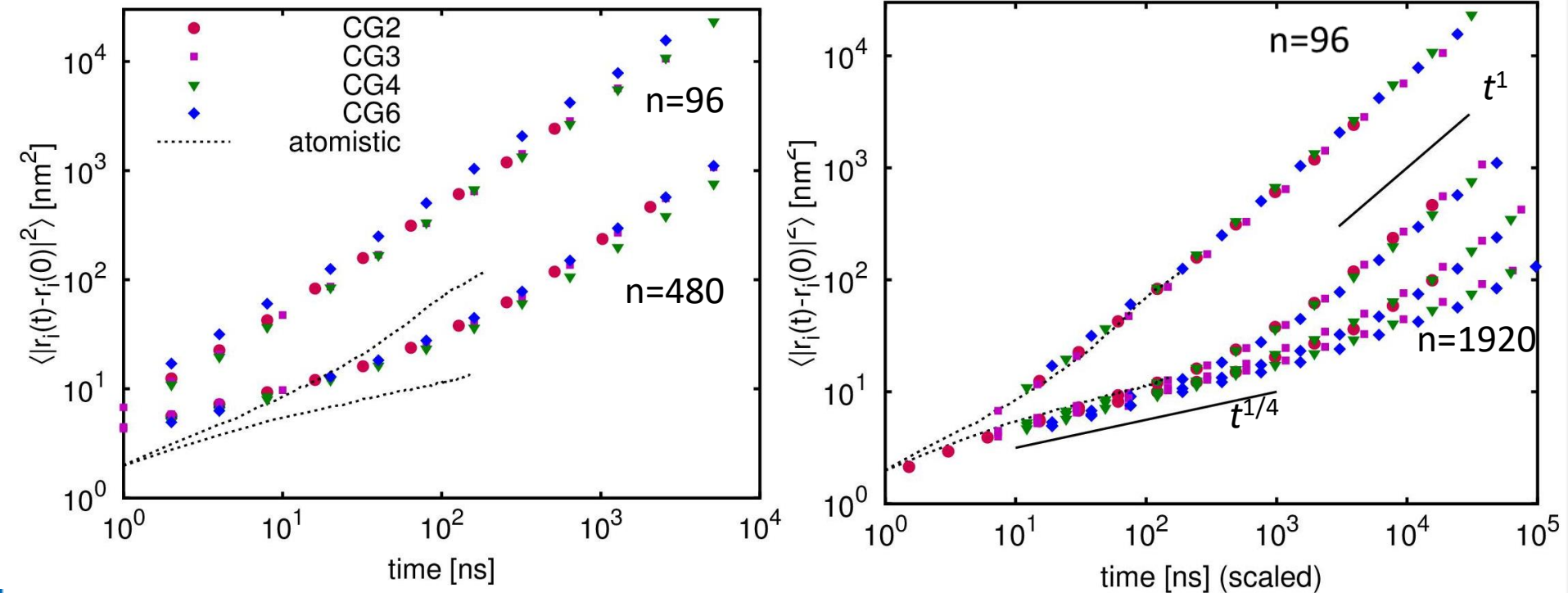
*<http://lammps.sandia.gov/>

Coarse-Grained Potentials



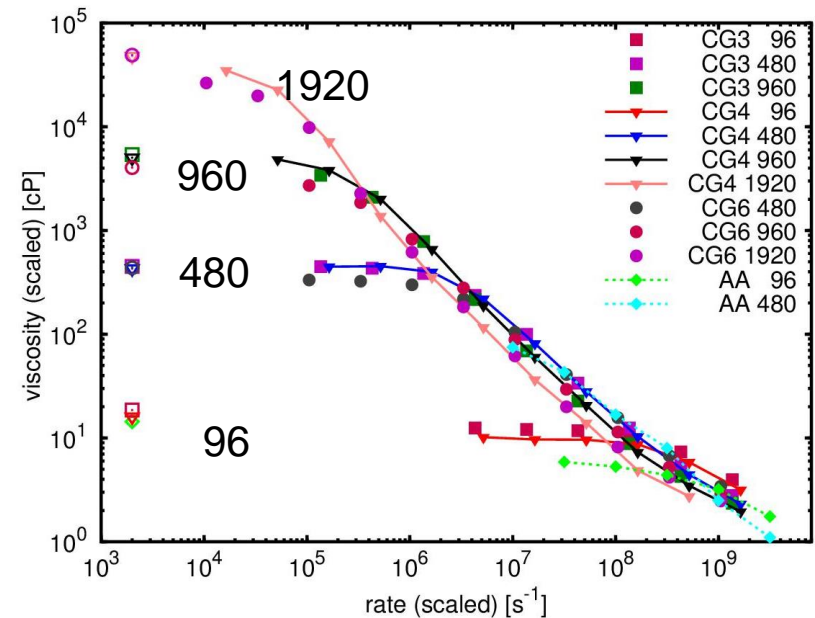
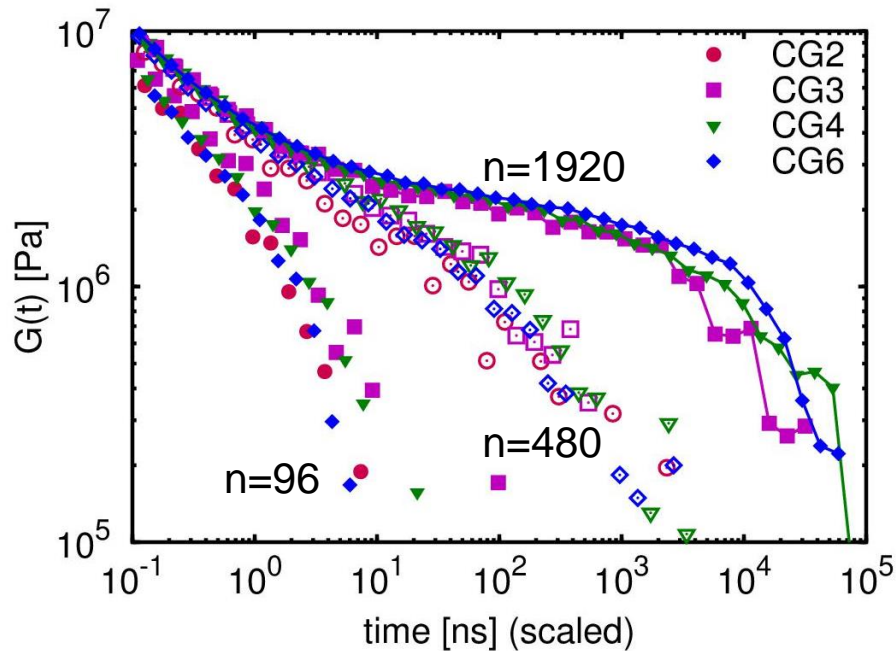
- Average bond length increases, bond distribution broadens as level of CGing increases
- Non-bonded potential softens as level of CGing increases
- Time step increases from 1 fs to 10 -20 fs for $\lambda \geq 3$
- ,Chains cut through each other for $\lambda = 6$
 - extra non-crossing constraint

Chain Mobility



- Coarse graining reduces the number of degrees of freedom, creating a smoother free-energy landscape
- Dynamics of Coarse Grained models 6-10 times faster
- Consistent scaling factor for $n = 96 - 1920$

Stress Relaxation and Viscosity



- Longer, more entangled chains form progressively more distinct plateau region
 - Plateau modulus in good agreement with experiment
- Viscosity versus shear rate show shear thinning at high shear rates, crossing over to shear independent regime
- Time and length scales not accessible by atomistic models

Summary/Outlook

- Atomistic Simulations ideally suited for phenomena on local scale
 - Present limitations 100's nanoseconds, 10's nanometers
- Bead-spring models ideally suited for addressing general polymer phenomena, testing basic theoretical models
 - Disregards atomistic details
 - Can not quantitatively describe properties like structure, local dynamics
- Systematic coarse grained models can bridge the gap of time and length scales while retaining atomistic characteristics
 - Reduces number of degrees of freedom and increases fundamental time step
 - Captures the atomistic detail needed for correct dynamics from monomer to polymer scale

Acknowledgements

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