

# *Dynamics of Entangled Polymers: Using LAMMPS for Large Scale Simulations*

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# Large Time/Length Scale Simulations

- Many systems one needs large times and length scales
  - Polymers, Membranes, Proteins, Nanoparticles
  - Acceleration algorithms are sometimes useful for structure but rarely for dynamics
- Step 1 Coarse grain – reduces number of atoms by  $\sim 10$  and increase time steps from fs to ps
- Step 2 Build initial state – sometimes easy/sometimes not
- Step 3 Run for a long time on as many processors as possible
  - Number beads  $> 500$ /processor

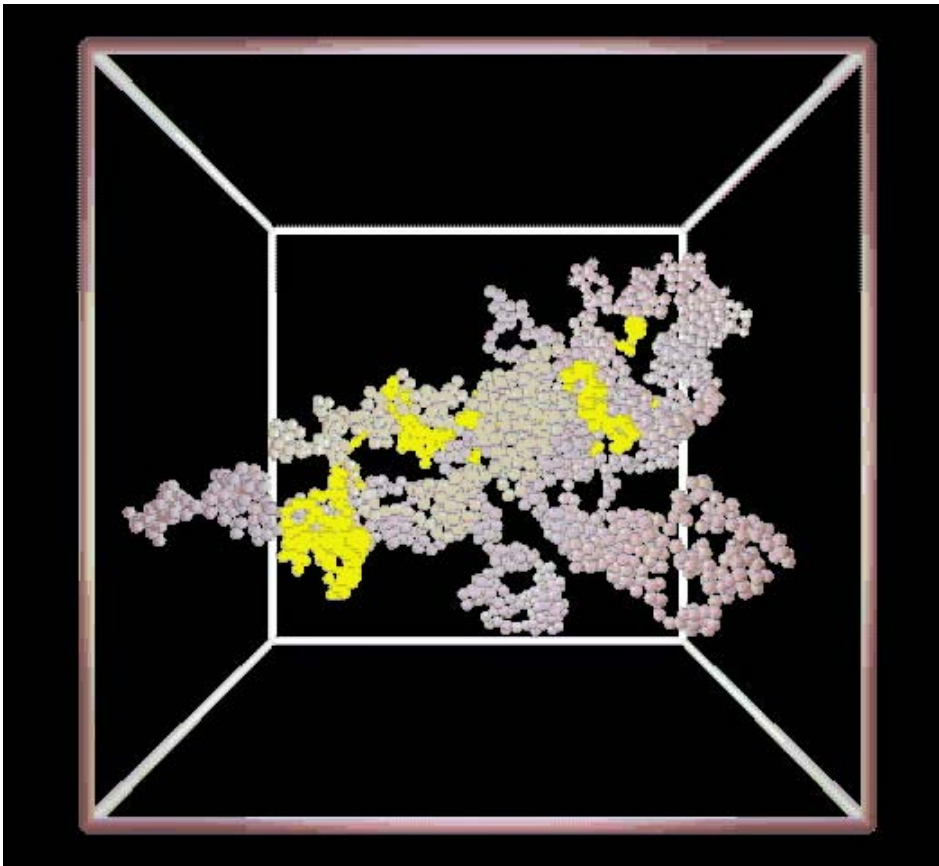
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Brute Force – nothing else will do

# Why are Polymers Interesting?

- Polymers can simultaneously be hard and soft
  - Unique Viscoelastic Behavior

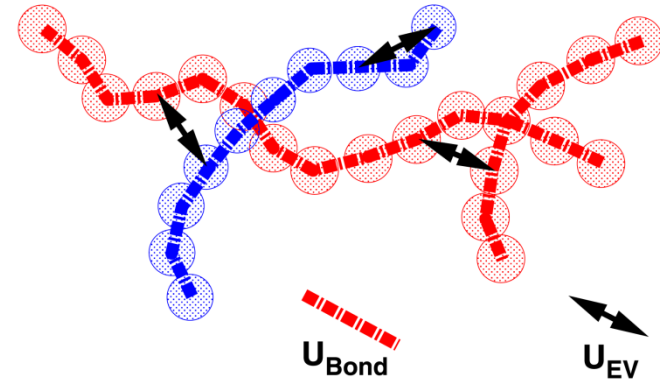


- Motion of a polymer chain is subject to complicated topological constraints

# Bead-Spring Model

- Short range - excluded volume interaction

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



- 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
- Chain stiffness can be included by addition of three-body terms
- MD – velocity-Verlet, time step 0.010 - 0.13 $\tau$

# Advancement Hardware/Software Past 25 Years

- Modified scalar code (Rahman)
- Vector MD – 4 processor Cray XMP – late 1980's
  - $10^4$  monomers – 10 million time steps
- Shared Memory – SGI cluster – 1990's
  - 4-8 processors
- Parallel MD Code – LAMMPS – late-1990's to present
  - 256-2048 processors – Intel and Sun Clusters most recent studies
  - $\frac{1}{2}$  to 1 million monomers – 1-2+ billion time steps
  - 5 million monomers – 300+ million time steps

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- **WARNING – time\_step > 2.14 billion steps bad things happen with LAMMPS**

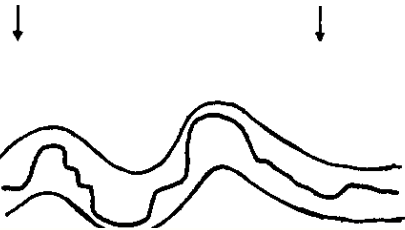
# Motion of Polymers



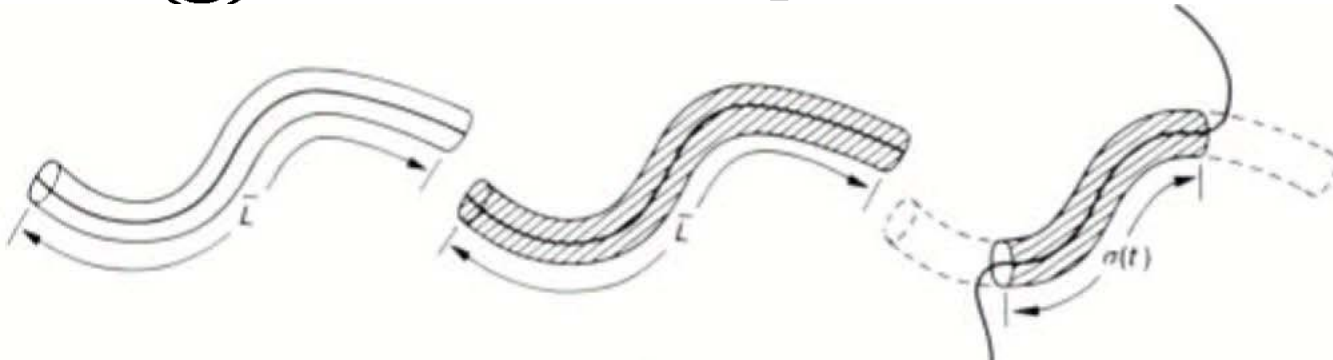
- Statistical Mechanics Approach



- Tube model (Edwards 1967)



- Reptation (de Gennes, 1971)





# Polymer Diffusion

- Simple Liquids

- $D \sim M^{-1}$  ,  $\eta \sim M$

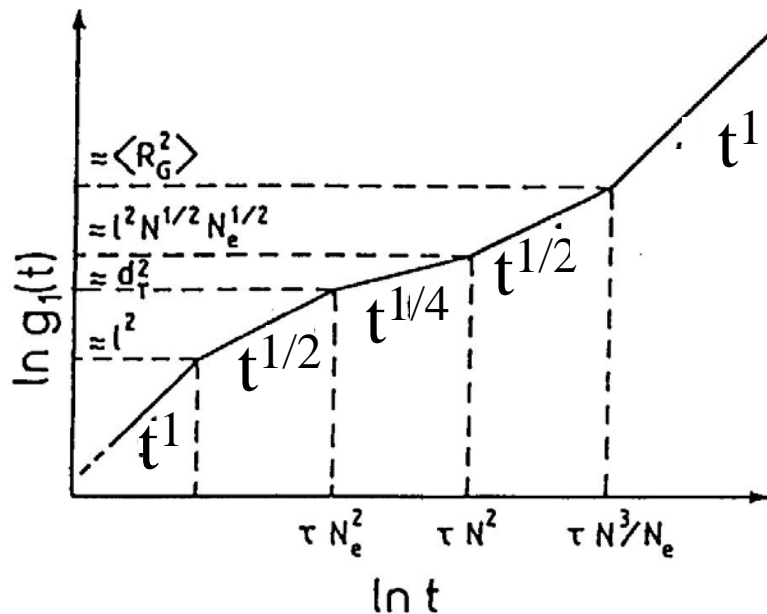
- Short Polymer Chains ( $M < M_e$ )

- Longest relaxation time  $\tau_R \sim M^2$

- Intermediate  $t^{1/2}$  time regime in mean square displacement

- $D \sim M^{-1}$  ,  $\eta \sim M$

- Long Polymer Chains ( $M > M_e$ ) - Reptation



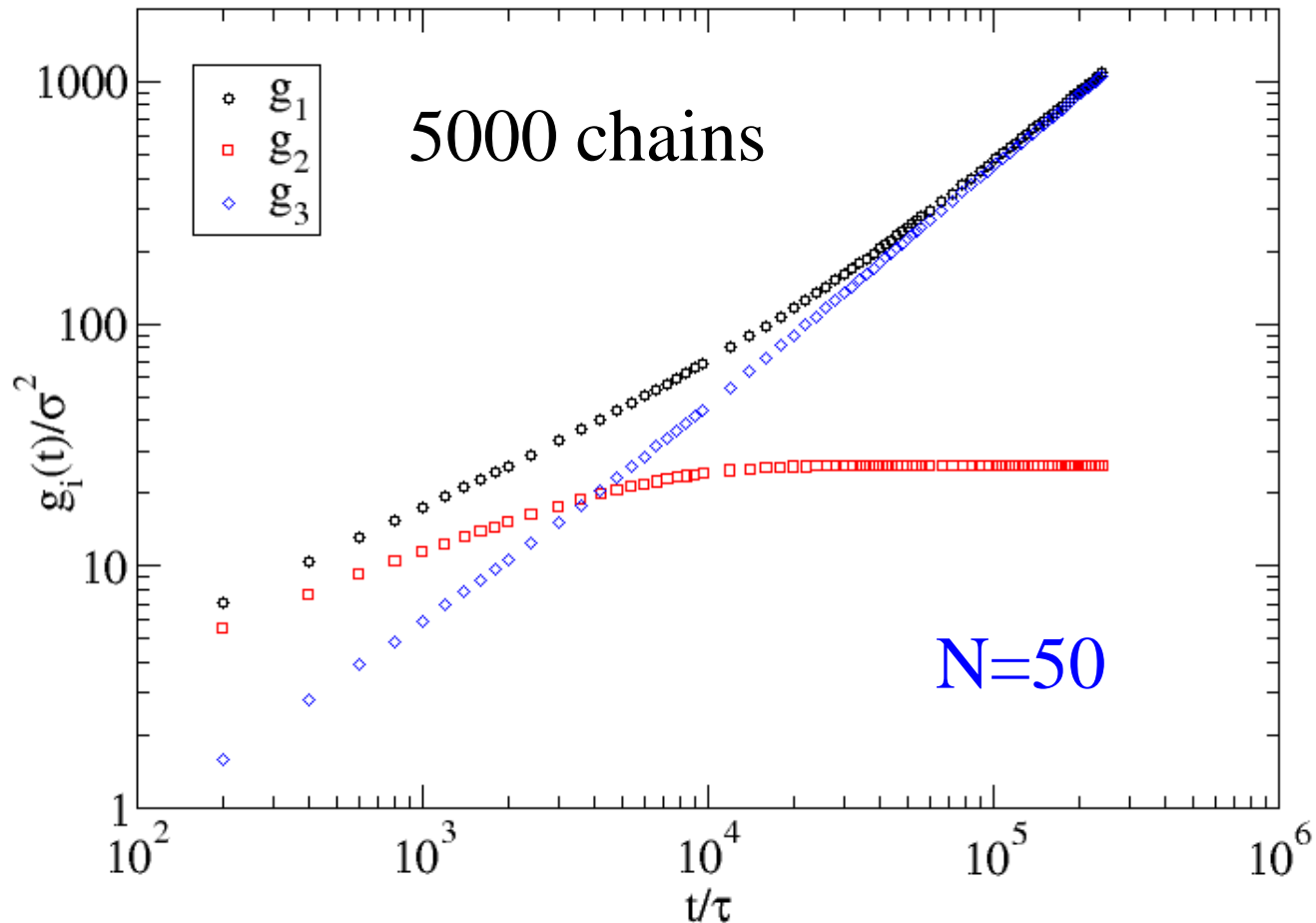
$$D \sim M^{-2}$$

$$\eta \sim M^3$$

$$\tau_d \sim M^3$$

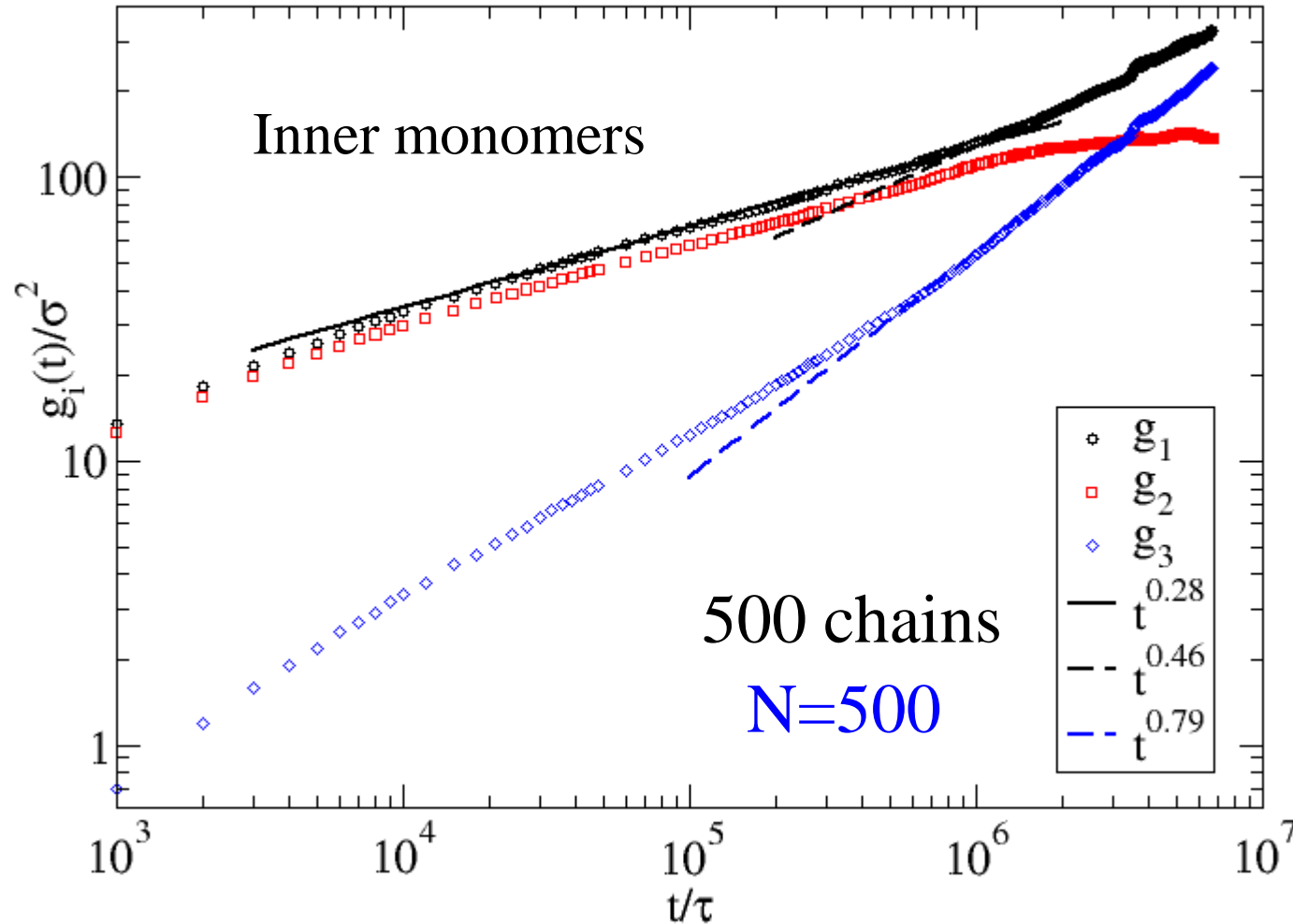
Characteristic signature –  
intermediate  $t^{1/4}$  regime

# Displacement of Unentangled Polymer



- Intermediate  $t^{1/4}$  regime of mass uptake corresponds to Rouse regime of mean squared displacement

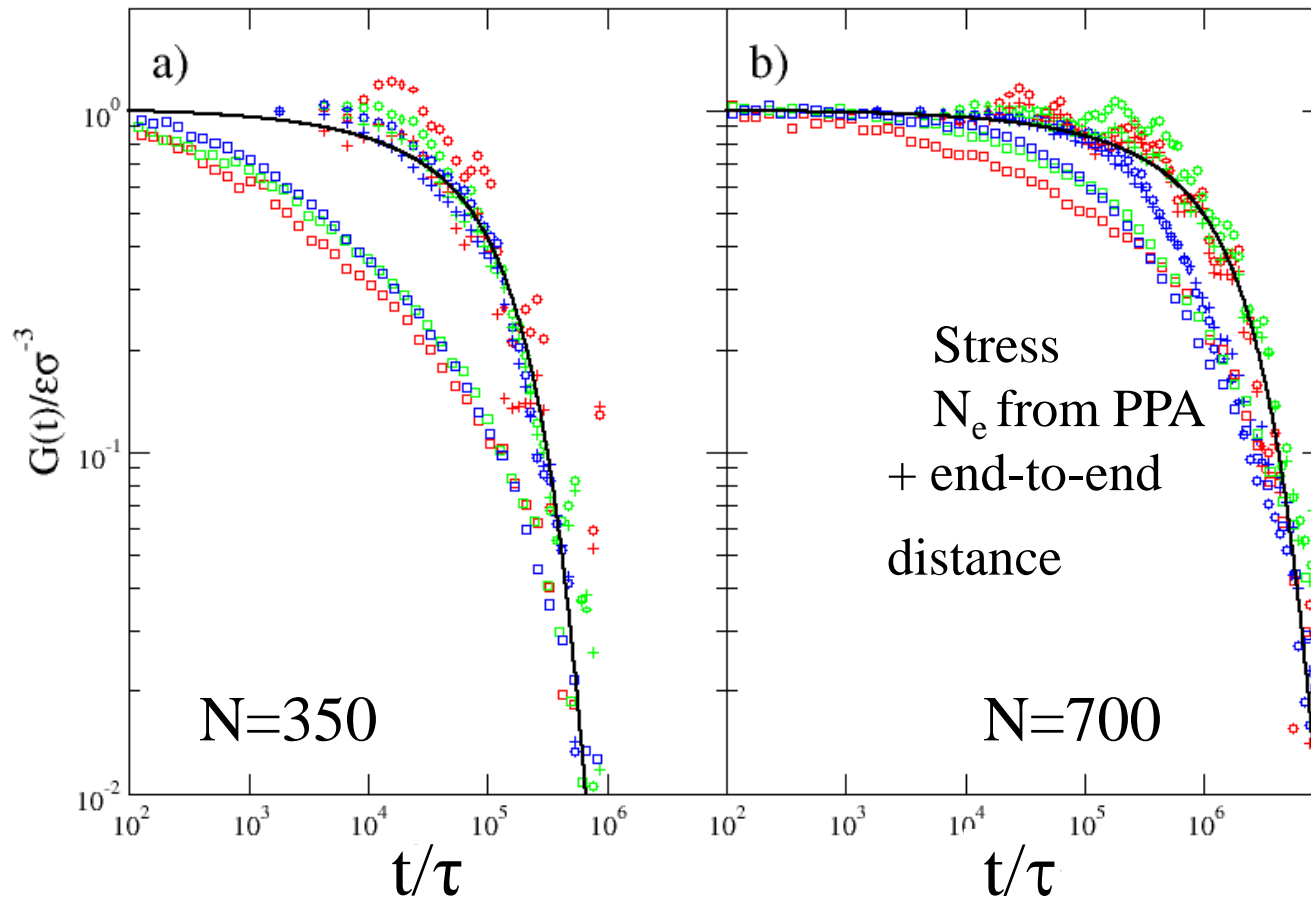
# Displacement of Entangled Polymer



- 100,000 processor hours

# Stress Relaxation - Melt

- Relaxation after elongation –  $\lambda = 2, 3, 4$  (red, green, blue)



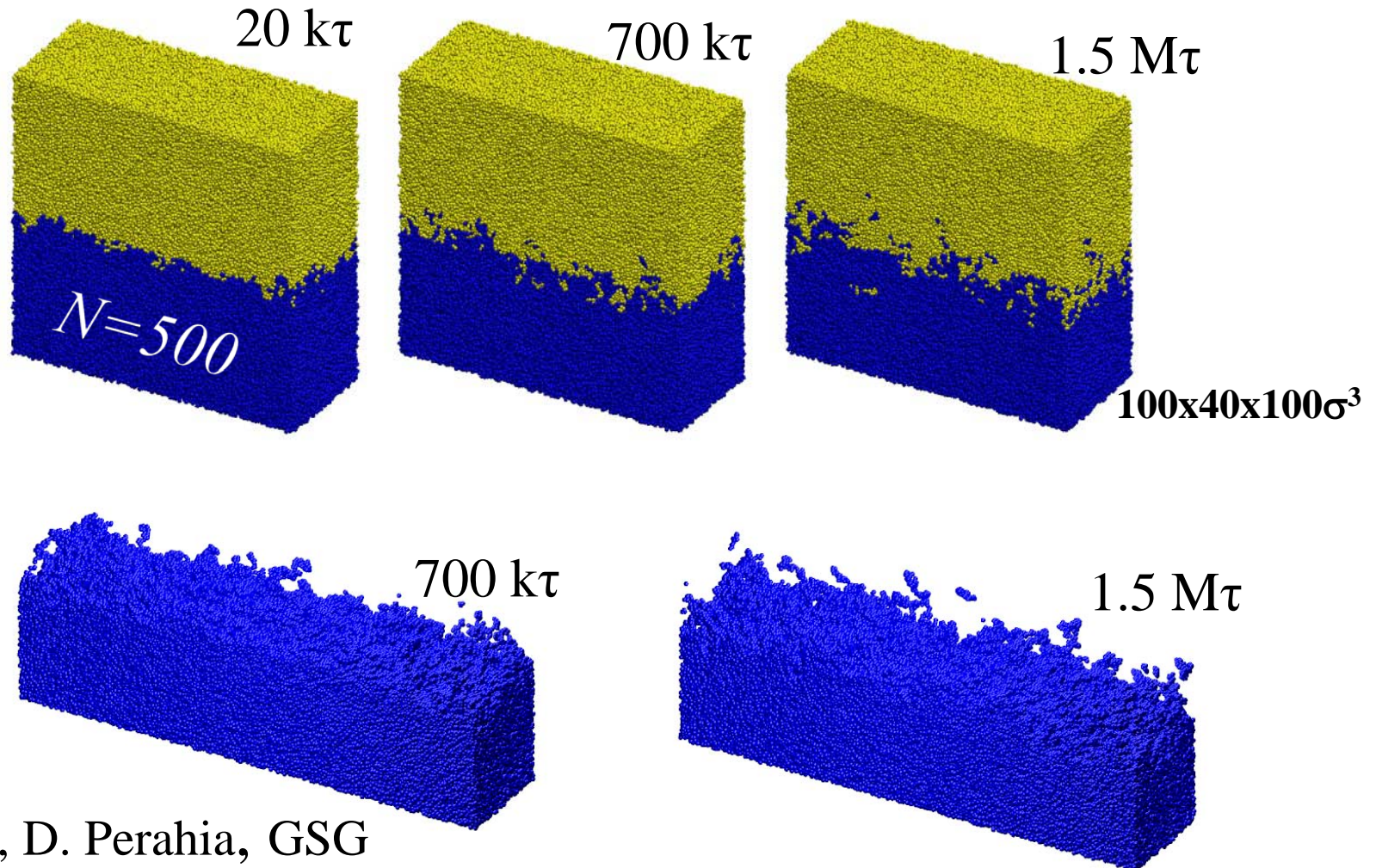
- $N=1000$
- similar results
- $N=3500$
- reach plateau

C. Svaneborg *et al.*,  
PRL (submitted)

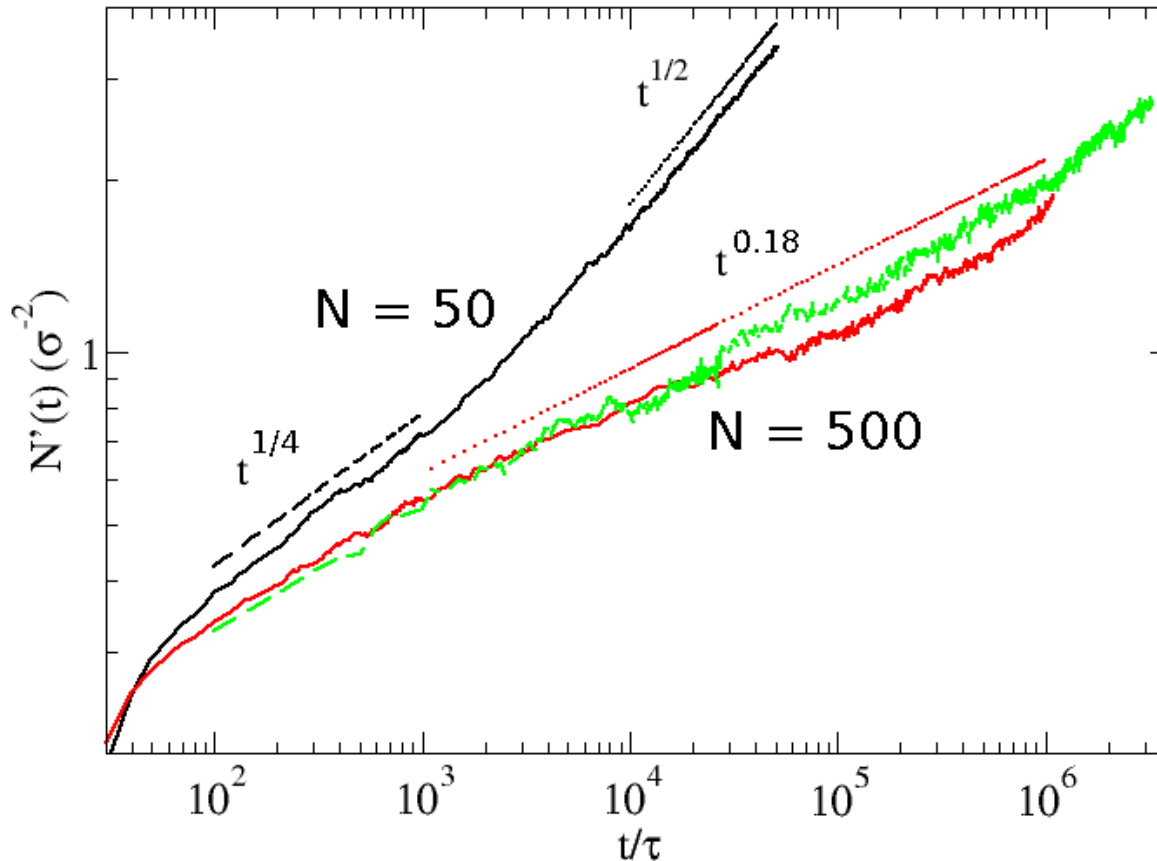
- Relaxation time independent of  $\lambda$ . same terminal relaxation time for all 3 quantities
- Experimental Strains –  $\lambda=1.01$  to  $1.05$

# Self-Healing of Polymer Films

- Development of Entanglements Across an Interface



# How Fast Does a Crack Heal?



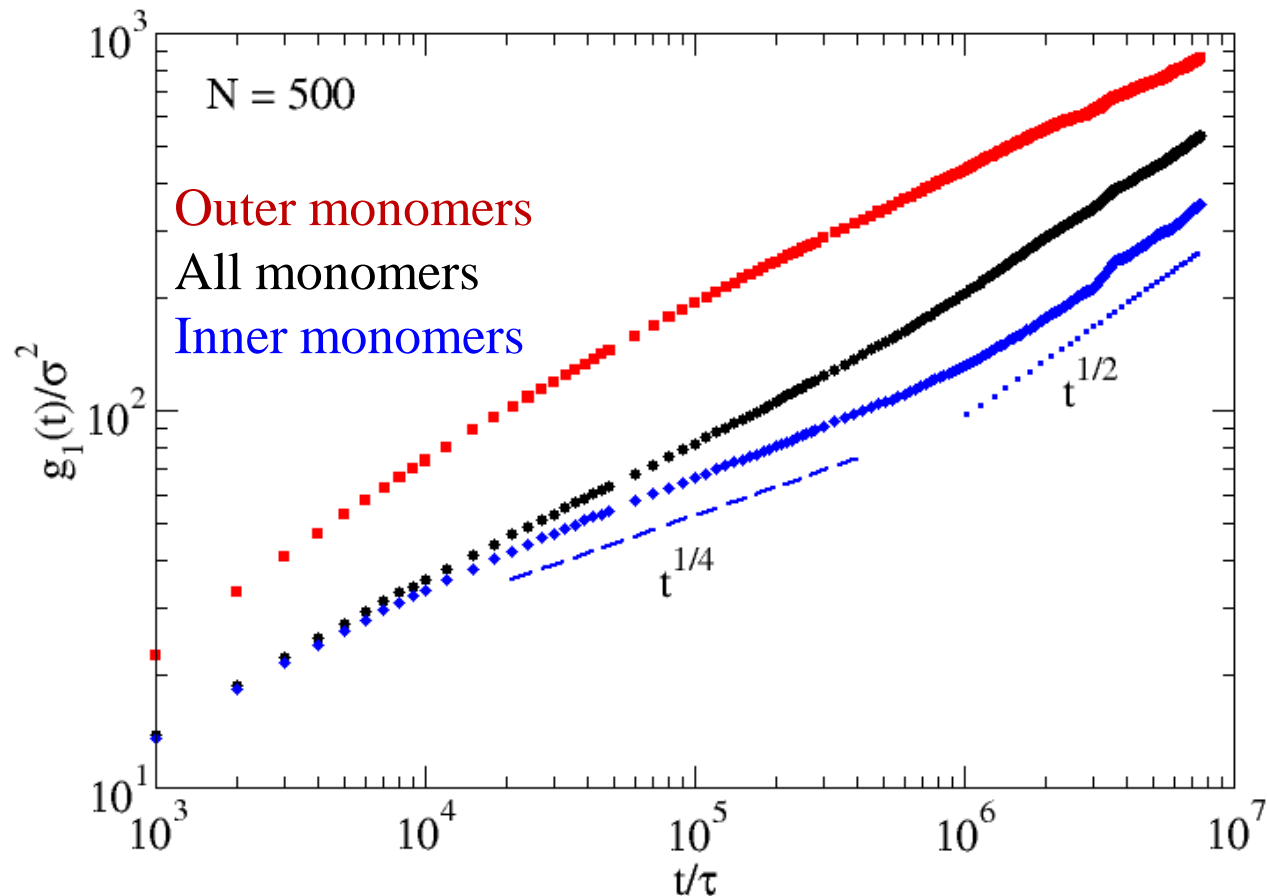
4.8 million  
monomers

3.5 million  
steps/day  
1024 cores

Longest run –  
2.5 million  
node hours

Mass Uptake, Penetration Depth  $z \sim \langle r(t)^2 \rangle^{1/2}$

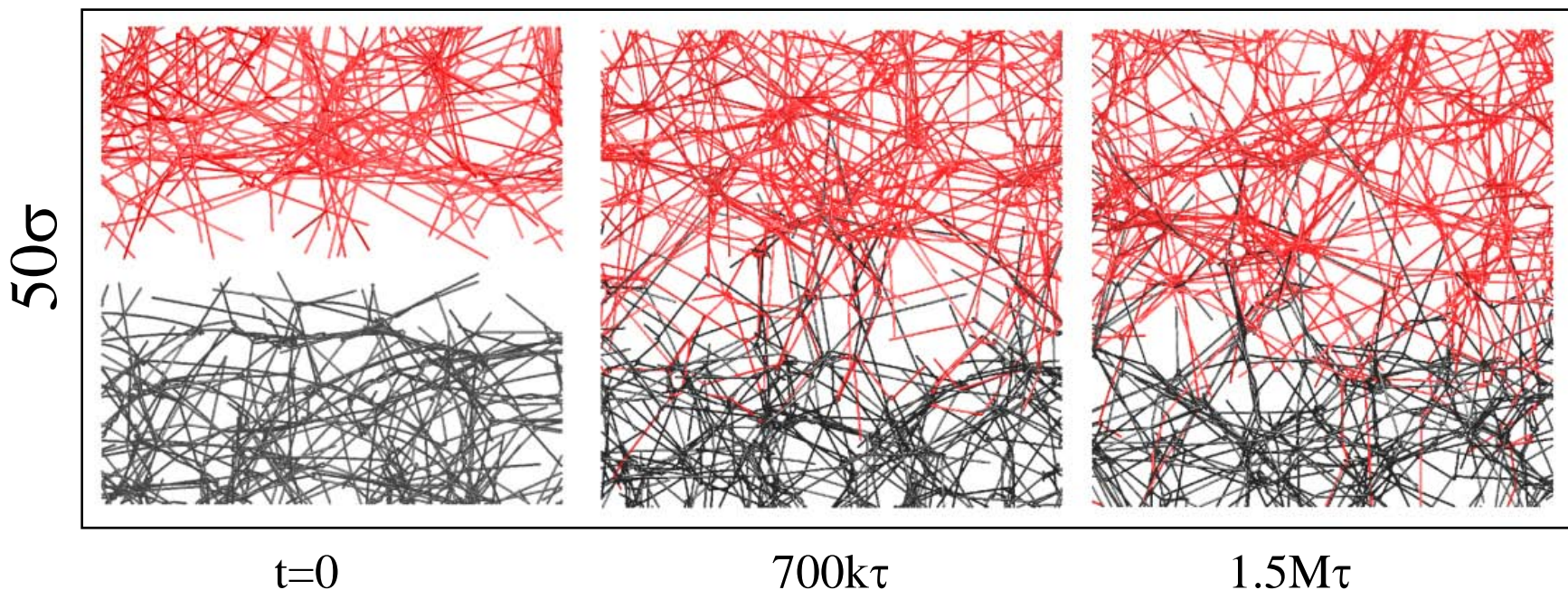
# Healing Depends on Motion of Chain Ends



- Chain ends move much faster, resulting in much faster self-healing than predicted by theory

# Entanglements at Interface

- Primitive Path Analysis
  - Contact all chains simultaneously with ends fixed
  - $(n_{\text{dec}} - 1)$  new beads are placed between adjacent beads on the original chains and process repeated -  $n_{\text{dec}}=4$





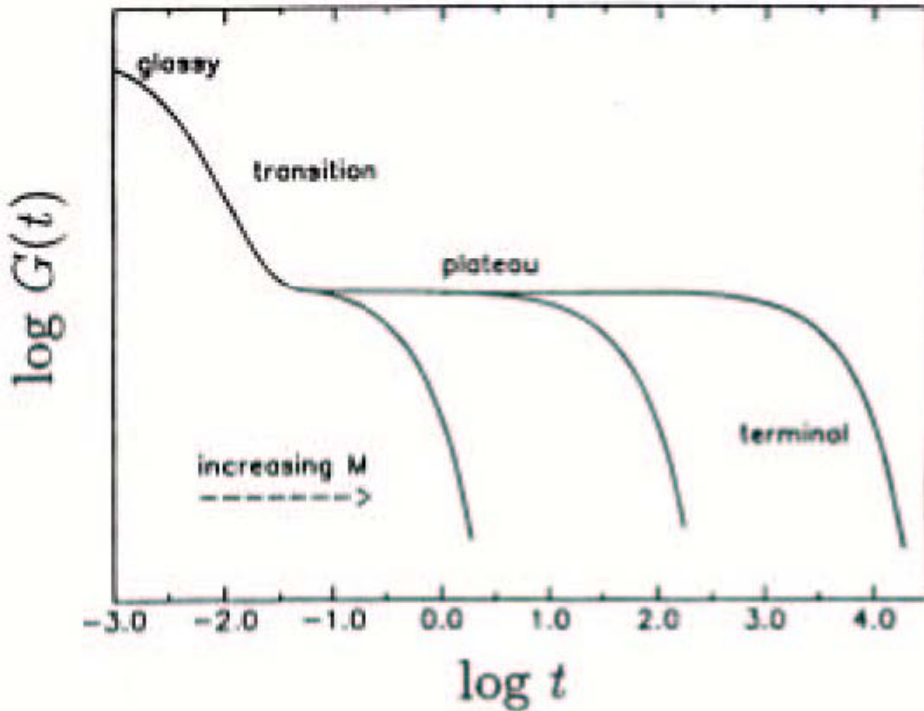
# Future Directions

- Outlook for computer modeling is exciting
  - Faster, cheaper computers
  - Efficient parallel MD codes
- Larger Systems, Longer Chains, Longer Times
- Smaller strain, shear rates
  - Viscosity
  - Relaxation after shear
- Constraint Release - Polydispersity
- Semidilute polymers – explicit solvent
- Primitive Path Dynamics – Melts/Networks
- Branched Polymers, Stars, .....

# Acknowledgement

- Department of Energy – Contract No. ER46456
- Center for Integrated Nano Technologies – Sandia
- New Mexico Computer Applications Center

# Viscoelasticity of Entangled Polymer Melt



Stress Relaxation  $G(t)$  after strain

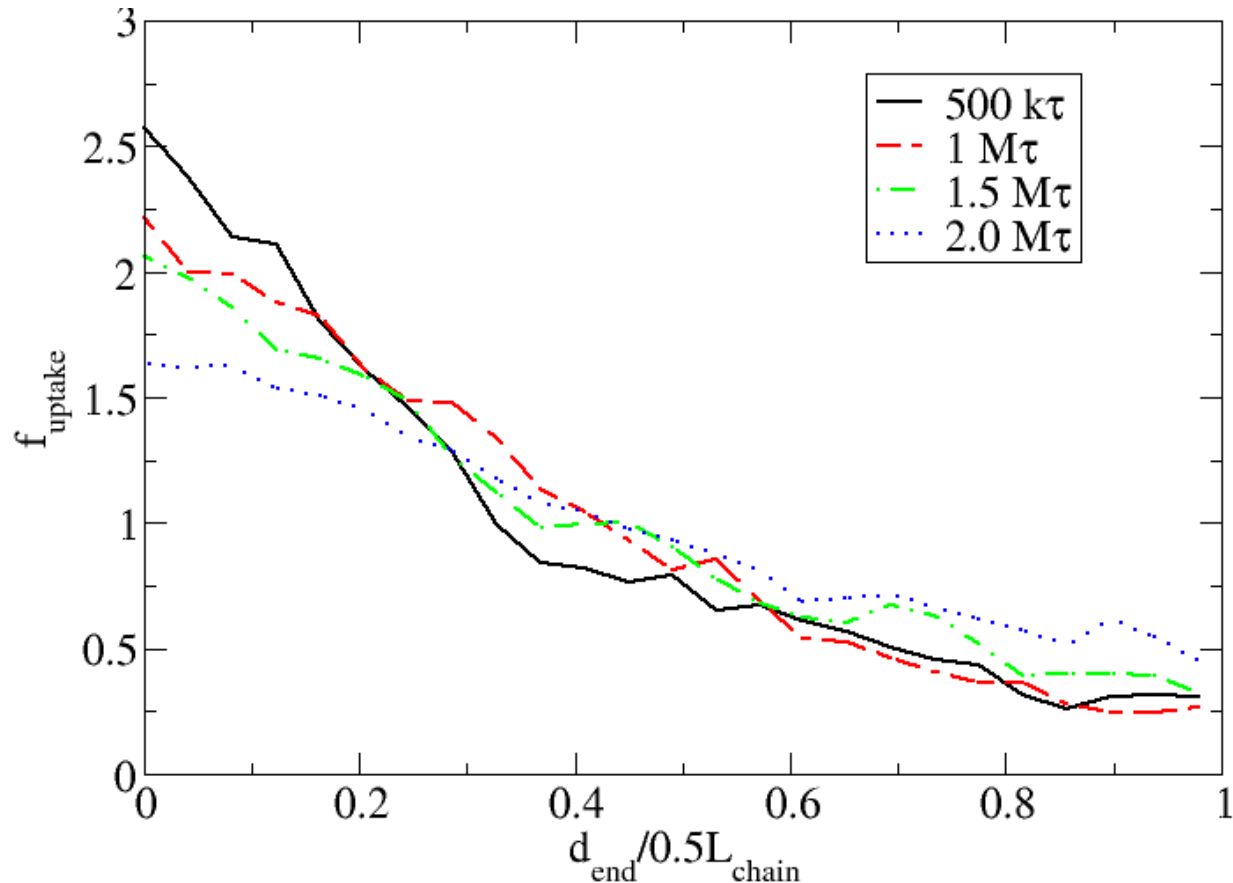
- **Macroscopic**

- Intermediate frequency, time polymer melt acts as a solid
- Long time, low frequency polymer acts as a liquid

- **Microscopic**

- Gaussian coils,  $R \sim N^{1/2}$
- Stress is due to entropy loss of stretched chains
- Polymers as "entropic springs"
- Stress relaxation due to Brownian motion of topologically constraint chains

# What contributes to interdiffusion?



- Chain ends dominate early time interdiffusion
- Rouse motion of chain ends leads to faster mass uptake