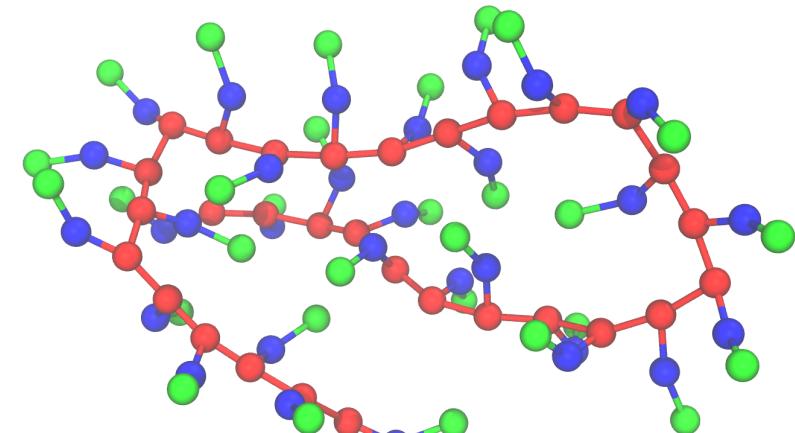
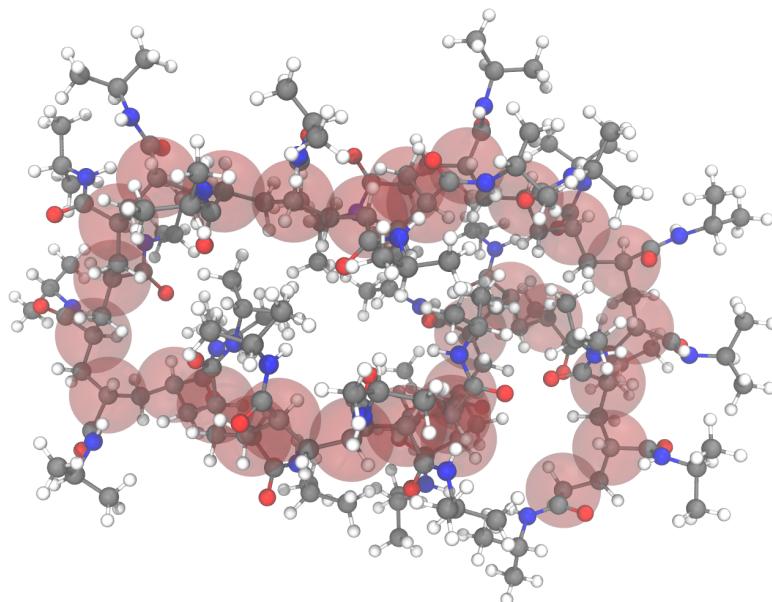


A Coarse-Grained Model for Thermoresponsive Poly(*N*-isopropylacrylamide)



Lauren J. Abbott and Mark J. Stevens

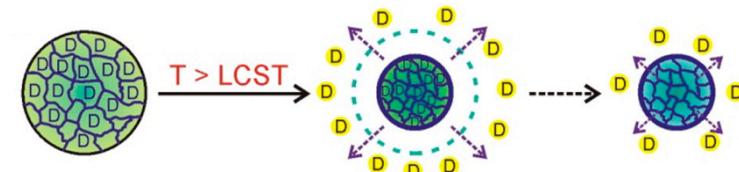
Temperature-responsive behavior useful for many applications

Responsive surface coatings
(e.g., sensors, catalysis)



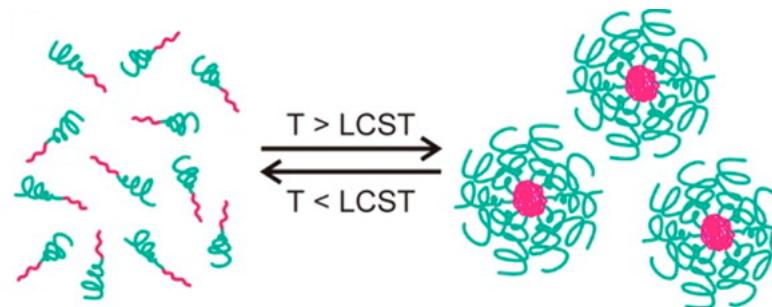
Gibson, *Chem. Soc. Rev.*, 2013

Responsive hydrogels
(e.g., drug delivery, tissue engineering)



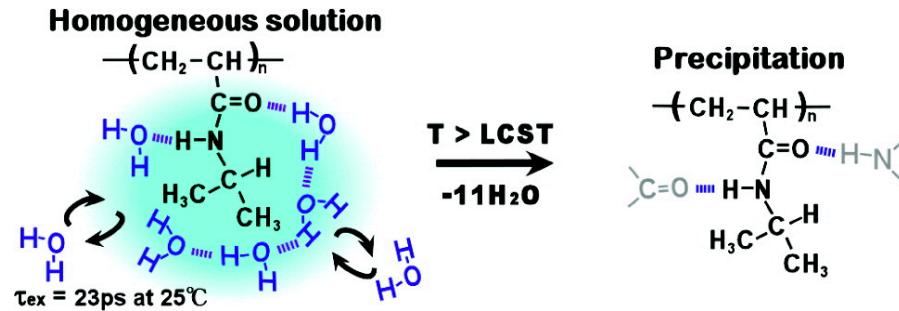
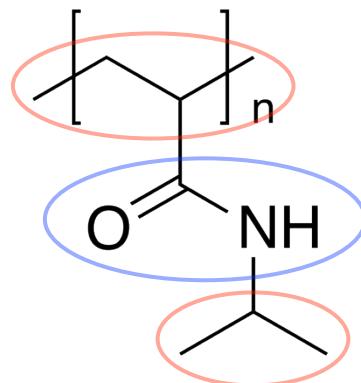
Lee, *Macromolecules*, 2013

Responsive self-assemblies

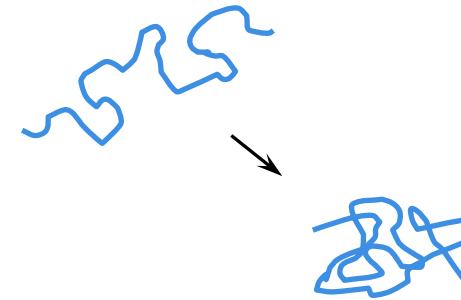
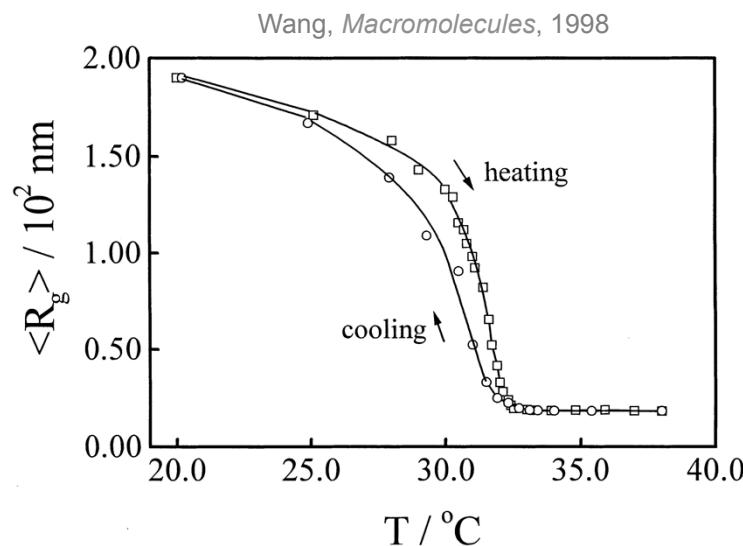


Lee, *Macromolecules*, 2013

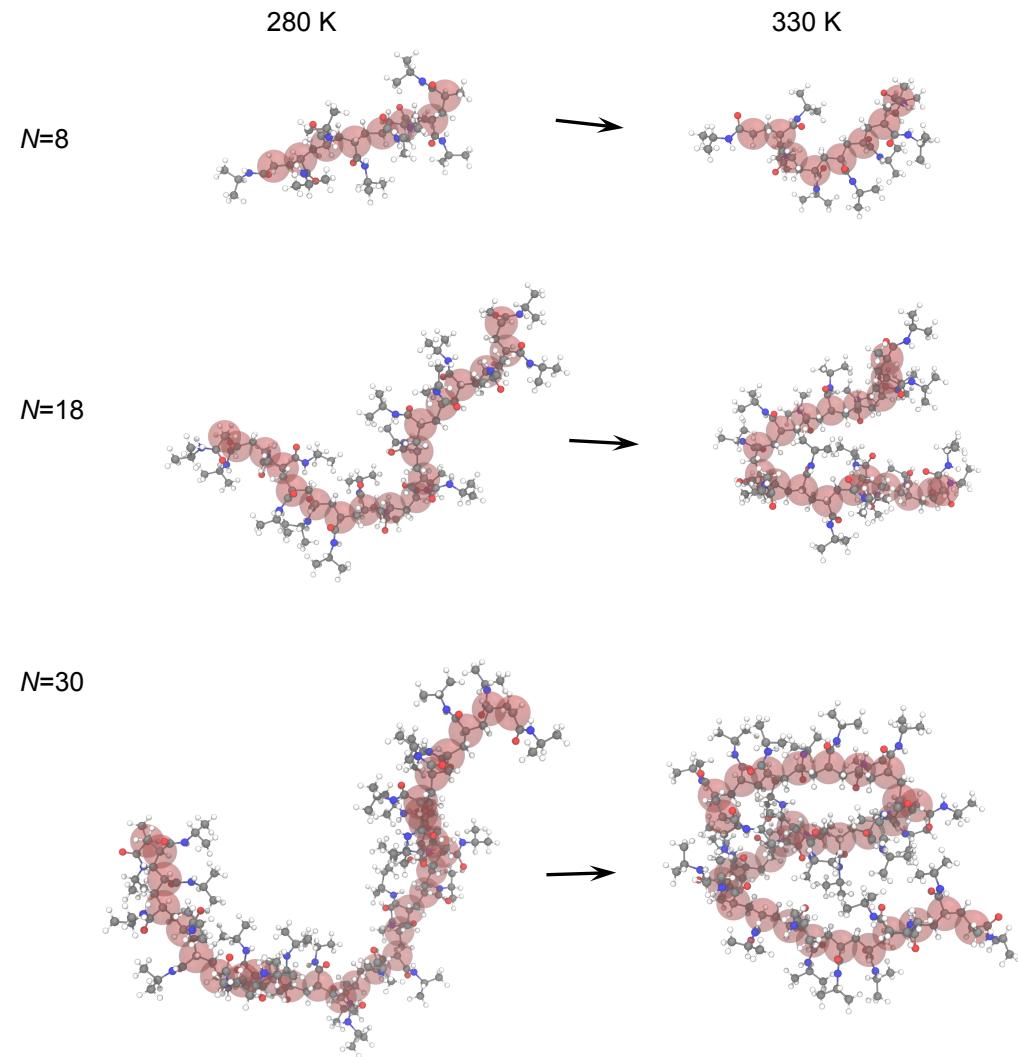
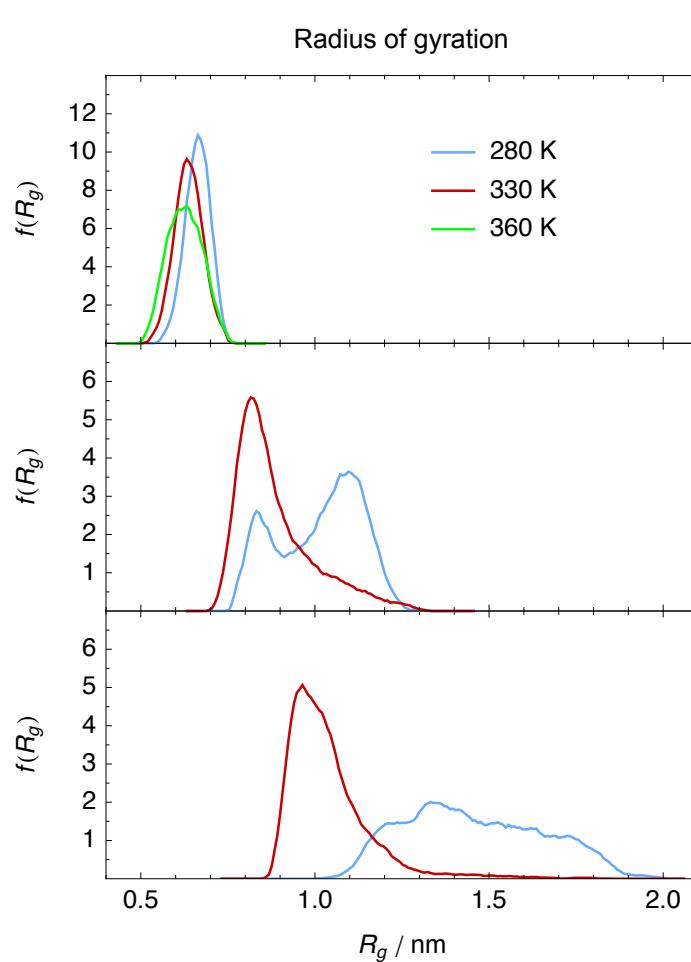
PNIPAM displays a sharp transition at its LCST ~ 32 °C



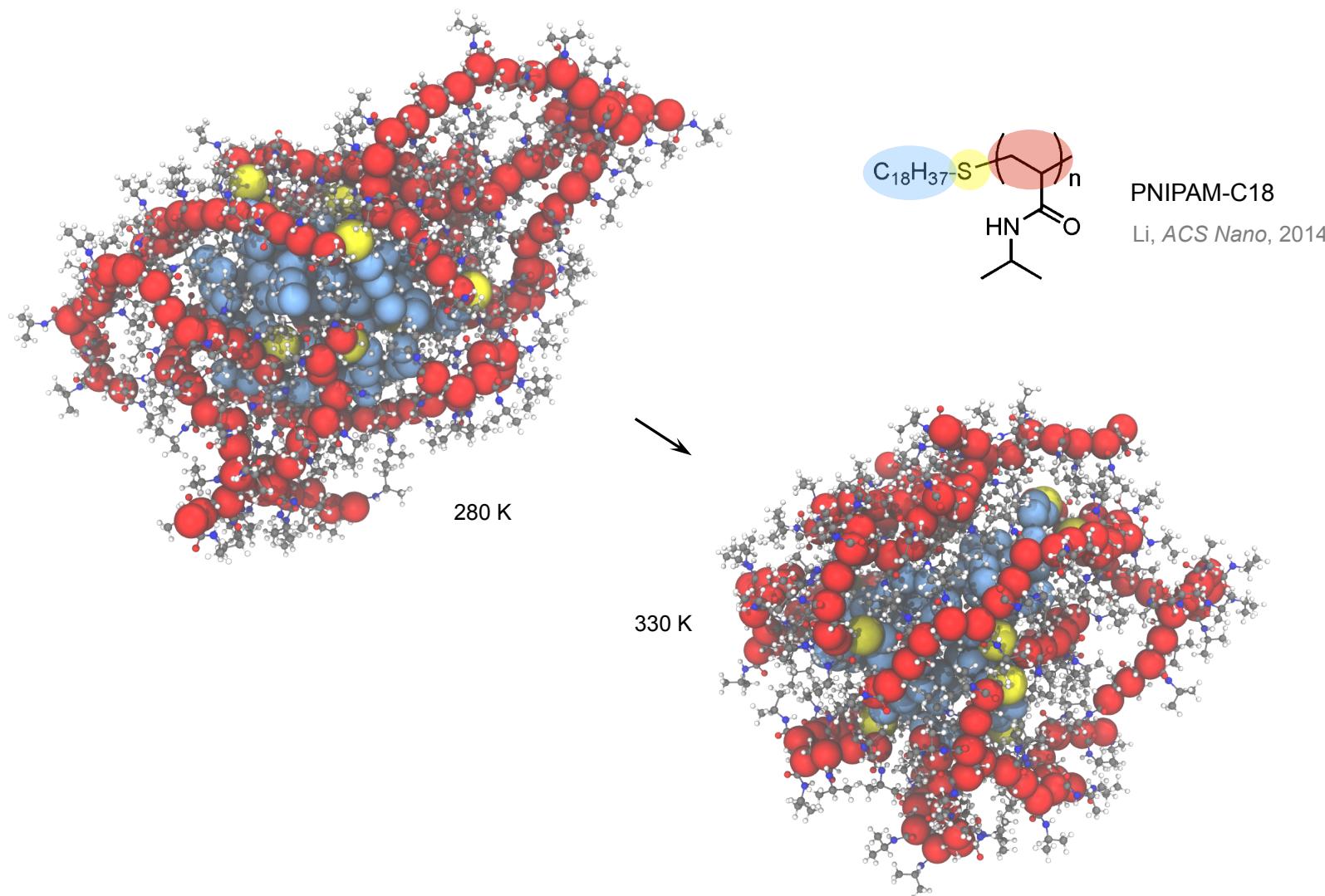
Ono, *J. Am. Chem. Soc.*, 2006



Atomistic simulations capture single-chain behavior



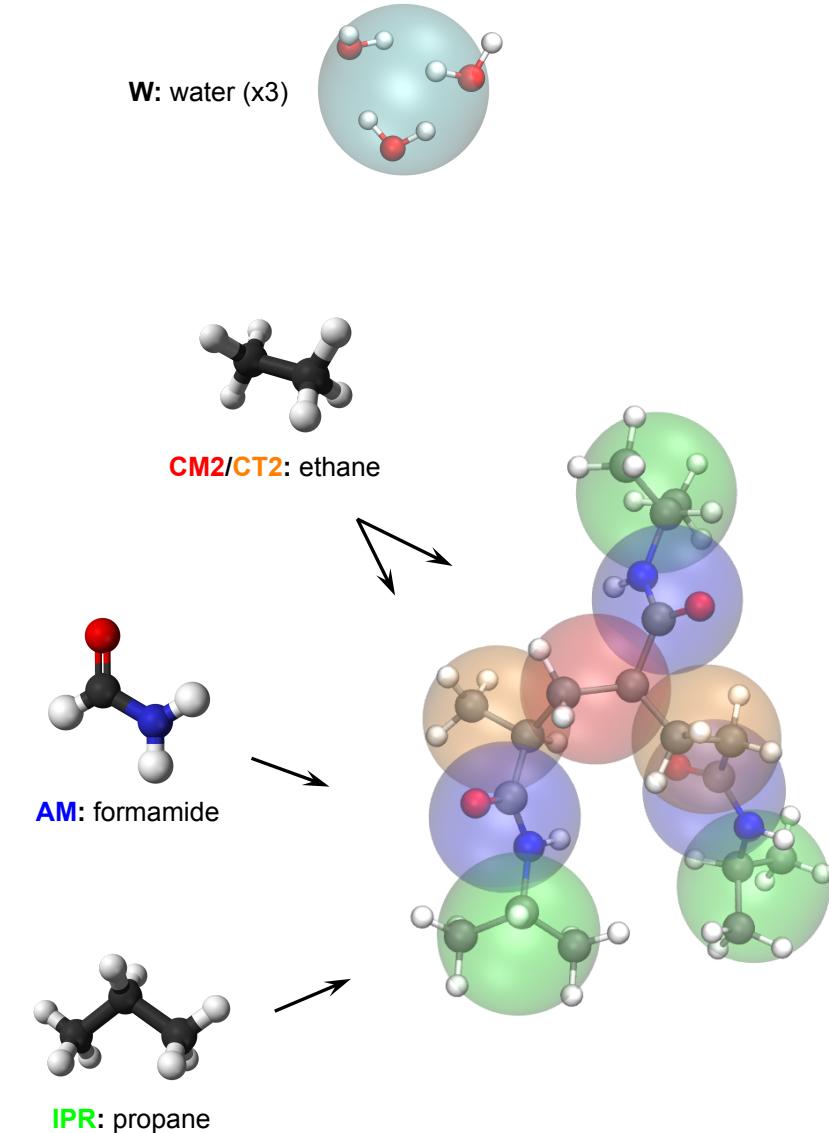
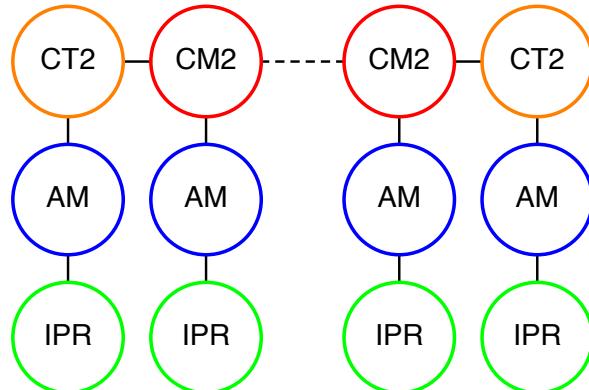
Micelles show temperature-responsive behavior



CG model for PNIPAM

Simulation details:

- Following methodology of Shinoda-DeVane-Klein (SDK) CG force field
- CG beads correspond to ~3 heavy atoms
- Explicit water bead = 3 water molecules
- Nonbond: LJ potentials fit to experimental data (e.g., surface/interfacial tension)
- Bonded: Gaussian-based potential fit to distributions from atomistic simulations
- Parameterization performed against $N=30$ oligomer
- MD with HOOMD-blue on GPUs

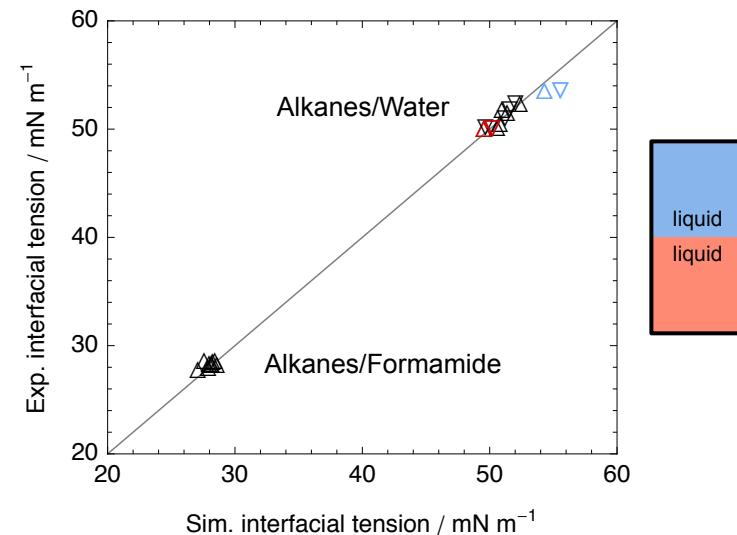
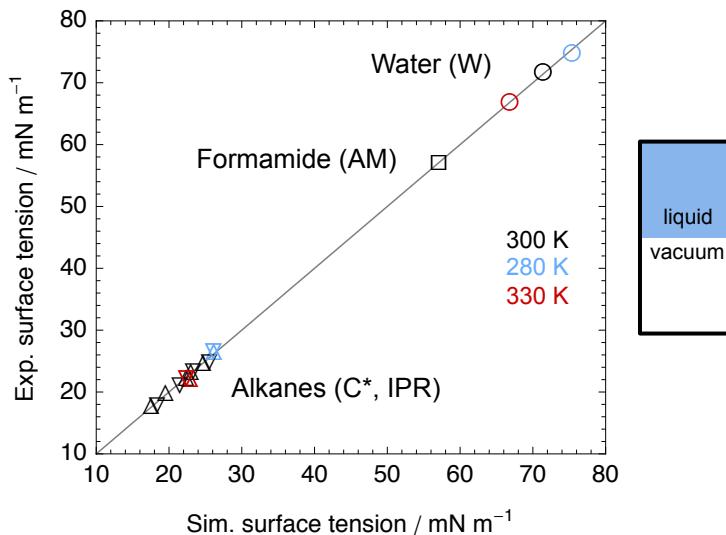
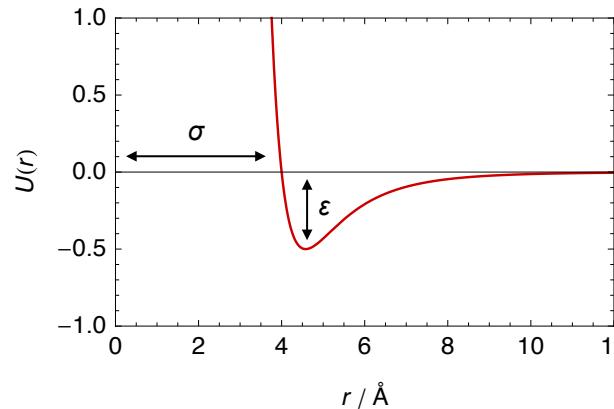


Nonbonded parameters fit to experimental data

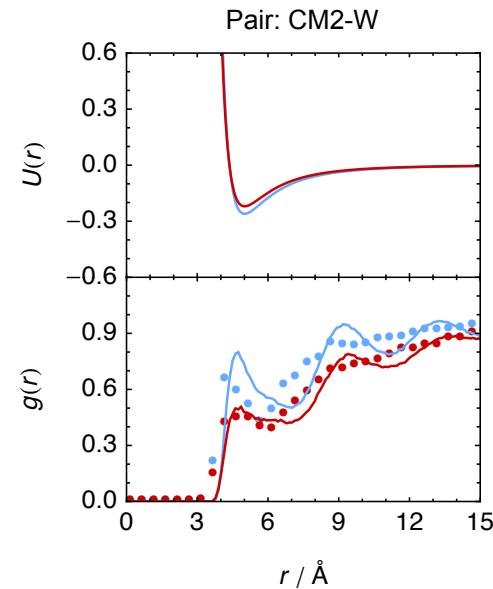
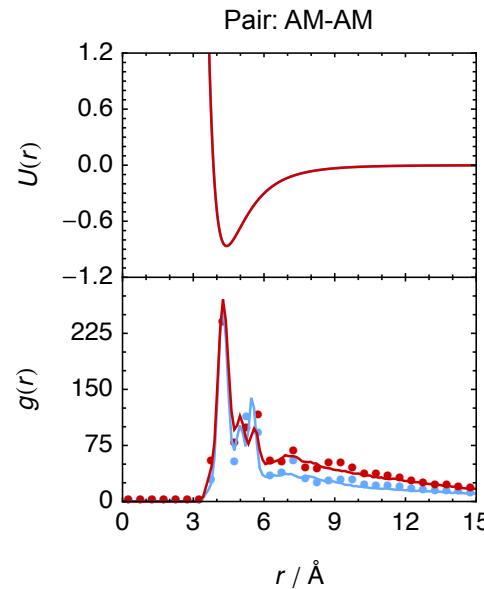
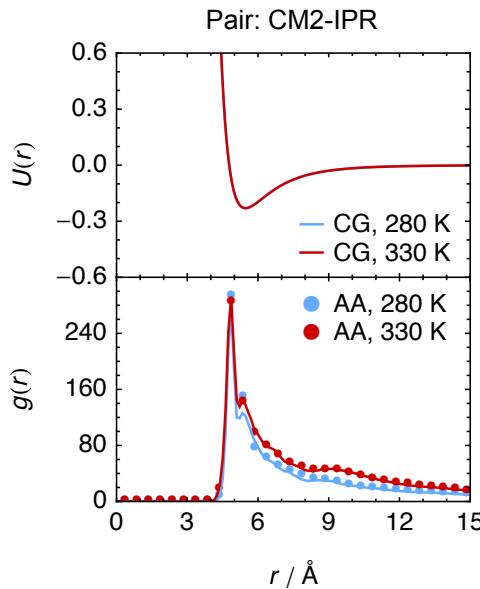
12-4/9-6 Lennard-Jones potentials:

$$U_{\text{LJ}}^{a-b}(r_{ij}) = c_{ab}\epsilon_{ij} \left[\left(\frac{\sigma_{ij}}{r_{ij}} \right)^a - \left(\frac{\sigma_{ij}}{r_{ij}} \right)^b \right]$$

Shinoda, *Mol Simul*, 2007



Adjusted for PNIPAM model to better match RDFs



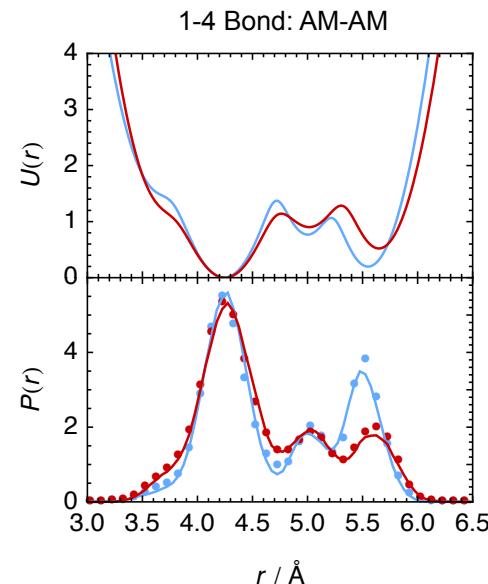
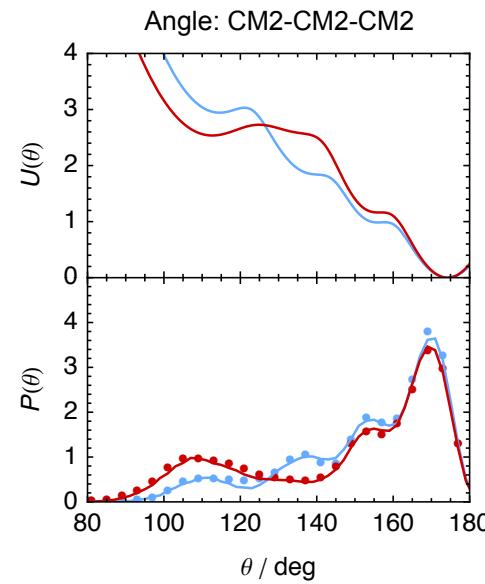
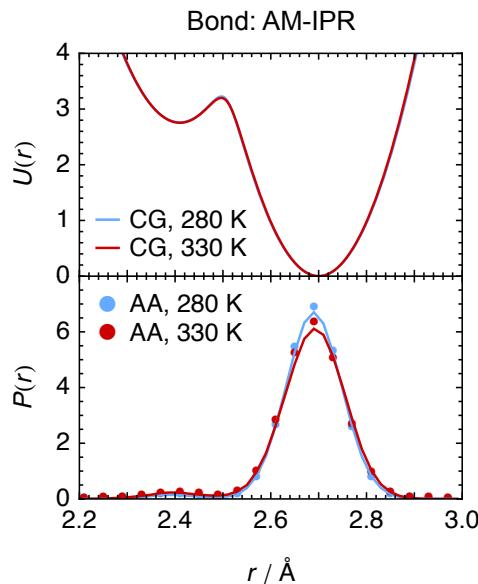
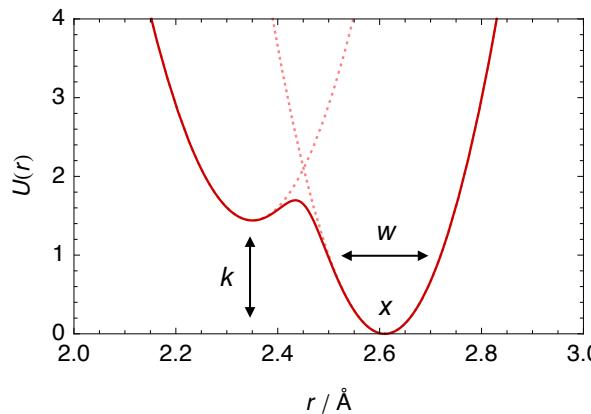
Temperature-dependent nonbonded parameters:
CM2-W, CT2-W, and IPR-W (hydrophobic beads with water)

Bonded parameters fit to atomistic simulations

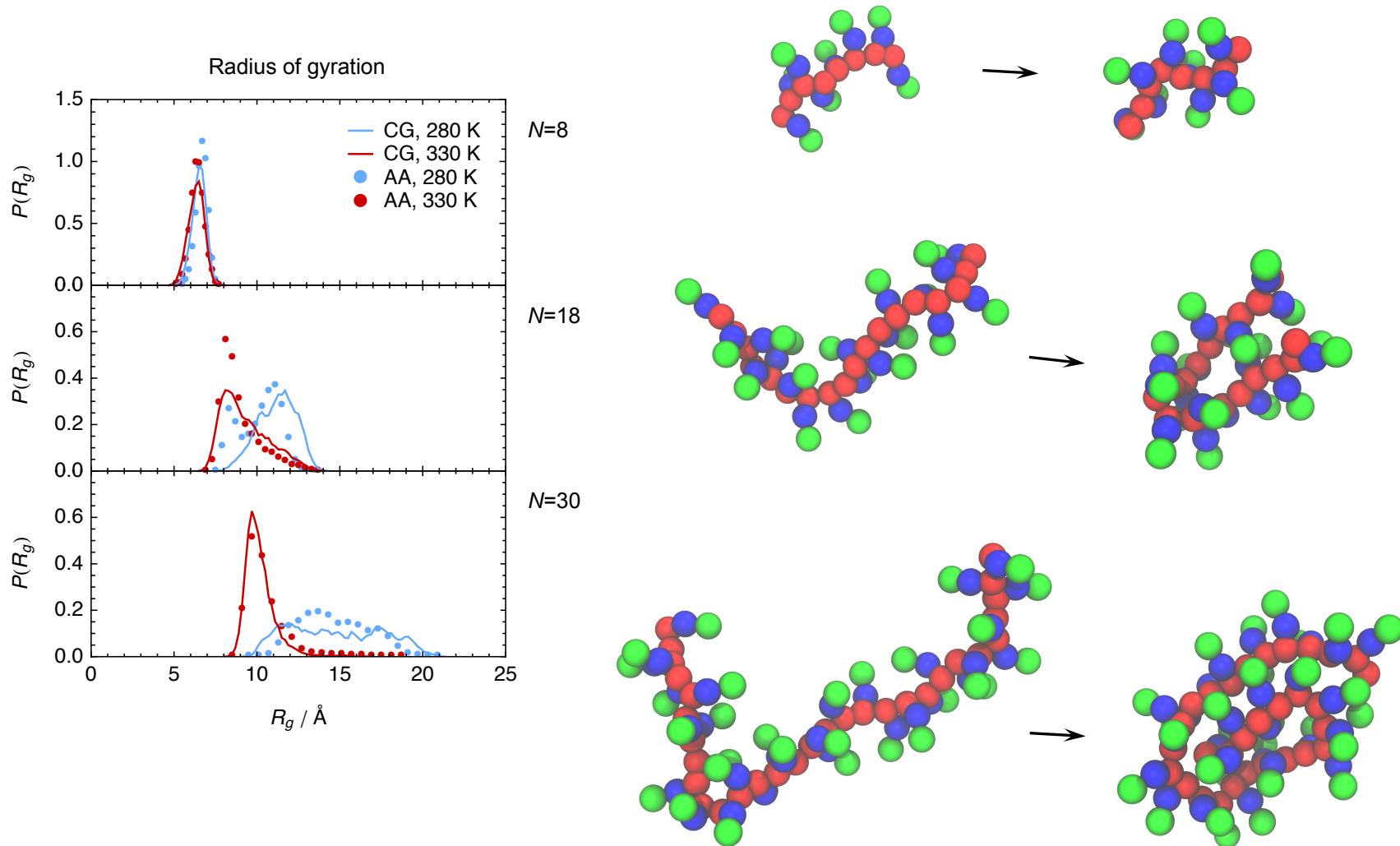
Gaussian-based potentials:

$$U(x, T) = c - k_B T \ln \sum_n k_n \exp \left[-\frac{(x - x_n)^2}{2w_n^2} \right]$$

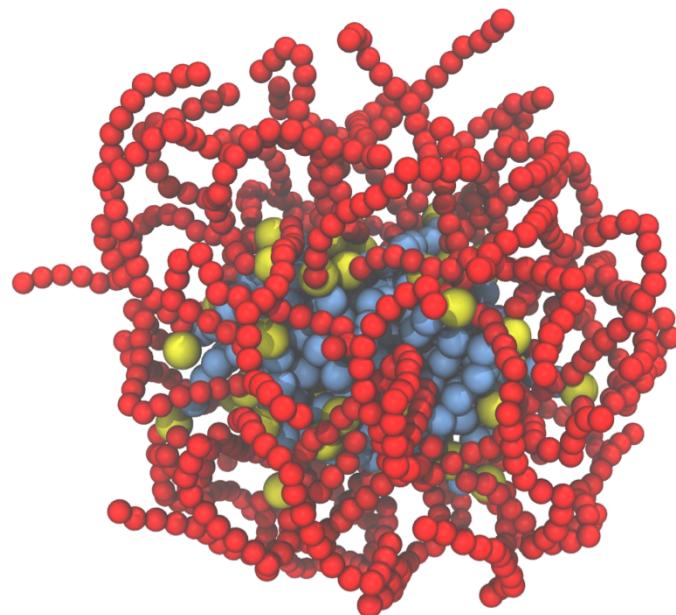
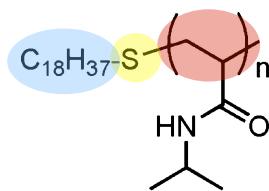
Milano, *J Polym Sci Part B*, 2005



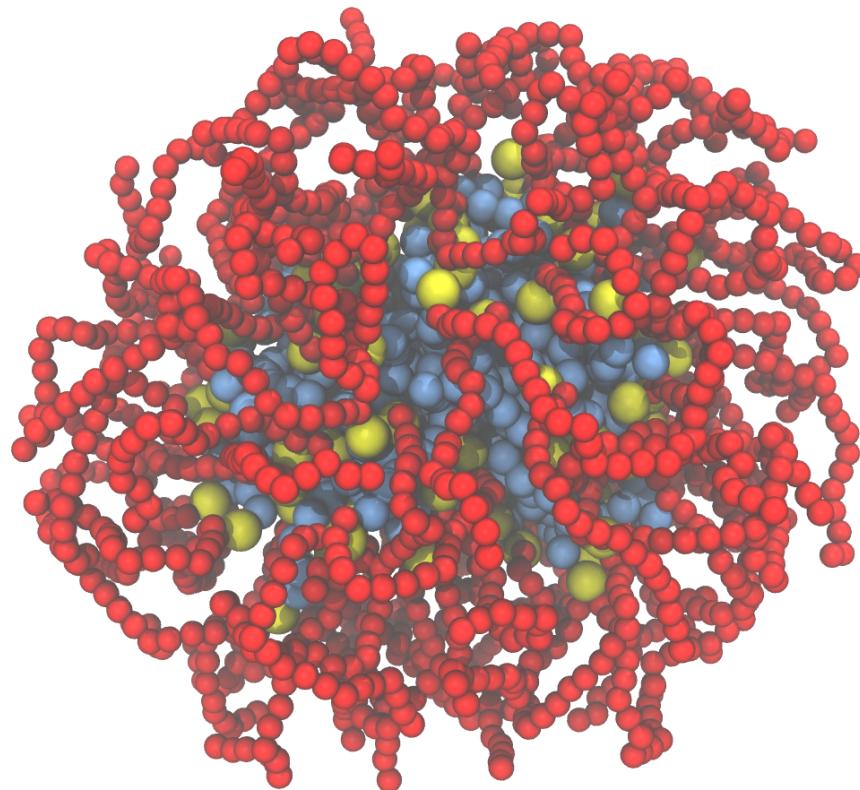
CG model yields correct single-chain behavior



CG model able to simulate micelle self-assembly



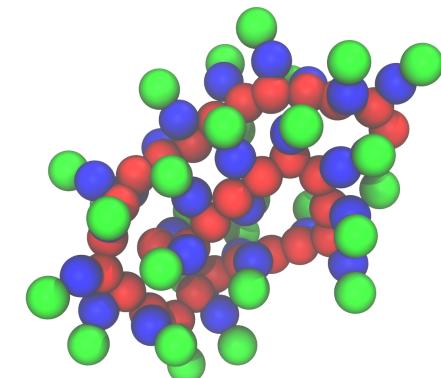
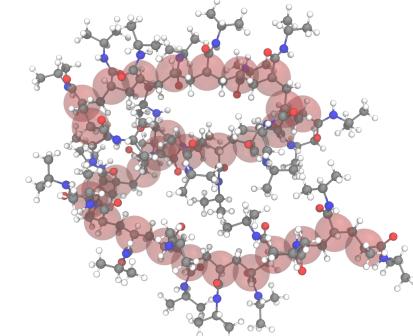
$M = 50,330$ K



$M = 100, 330 \text{ K}$

Summary

- Atomistic simulations of PNIPAM oligomers produce correct single-chain behavior and chain-length dependence
- Small micelles show temperature-dependence, but at the limit of what can be achieved with atomistic models
- Following SDK CG methodology, PNIPAM CG model fit to experimental data and atomistic simulations
- CG model reproduces single-chain behavior and chain-length dependence of atomistic simulations
- Possible to study self-assembly in larger systems with CG model



(1) LJ Abbott et al, *J Phys Chem B*, **2015**, 119, 3837

(2) LJ Abbott and MJ Stevens, *J Chem Phys*, **2015**, 143, 244901