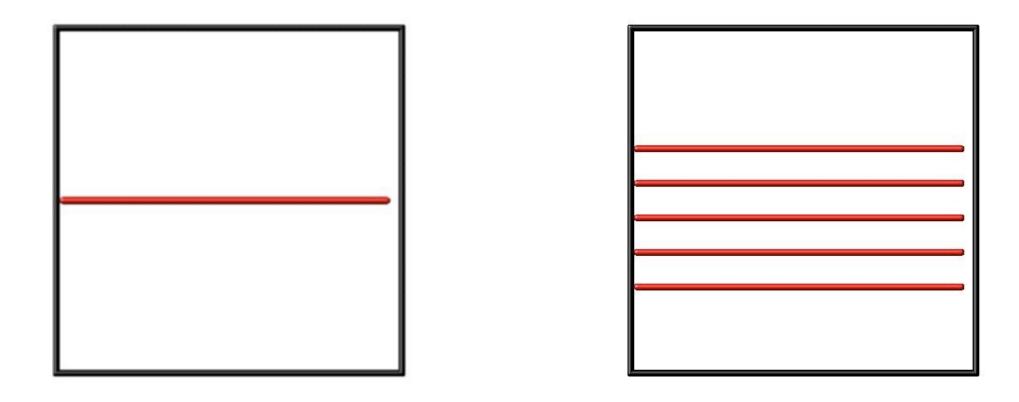
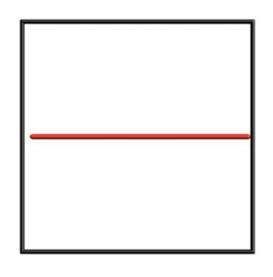
Single and multiple polymer simulations



Single and multiple polymer simulations



$$E = 4\epsilon \left[\left(rac{\sigma}{r}
ight)^{12} - \left(rac{\sigma}{r}
ight)^{6}
ight] \qquad r < r_c \qquad \sigma$$
 (distance units)

 ϵ (energy units)

LJ cutoff (distance units)

#FENE type bond = to define a finite extensible nonlinear elastic (FENE) potential, used for bead-spring polymer models bond_style fene bond_coeff 1 30.0 1.5 1.0 1.0 special_bonds fene

$$R_0$$
 (distance) ϵ (energy)

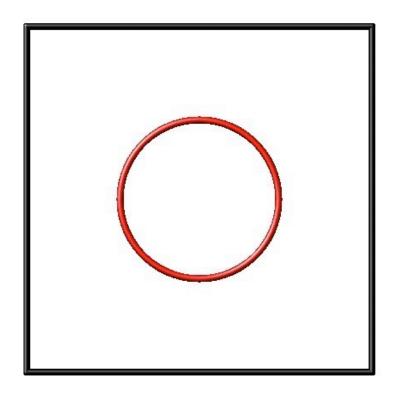
 σ (distance)

K (energy/distance^2)

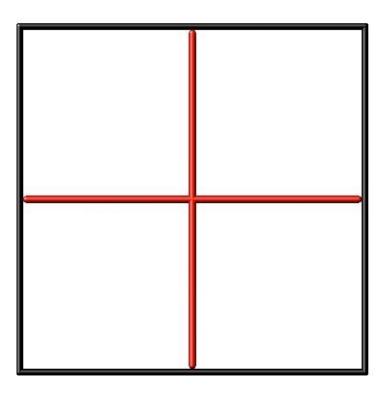
$$E = -0.5 K R_0^2 \ln \left[1 - \left(rac{r}{R_0}
ight)^2
ight] + 4 \epsilon \left[\left(rac{\sigma}{r}
ight)^{12} - \left(rac{\sigma}{r}
ight)^6
ight] + \epsilon$$

atom_style molecular

Circular polymer

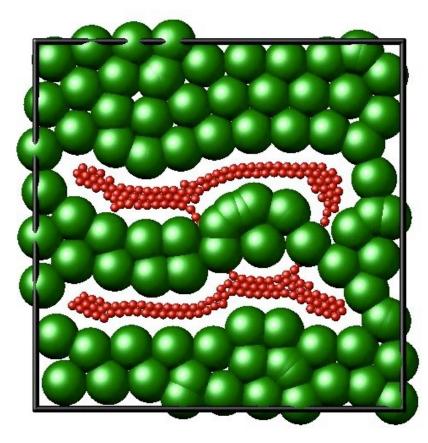


Cross-shaped polymer

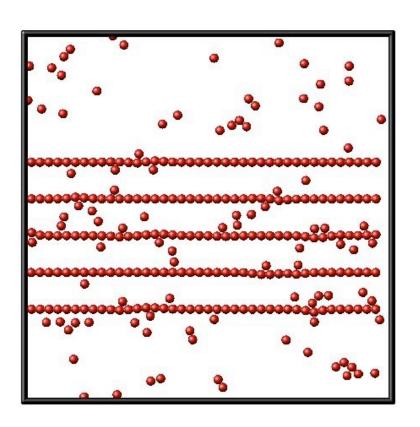


Putting it all together

Polymers with solvent

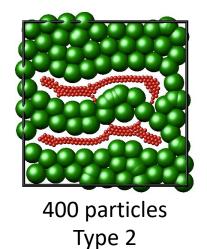


400 particles Type 2



100 particles Type 2

Polymers with solvent



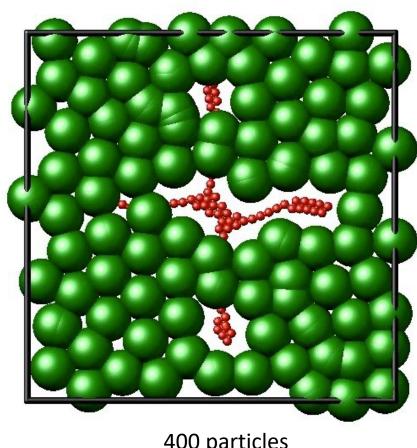
100 particles Type 2

$$E = A iggl[1 + \cos \left(rac{\pi r}{r_c}
ight) iggr] \qquad r < r_c$$

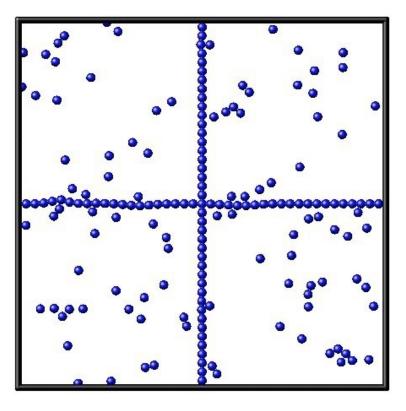
A (energy units)

cutoff (distance units)

Star polymer with particles

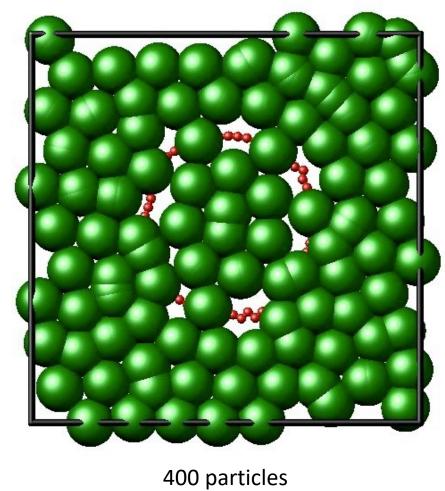


400 particles Type 2

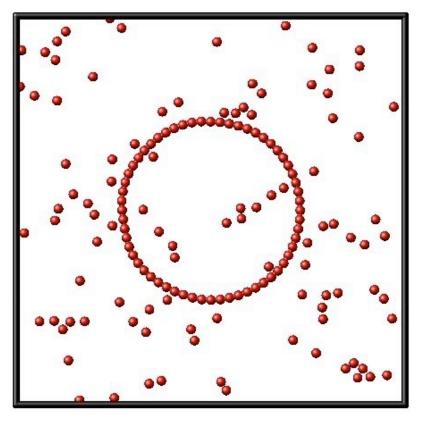


100 particles Type 2

Circular polymer with particles



Type 2



100 particles Type 2

Future Plans

- Incorporate Martini Model
- Establish a bilayer membrane
- Insert membrane protein from RCSB Protein data bank
- Study Radius of gyration in response to changes in environment or biological structures such as membrane
 - pH and charge differences