DOW-Caltech

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Summary of Modifications to CG FF

- Remove Urey-Bradley term (simulations showed it had very little effect)
- Refit nonbond parameters with shorter (10A) cutoffs
- Fit separate polymer-polymer nonbond parameters for phobic, philic. Fit each case to its respective RDF (necessary to get correct density, structure) via genetic algorithm
- Use water-polymer parameters straight from "spring" calculations (slide 4); no optimization

First Approximation to Nonbond Interactions

Make atomistic models of each bead

0

- 2 A beads shown left
- Consider them pairwise alone (in vacuum)
- Attach the centers of the beads by a spring (red arrow left) at a certain distance
- Do some dynamics (let beads move around).
 Calculate average net nonbond interaction between beads at the fixed spring distance
- Shorten spring, repeat (0.75 < r < 30 A)
- Fit a Morse function to E(r)
- Optimize parameters to give correct total and partial distribution functions (genetic algorithm)

Coarse-Grain Parameters: Bond (1-2)

Bond Type	K	r0
MA-Me	73.534	2.507
MA-MB1	61.947	2.492
MB1-MB2	2.047	4.713
EB-MB2	4.091	5.065
MB2-MB2	2.549	5.062
MA-MA	89.374	2.863
A-MA	76.578	2.772
MA-OH	109.422	2.519
A-MB1	55.244	2.472
A-A	83.514	2.706
MB1-EB	2.409	4.855

Coarse-Grain Parameters: Polymer-Polymer for Phobic (Morse Potential)

	A-A	A-MB1	A-EB	A-MA	A-Me	A-OH	A-MB2
D	0.71604	1.04719	1.08090	2.34742	0.88781	28.37712	1.45580
α	1.02535	1.86598	1.12633	2.26192	2.36577	0.40997	1.47176
r0	6.37085	5.30797	4.45610	4.50869	4.88589	2.56688	3.53001
	MB1-MB1	MB1-EB	MB1-MA	MB1-Me	MB1-OH	MB1-MB2	EB-EB
D	0.02668	0.000004	2.07183	0.00756	0.111771	0.00125	0.00004
α	1.02669	2.65552	1.19196	1.67879	2.40114	2.00946	3.26715
r0	3.93748	4.79364	3.94293	8.90981	4.76278	5.83608	5.93878
	EB-MA	EB-Me	EB-OH	EB-MB2	MA-MA	MA-Me	MA-OH
D	1.15628	0.00428	0.00076	0.00036	4.19944	1.43461	19.90436
α	1.14629	2.50045	2.43605	3.24620	0.53061	1.28800	0.42935
r0	5.27837	5.52156	6.36716	4.75927	3.64588	3.71336	3.33133
	MA-MB2	Me-Me	Me-OH	Me-MB2	OH-OH	OH-MB2	MB2-MB2
D	1.54776	3.93E-2	8.87E-1	6.07E-1	1.21E-2	4.13E-2	5.67E-1
α	1.17607	1.37015	2.89465	1.84133	2.28496	2.67778	1.26988
r0	7.09493	6.75477	4.40895	4.14551	4.13427	4.32086	6.56236

Water-Polymer Nonbond Parameters

	A-W	MB1-W	EB-W	MA-W	Me-W	OH-W	MB2-W
D	1.27E-006	3.27985	0.00018	9.42E-7	7.81E-9	6.31029	0.00001
α	3.03569	3.26571	3.19533	2.99605	3.56872	1.91161	3.19026
r0	5.13924	2.53969	4.19894	5.23491	5.23188	2.71191	4.74360

Coarse-grain versus Atomistic Modeling: Phobic system





CG: 5 chains, DP = 3000 44,455 beads, 10fs timestep Atomistic: 4 chains, DP = 300 28670 atoms, 1fs timestep

Coarse-grain versus Atomistic Modeling: Philic system





CG: 5 chains, DP = 300038,082 beads, 10fs timestep Atomistic: 4 chains, DP = 300 21,983 atoms, 1fs timestep

Tentative Preliminary Results

Large system for phobic: Density ~0.95 Large system for philic: Still equilibrating

Both systems are still running.



This result is still very preliminary, but so far it looks like our force field reproduces the target structure quite well in the large system



Bond coefficient (harmonic)

$$E_{bond} = K (r - r_0)^2$$

4	5	73.5344	2.50651
4	2	61.9467	2.49174
4	4	89.3737	2.86326
1	4	76.5779	2.77158
4	6	109.422	2.5192
1	2	55.2436	2.47211
1	1	83.5141	2.70619
2	3	2.4094	4.8548

Non-bond parameters (Morse)

$$E_{nonbond} = D \left(e^{-2\alpha (r - r_0)} - 2 e^{-\alpha (r - r_0)} \right)$$

Type 1	Type 2	D (kcal/ mol)	α (A ⁻¹)	r ₀ (Angstrom)
1	1	0.271815	0.998861	4.111223
1	2	0.648291	0.546075	8.616739
1	3	0.711131	1.023365	5.411805
1	4	3.807914	1.598482	1.949401
1	5	0.847399	2.820771	5.011325
1	6	35.77229	0.699786	2.085994
1	7	2.677882	2.239555	3.630308
2	2	0.039274	1.609902	2.021534
2	3	0.000004	1.380162	3.806448
2	4	0.660678	1.588386	3.476449
2	5	0.007617	1.226333	5.534197

Non-bond parameters (Morse)

$$E_{nonbond} = D \left(e^{-2\alpha (r - r_0)} - 2 e^{-\alpha (r - r_0)} \right)$$

Type 1	Type 2	D (kcal/ mol)	α (A ⁻¹)	r ₀ (Angstrom)
2	6	0.163596	2.133694	2.981971
2	7	0.000757	1.80412	6.953686
3	3	0.000014	2.641531	6.290063
3	4	2.054321	0.574267	2.163598
3	5	0.004577	3.486313	4.69669
3	6	0.001231	1.985388	7.062118
3	7	0.000299	2.539202	7.217903
4	4	3.50484	0.396974	2.745834
4	5	1.867358	1.477765	2.369911
4	6	21.7325	0.487019	3.89987
4	7	1.208406	1.405692	6.003881



Could confinement of water in polymers lead to cavitation, hence degradation?

Transduction of subsaturation in the vapor phase of water into negative pressures in the liquid phase

Engineering demonstration



PV curve for water (IAPWS)

When P_{root} , $P_{leaf} < p_{w,sat}$ liquid water is mechanically stable and thermodynamically metastable with respect to the vapor state.





Stroock & Co-workers, Nature (2008) Vol 455 Wagner & Prub, The IAPWS Formulation 1995 for the Thermodynamic Properties of **D**Pdinary Water Substance for General and Scientific Use, JCP (2002), Vol. 31, No. 2,

How does water climb up in a tree?



Directions

- 1. Plants can regenerate their tissues and polymers can not.
- 2. Cavitation could be responsible for degradation of polymer exposed to rain
- 3. Can we predict a PV curve for negative pressures?
- 4. Can we capture the process of cavitation as a function of sub-saturated vapor pressure?
- 5. Goddard-Molinero Water model vs New Molinero model

Ongoing Work

Finish big phobic, big philic, check results

Test water-polymer parameters, moving toward GCMC calculations for equilibrium water content and wetting/drying simulations