

Interfacial Properties of Polydimethylsiloxane/Silica/Water Systems

Ahmed E. Ismail, Gary S. Grest, David R. Heine,
Mesfin Tsige,* and M. J. Stevens

Sandia National Laboratories, *Department of Physics, Southern Illinois University

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Outline

- 1 Motivation
- 2 Methodology
- 3 Interfacial behavior
- 4 Diffusion properties
- 5 Conclusions

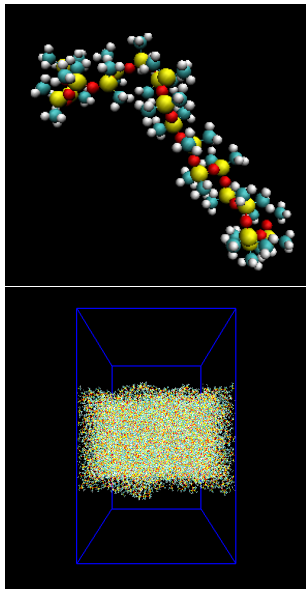


- Siloxanes are commercially valuable silicon polymers
 - ▶ Adhesives and sealants
 - ▶ Lubricants
 - ▶ Heat transfer fluids
- Study the interaction of water and siloxanes – in particular polydimethylsiloxane (PDMS)
 - ▶ Diffusion of water through bulk PDMS
 - ▶ Dynamics of wetting a PDMS surface
 - ★ Surface tensions both PDMS and water
 - ★ Not widely available for PDMS; many studies for water



Simulation details

PDMS surface tension

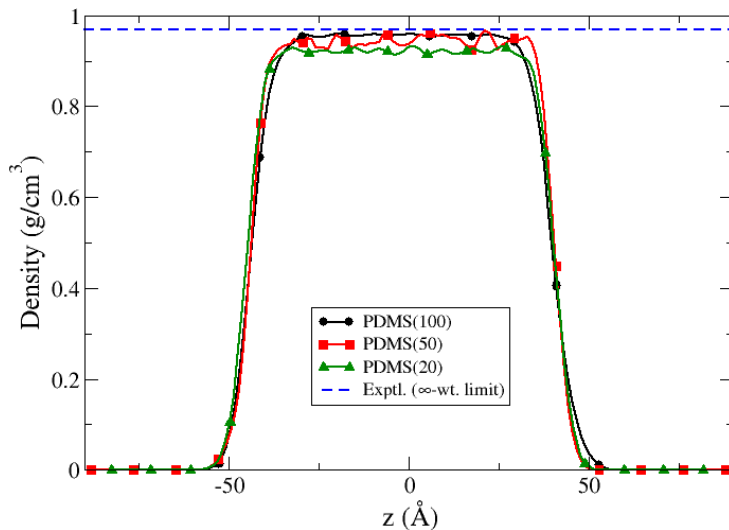


- 500 20-mers (103500 atoms) or 100 100-mers (107000 atoms)
- Simulation box: $100 \text{ \AA} \times 100 \text{ \AA} \times 170 \text{ \AA}$
- Buckingham (Exp-6) potential of Smith et al. (2004)
 - ▶ 10 Å cutoff
 - ▶ PPPM Ewald for electrostatics
- All-atom NVT simulation at 300 K using RESPA with 10^6 time steps of 2 fs
- Total time: 2 ns



Simulation results: liquid-vapor density

Density of longer chains rapidly approaches bulk limit



Simulation results: surface tension

Density of longer chains rapidly approaches bulk limit

Surface tension includes results of simulation plus long-range correction:

$$\gamma = \gamma_p + \gamma_{tail}$$

where

$$\begin{aligned}\gamma_p &= \frac{L_z}{2} \left(p_z - \left(\frac{p_x + p_y}{2} \right) \right) \\ \gamma_{tail} &= \frac{\pi}{2} \int_{-\infty}^{\infty} \int_{-1}^1 \int_{r_c}^{\infty} r^3 \frac{dU(r)}{dr} g(r) (1 - 3s^2) \times \\ &\quad \left(\rho(z) \rho(z - sr) - (\rho_G(z))^2 \right) dr ds dz\end{aligned}$$



Simulation results: surface tension of PDMS

Surface tension increases with molecular weight for -CH₃ PDMS

Chain type	Surface tension, mN/m		
	γ_p	γ_{tail}	$\gamma = \gamma_p + \gamma_{tail}$
20-mer -CH ₃	15.0	5.5	20.5
50-mer -CH ₃	18.0	5.7	23.7
100-mer -CH ₃	18.5	5.4	23.9
20-mer -OH	16.4	5.2	21.6

Exptl. value, infinite molecular weight:

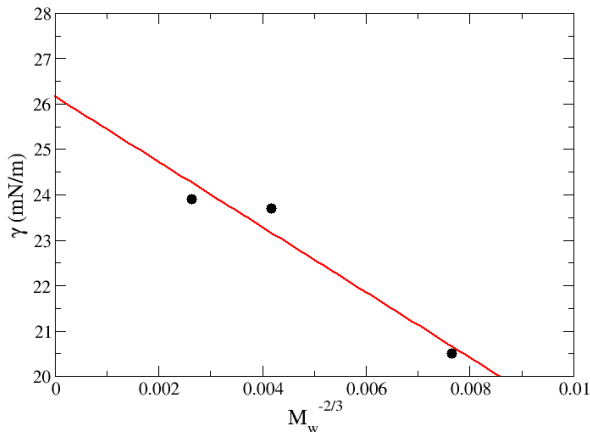
$$\gamma = 21.0 \text{ mN/m.}$$

[Dee and Sauer, *Macromolecules*, **26**, 2771 (1993).]



Simulation results: surface tension of PDMS

–2/3-power law dependence

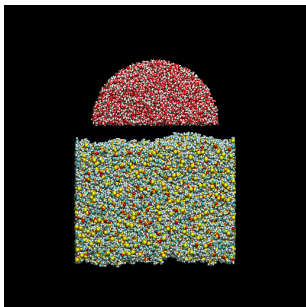


$$\gamma(M_w = \infty) = 26.2 \text{ mN/m (simulation)}$$
$$\gamma(M_w = \infty) = 21.0 \text{ mN/m (experimental)}$$

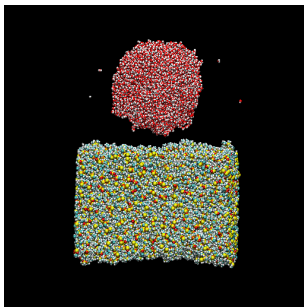


Surface tension of PDMS-water interface

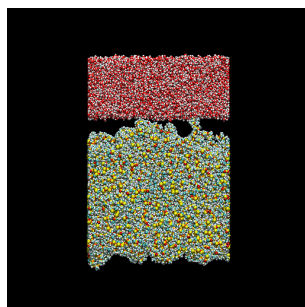
Three different geometries tested



Hemispherical drop



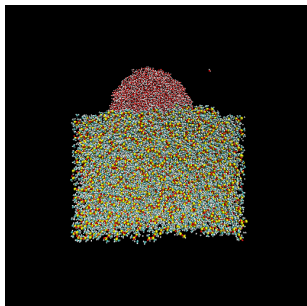
Spherical drop



Slab

Surface tension of PDMS-water interface

Three different geometries tested



Hemispherical drop

- 60 Å radius hemispherical drop (~ 10000 molecules) placed on top of PDMS slab
- 2 ns simulation in *NVT* ensemble at 300 K
- Measure contact angle as a function of PDMS chain length

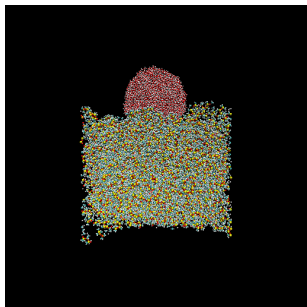
Chain length	Contact angle (deg.)
20-mer -CH ₃	108 ± 10
50-mer -CH ₃	108 ± 10
100-mer -CH ₃	108 ± 10

- Exptl. contact angle: 105-120 degrees



Surface tension of PDMS-water interface

Three different geometries tested



Spherical drop

- 50 Å radius spherical drop (~ 10000 molecules) placed on top of PDMS slab
- 2 ns simulation in *NVT* ensemble at 300 K
- Measure contact angle as a function of PDMS chain length
- Use Young's equation to determine surface tension of PDMS-water interface:

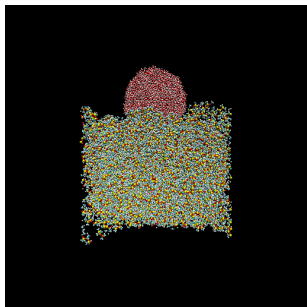
$$\gamma_{PW} = \gamma_P + \gamma_W \cos \theta$$

where $\gamma_W = 61$ mN/m is the surface tension of SPC/E water



Surface tension of PDMS-water interface

Three different geometries tested



Spherical drop

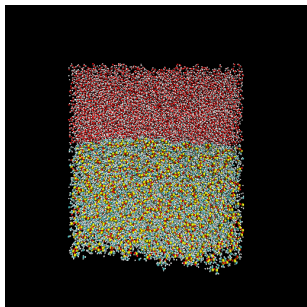
Chain length	Contact angle (deg.)
20-mer -CH ₃	108 ± 10
50-mer -CH ₃	108 ± 10
100-mer -CH ₃	108 ± 10

- Using Young's equation, we find $\gamma_{PW} = 39 \pm 8$ mN/m.
- Exptl. value ~ 40 mN/m



Surface tension of PDMS-water interface

Three different geometries tested



Slab

- 50-Å thick slab of water (~ 20000 molecules) placed on top of PDMS slab
- 2 ns simulation in NVT ensemble at 300 K
- Measure total surface tension
- γ for PDMS-water surface tension given by overall γ , less water-vapor, PDMS-vapor interfaces:

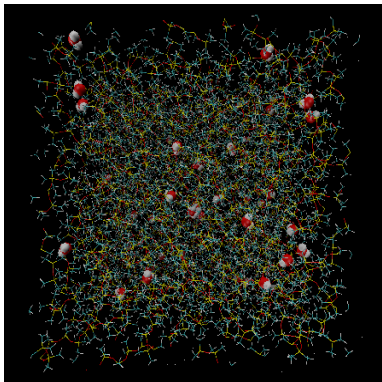
$$\gamma_{PW} = \gamma_{tot} - (\gamma_P + \gamma_W)$$

- For 100-mer -CH₃ PDMS,
 $\gamma_{PW} = 127 - (24 + 61) = 42\text{mN/m}$



Diffusion of water through PDMS

Simulation methodology



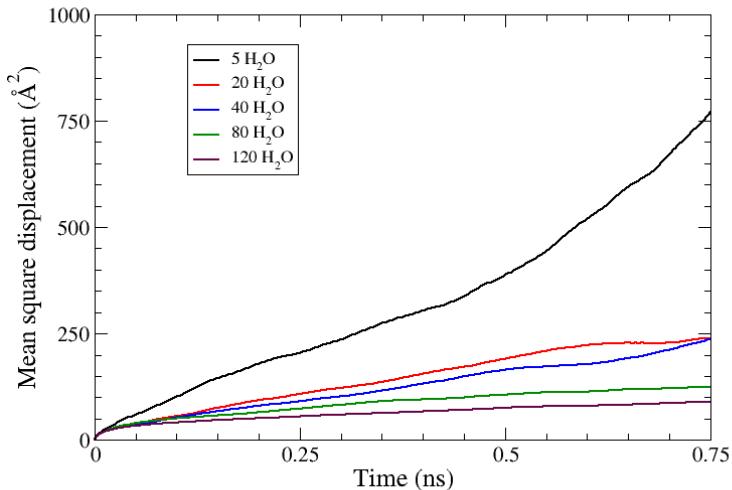
- 5-120 water molecules placed in bulk sample of 100 20-mers or 20 100-mers ($-\text{CH}_3$ and $-\text{OH}$ terminated)
- > 2 ns MD simulations in the NVT at $T = 300$ K.
- Measure rms displacement of waters as a function of time
- Goals:
 - ▶ Determine diffusion constant
 - ▶ Observe phase separation



Diffusion of water through PDMS

Strong concentration effects on diffusion

Diffusion of water through 100-mer CH_3 -terminated PDMS



Diffusion of water through PDMS

Concentration, terminal-group effects on diffusion

Chain type	Diffusion constant, D ($\text{\AA}^2 \text{ps}^{-1}$)	
	5 H ₂ O	20 H ₂ O
20-mer -CH ₃	0.2742	0.0531
20-mer -OH	0.1524	0.0513
100-mer -CH ₃	0.1490	0.0518
100-mer -OH	0.1197	0.0808

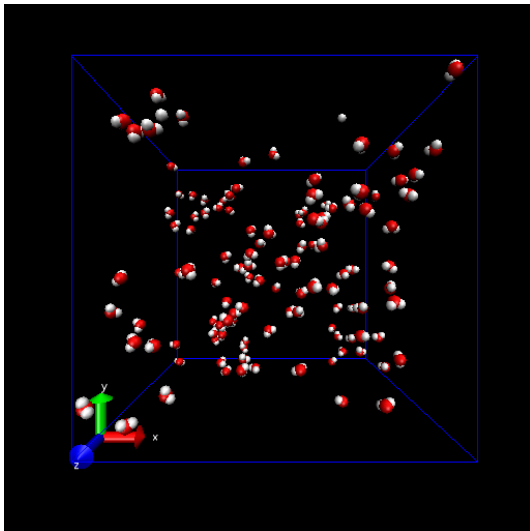
Exptl. value, bulk H₂O: $0.2272 \text{ \AA}^2 \text{ps}^{-1}$

Phase separation observed for concentrations > 20 water molecules



Diffusion of water through PDMS

Water molecules quickly form clusters



Conclusions

- PDMS, water surface tensions underestimated by surface tension
- Diffusion constants reasonable
 - ▶ Some termination-group effects
 - ▶ Phase separation occurs for intermediate to high concentrations of water
- Contact angle of PDMS-water interface
 - ▶ Different geometries yield almost identical contact angles
 - ▶ Results in reasonable agreement with experimental data
- PDMS-water interfacial tension
 - ▶ Different approaches to calculating PDMS-water interfacial tension yield comparable results
 - ▶ Also in good agreement with experimental observations



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- Scott W. Sides (prior PDMS studies)
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