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WEBINAR

Molecular Simulation of Fluids: The SAFT Coarse Graining Technique

Professor Erich Müller
Imperial College London

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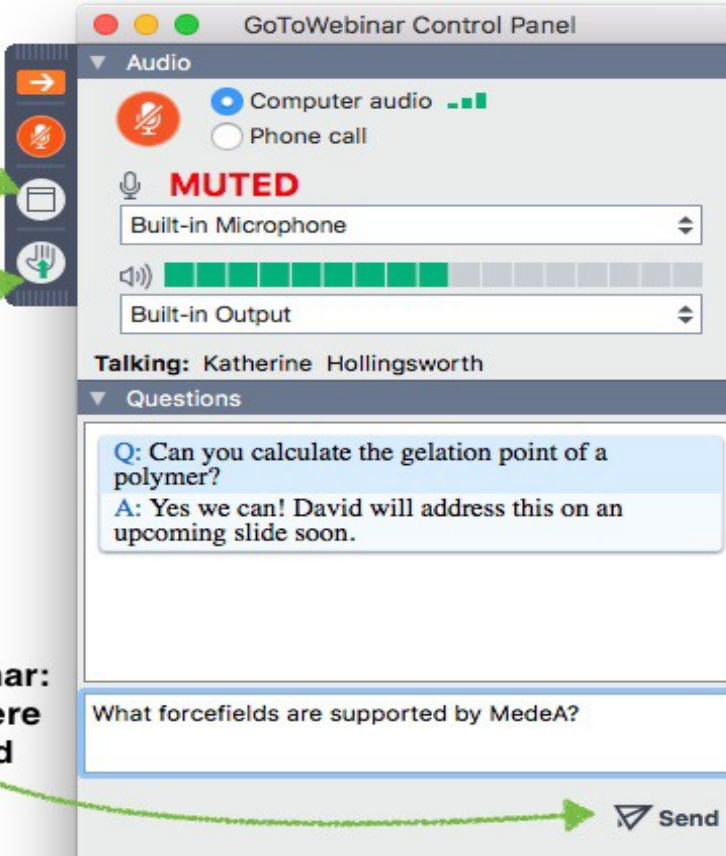
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Use the raise hand icon to bring attention to your question



The screenshot shows the GoToWebinar Control Panel interface. On the left, a vertical toolbar contains several icons: a right-pointing arrow, a microphone with a slash, a window icon, and a hand with the index finger raised. Green arrows point from the text 'full screen' to the window icon, and from 'during discussion:' to the hand icon. The main panel is titled 'GoToWebinar Control Panel' and has a 'MUTED' status. It includes sections for 'Audio' (with 'Computer audio' selected and 'Phone call' unselected), 'Talking: Katherine Hollingsworth', and 'Questions'. The 'Questions' section shows a question: 'Q: Can you calculate the gelation point of a polymer?' and an answer: 'A: Yes we can! David will address this on an upcoming slide soon.' Below this is a text input field containing 'What forcefields are supported by MedeA?' and a 'Send' button with a paper plane icon. A green arrow points from the text 'any time during webinar: type your question here and then press Send' to the 'Send' button.

full screen

during discussion:

any time during webinar:
type your question here
and then press Send

Webinar Speakers



Professor Erich Müller

Imperial College London

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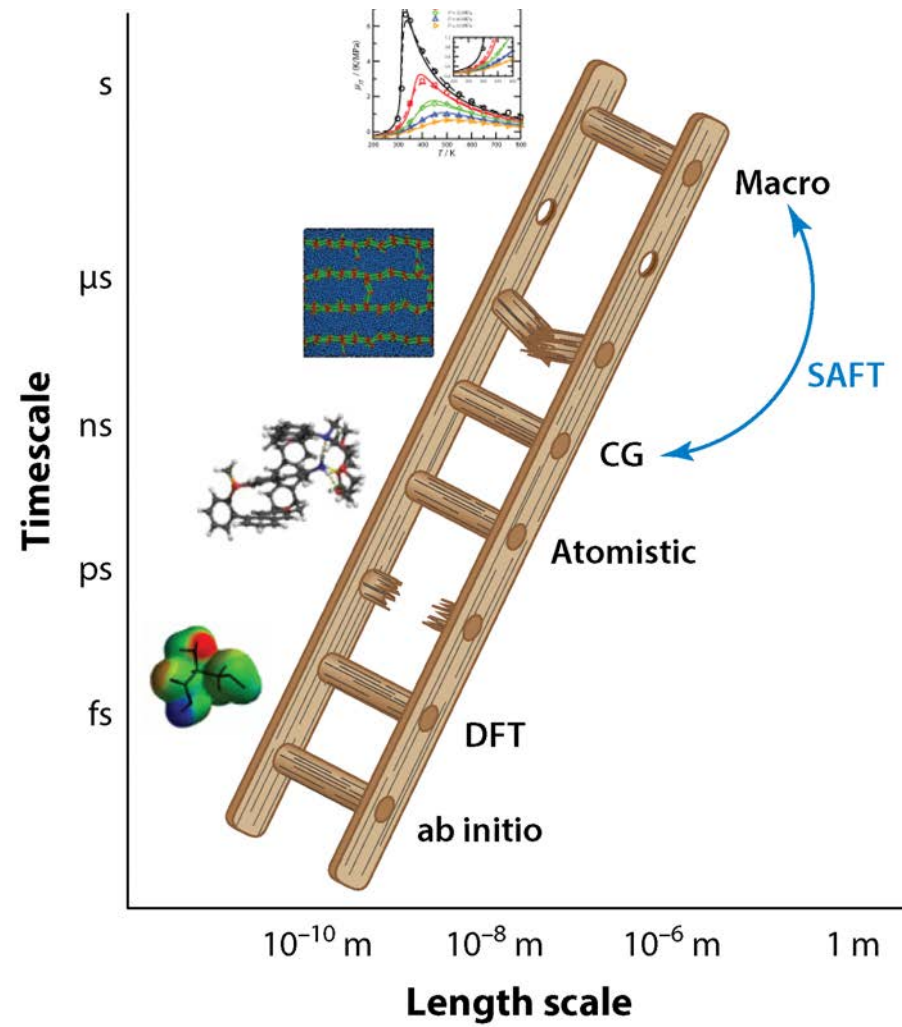
myiannourakou@materialsdesign.com

Molecular simulation of fluids:
The SAFT coarse graining technique

Erich A. Müller

Department of Chemical Engineering

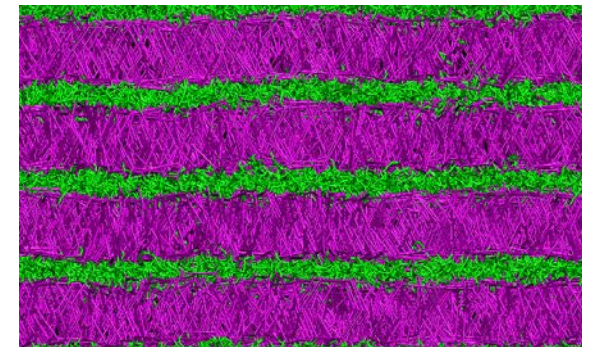
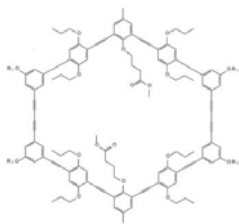
Imperial College London, U.K.



The molecular simulation ladder

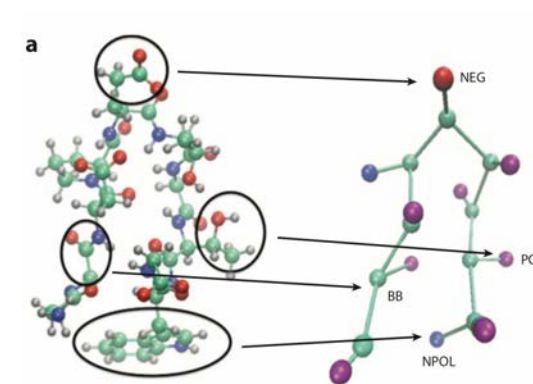
Coarse Graining

- By removing the detail in the molecular model, significant gains in both the speed of the calculations and the accessible size of the system can be procured.
- Essential if large molecules and/or large time scales are needed e.g. polymer dynamics, multicomponent mixtures, self-assembly of soft matter,
- Risk of GIGO (garbage in, garbage out) as no universally accepted procedure or methodology exists.
- Integrating out degrees of freedom (bottom-up approach) inevitably results in a loss of information with problems of transferability



Linking scales: Bottom-up approaches

- Iterative-Boltzmann technique: Attempts to match the radial distribution function of an atomistically-detailed model [1]
- Force matching : Attempts to equate the effective intermolecular forces (and angles) between atomistic detailed models and larger force centres. [2]
- Relative Entropy scaling : employs a measure of the entropy to obtain matches between scales [3]
- Machine learning : [4].



Coarse-grained model of a peptide with at least one CG site per amino acid residue. From [2]

[1] T. C. Moore, C. R. Iacovella, and C. McCabe, "Derivation of coarse-grained potentials via multistate iterative Boltzmann inversion," *J. Chem. Phys.* 140, 224104 (2014)

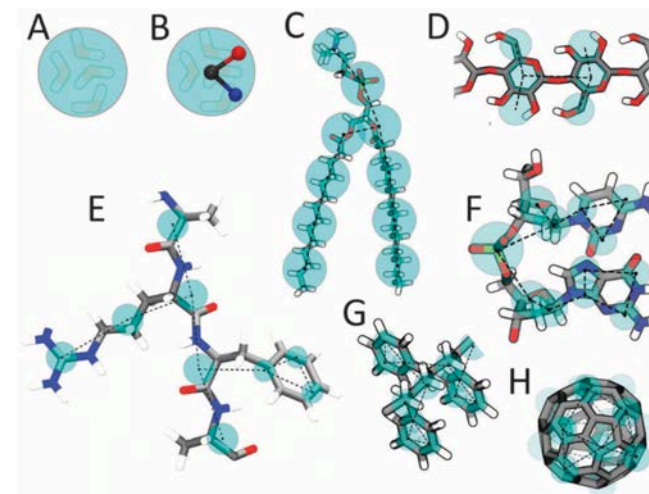
[2] M. G. Saunders and G. A. Voth, "Coarse-graining methods for computational biology.," *Annu. Rev. Biophys.*, 42, 73–93, (2013).

[3] M. S. Shell, "Coarse-graining with the relative entropy," in *Advances in Chemical Physics* (John Wiley & Sons, Inc., 2016), pp. 395–441

[4] P. Gao, X. Yang, and A. M. Tartakovsky, "Learning Coarse-Grained Potentials for Binary Fluids.," *J. Chem. Inf. Model.*, 60(8), 3731–3745, (2020)

Linking scales: Top-down approaches

- MARTINI: guides the parameter estimation by matching the experimental water/octanol partition coefficient. Employs a fixed set of equal-sized beads [1]
- SAFT: maps the size, energy and range of the beads to the expected macroscopic volumetric properties of the fluids [2]
- DPD: requires ad-hoc fitting of parameters, e.g. to infinite dilution coefficients [3]

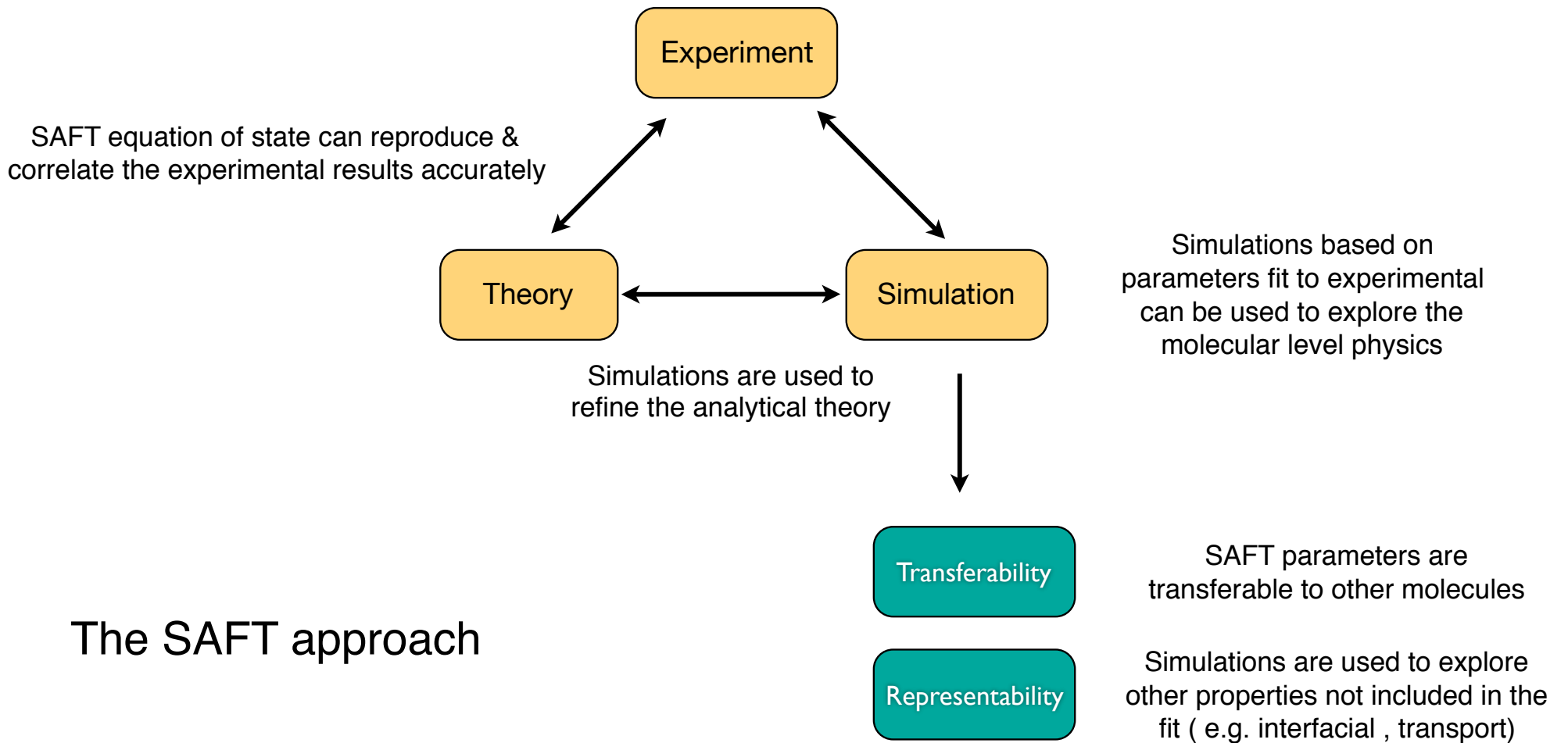


MARTINI mapping of selected molecules. From [1]

[1] S. J. Marrink and D. P. Tieleman, "Perspective on the Martini model," *Chem. Soc. Rev.*, 42(16) 6801–22, 2013.

[2] E. A. Müller and G. Jackson, "Force-Field Parameters from the SAFT- γ Equation of State for Use in Coarse-Grained Molecular Simulations," *Annu. Rev. Chem. Biomol. Eng.*, 5, 405–427, 2014.

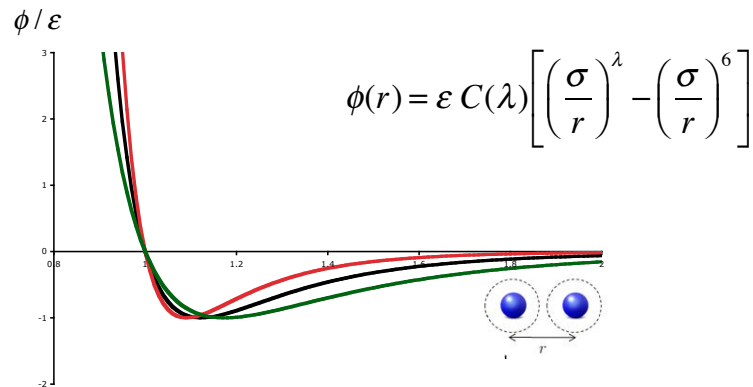
[3] A. Vishnyakov, M.-T. Lee, and A. V. Neimark, "Prediction of the Critical Micelle Concentration of Nonionic Surfactants by Dissipative Particle Dynamics Simulations," *J. Phys. Chem. Lett.*, 4(5) 797–802, 2013.



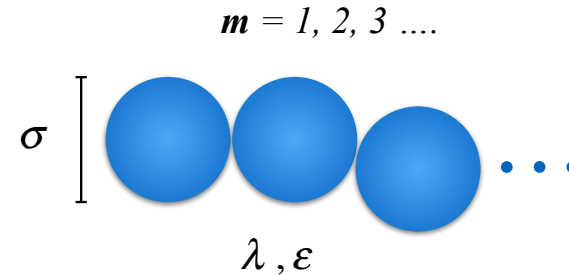
The SAFT approach

SAFT- γ -Mie : Equation of state

Mie potential



4 physical parameters

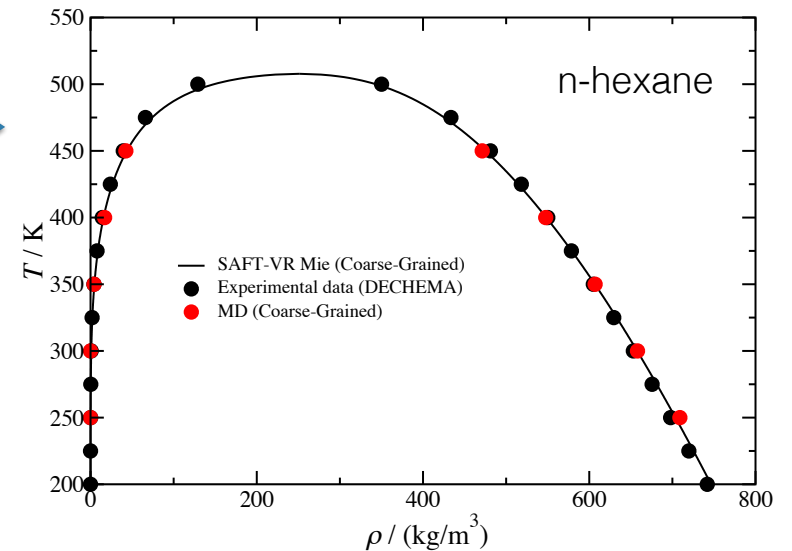


SAFT- γ formulation is unique as it directly relates a macroscopic equation of state with an underlying force field

third order T expansion

$$a = a^{IDEAL} + a^{MONO} + a^{CHAIN}$$

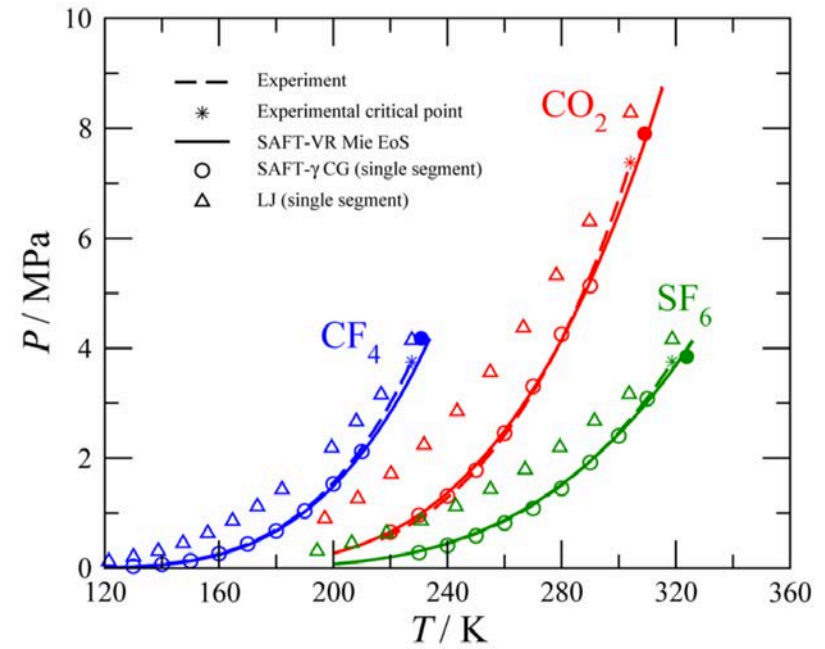
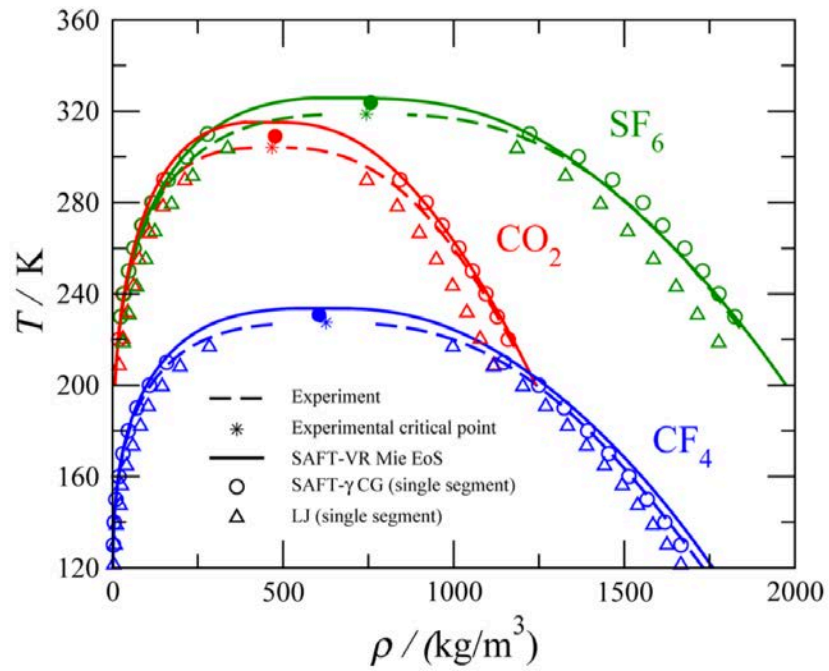
$$a^{MONO} = m \left[a^{HS} + \beta a_1 + \beta^2 a_2 + \beta^3 a_3 \right]$$



How to obtain the parameters for the EoS / forcefield?

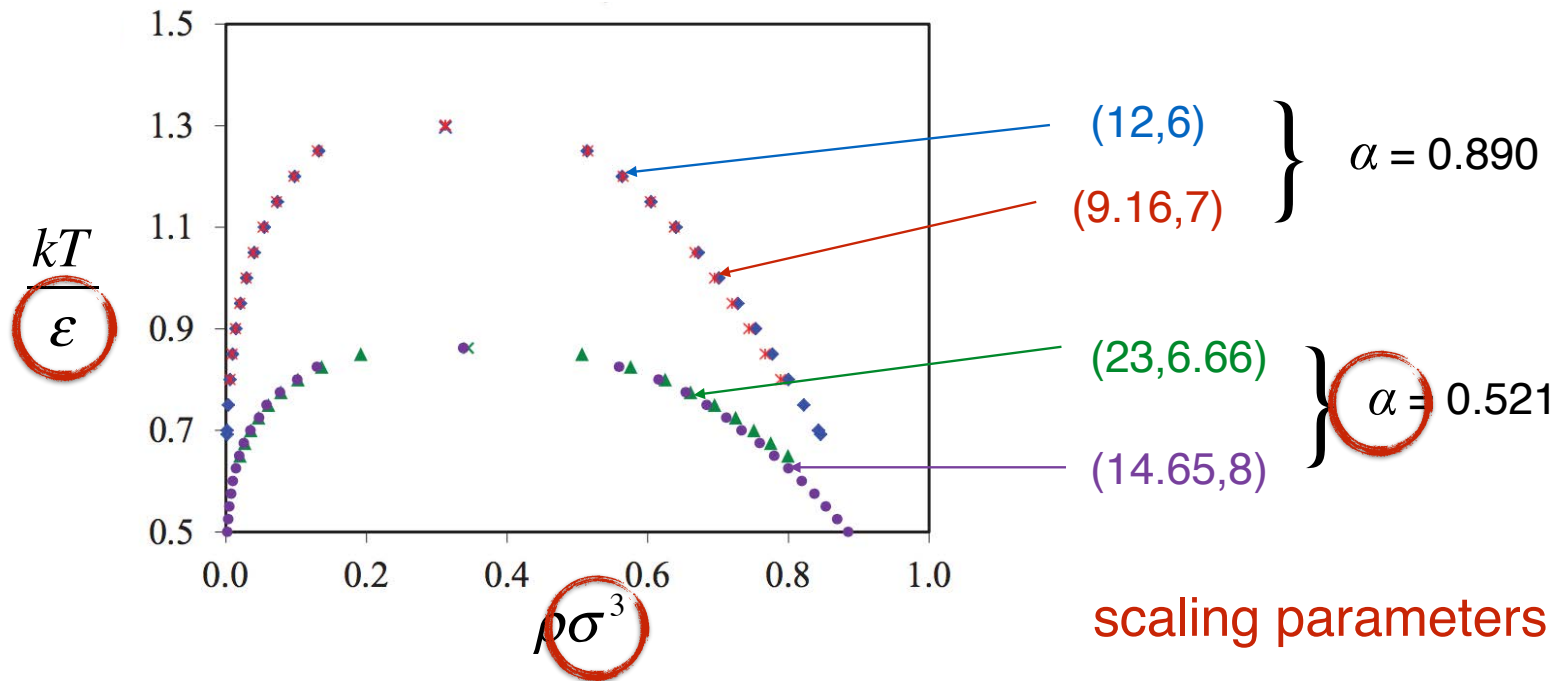
- Fit the EoS to experimental data. Usually information on the vapor-liquid densities and vapour pressures, although we can incorporate other properties (excess volumes, enthalpies, etc.)
- Formulate in terms of corresponding states, hence only need a few pieces of key critical data

Fitting to experimental VLE data



Conformality of the Mie (m,n) potential

$$u(r) = C\epsilon \left[\left(\frac{\sigma}{r} \right)^m - \left(\frac{\sigma}{r} \right)^n \right]$$

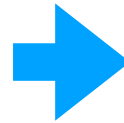
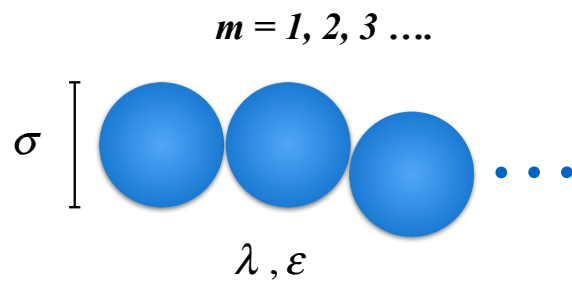


van der Waals parameter

$$\alpha = \frac{1}{\epsilon\sigma^3} \int_{\sigma}^{\infty} u(r)r^2 dr = \frac{\lambda}{3(\lambda-3)} \left(\frac{\lambda}{6} \right)^{6/\lambda-6}$$

for the $(\lambda,6)$ potential

Three (four) parameter corresponding states



$$m = f(\text{molecular geometry})$$

$$\lambda = f(\alpha) = f(\omega)$$

$$\epsilon = f(T^*) = f(T_c)$$

$$\sigma^3 = f(\rho^*) = f(\text{density})$$

Since SAFT is an equation of state it can be described in terms of corresponding states, hence **the molecular parameters can be obtained from critical properties.**

SAFT- γ Mie, bottled

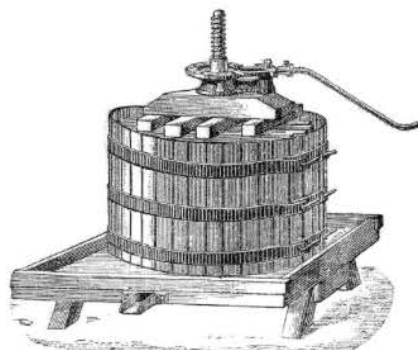
The SAFT- γ Mie force field is a powerful approach to coarse-grained molecular dynamics: the parameters for the model of a substance are computed from its thermophysical properties, removing the need for tuning models to atomistic simulations. The theory also provides a corresponding equation of state. Here we make the power accessible to anyone.



The bottle

"The bottle" is our collection of 6000+ molecules for which we have already computed the force field parameters. You can search by CAS number, name, or chemical formula.

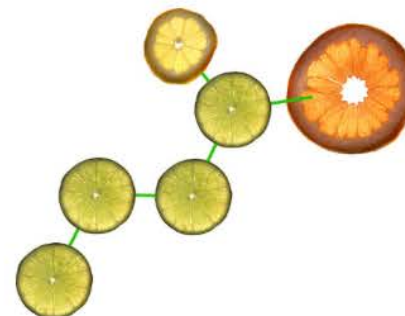
[Search the bottle](#)



Press your own

You can create your own model by providing critical properties for the compound you want to study. The M&M correlation instantly gives you the force field parameters.

[Run your own compound](#)



Heteronuclear models

We're hoping to extend this such that you can draw the molecule you want. This feature is still work in progress, for now we just have a demo of the interface that gives you back the SMILES.

[Build with the blocks](#)

The theory

The SAFT- γ Mie force field forms part of an established top-down coarse graining methodology that enables molecular dynamics simulations of a wide range of components in the vapor and liquid states [1]

References

[1] Erich A. Müller and Ge...

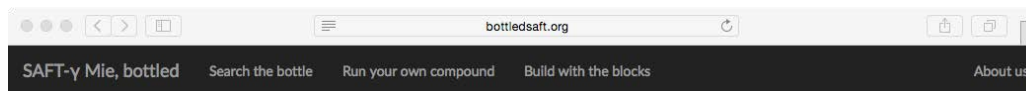
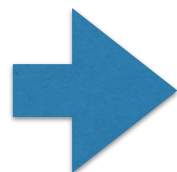
[2] Andrés Mejía, Carmelo Herdes and Brian P...

www.bottledsaft.org

Example

input

T_c	374.1 K
ω	0.3268
$\rho_{0.7}$	13044 mol/m ³



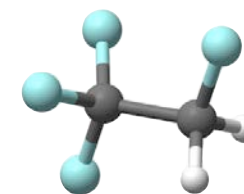
Final results

	1,1,1,2-tetrafluoroethane
CAS	811-97-2
formula	C2H2F4
Molar mass (g/mol)	102.03
Liquid density at T = 0.7 T _c (mol/m ³)	13107.37

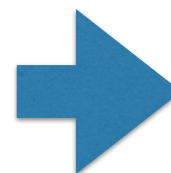
The parameters available are shown below. In this case the correlation gives sensible results for several segment numbers. Choosing the best one is a tradeoff. There should be roughly three atoms per segment, not counting hydrogen. The topology (e.g. the length to breadth aspect ratio) of the coarse-grained model should match the topology of the original molecule.

Clicking "raaSAFT script" gives you a Python implementation for simulating this with raaSAFT, our SAFT- γ Mie framework for running MD simulations with HOOMD-blue (well-supported) or GROMACS (new feature). See the [raaSAFT webpage](#) for how to install and use raaSAFT.

R134a



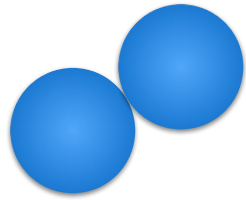
# segments	2
lambda_r	21.638
lambda_a	6
epsilon (K)	291.47
sigma (m)	3.6710e-10



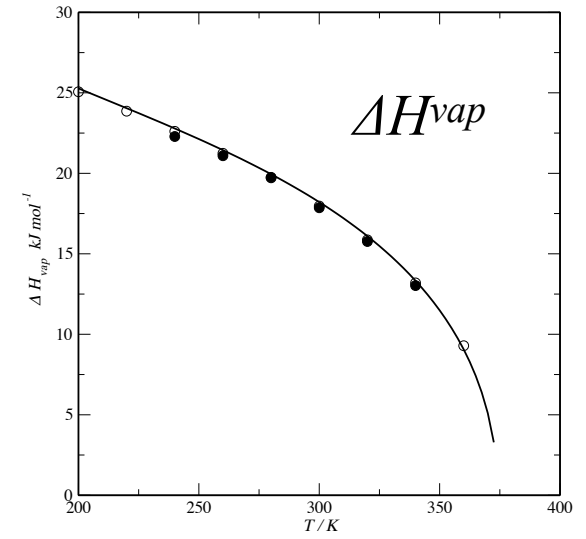
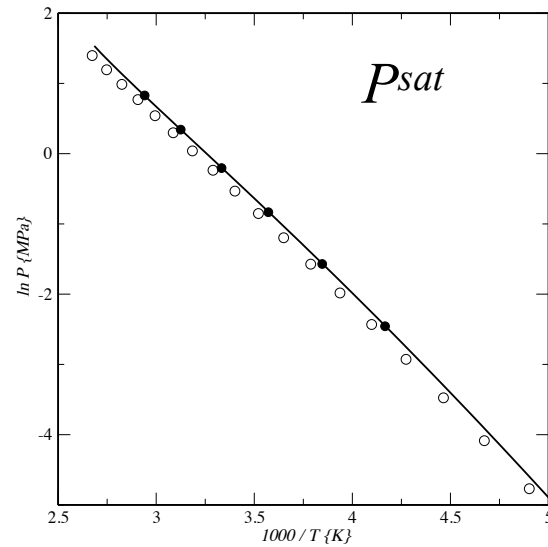
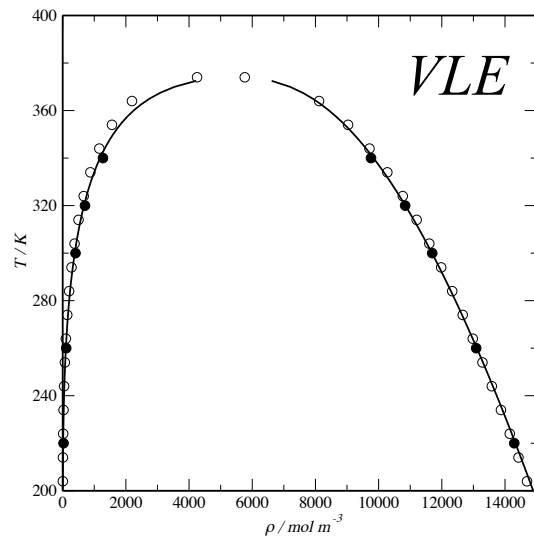
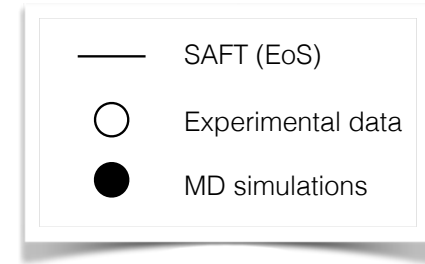
SAFT parameters

m	2
λ	21.6
ϵ (K)	291.5
σ (Å)	3.671

R134a



m	2
λ	21.6
ε (K)	291.5
σ (Å)	3.671



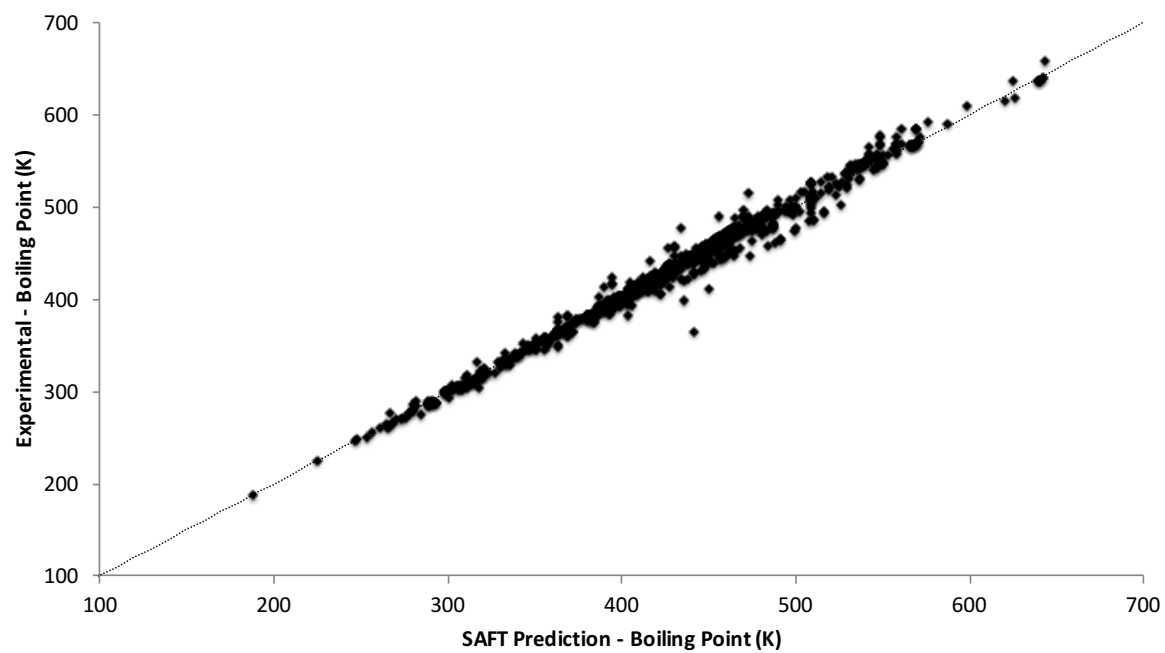
Parameters for 6000+ well-defined chemicals

	m	λ	ϵ/k_B [K]	σ (nm)
<i>n-alkanes</i>				
methane	1	16.39	170.75	0.3752
ethane	1	27.30	330.25	0.4349
propane	1	34.29	426.08	0.4871
butane	2	13.29	256.36	0.3961
pentane	2	16.06	317.50	0.4248
hexane	2	19.57	376.35	0.4508
heptane	2	23.81	436.13	0.4766
octane	3	16.14	333.70	0.4227
nonane	3	18.31	374.21	0.4406
decane	3	20.92	415.19	0.4584
undecane	4	16.84	348.90	0.4216
dodecane	4	18.41	378.56	0.4351
tetradecane	5	17.66	363.06	0.4183
hexadecane	5	21.20	418.13	0.4432
octadecane	6	19.53	393.74	0.4262
eicosane	6	24.70	453.10	0.4487
<i>aromatics</i>				
benzene	2	14.23	353.93	0.3978
toluene	2	16.95	411.87	0.4266
ethylbenzene	3	12.80	309.69	0.3837
naphthalene	3	12.84	376.50	0.3932
<i>light gases</i>				
nitrogen	1	20.02	122.85	0.3653
carbon dioxide	2	14.65	194.94	0.2848

oxygen	1	17.93	144.02	0.1295
hydrogen sulphide	1	27.38	403.93	0.3801
sulphur dioxide	2	16.06	291.10	0.3091
carbon monoxide	1	21.49	132.83	0.3687
helium	1	14.84	4.44	0.3353
argon	1	14.85	132.04	0.3414
<i>heterocyclics</i>				
pyridine	2	15.52	410.46	0.3899
pyrrolidine	2	19.74	426.12	0.3914
thiolane	2	13.83	391.03	0.4012
thiophene	2	13.58	354.34	0.3832
<i>branched and cyclo-alkanes</i>				
isopentane	2	14.92	298.38	0.4236
isobutane	2	12.94	241.57	0.3974
cyclopentane	2	13.52	312.00	0.3992
cyclohexane	2	14.05	345.94	0.4234
<i>unsaturates</i>				
ethylene	1	25.62	299.49	0.4180
propylene	1	33.65	417.60	0.4721
1-pentene	2	17.50	328.86	0.4183
1-decene	3	18.93	394.49	0.4516
<i>solvents / others</i>				
water	1	8.395	378.87	0.2915
tetrahydrofuran	2	14.85	348.92	0.3840

- Mejía, A., et al. (2014). Force Fields for Coarse-Grained Molecular Simulations from a Corresponding States Correlation. *Ind. Eng. Chem. Res.*, 53(10), 4131–4141.

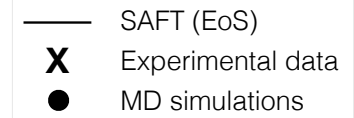
A quick check



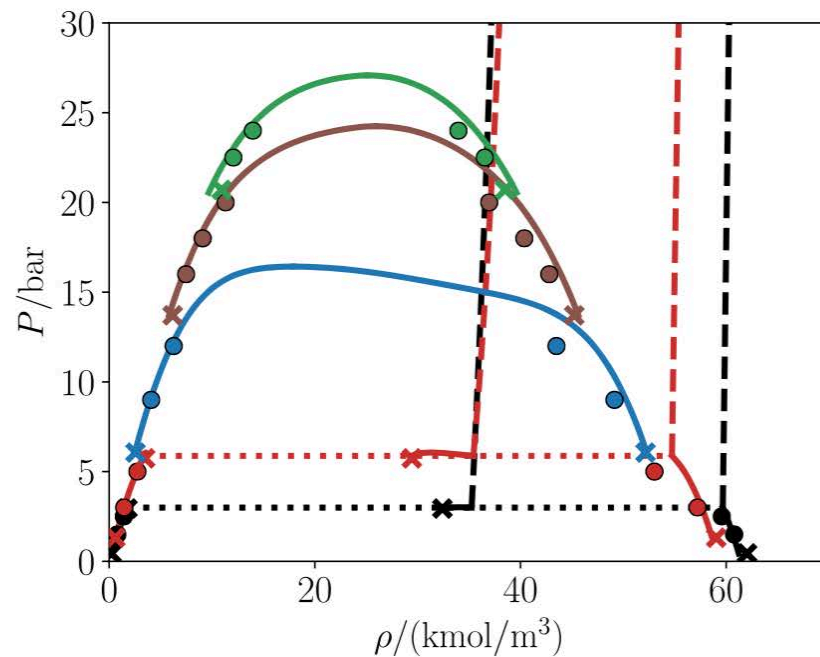
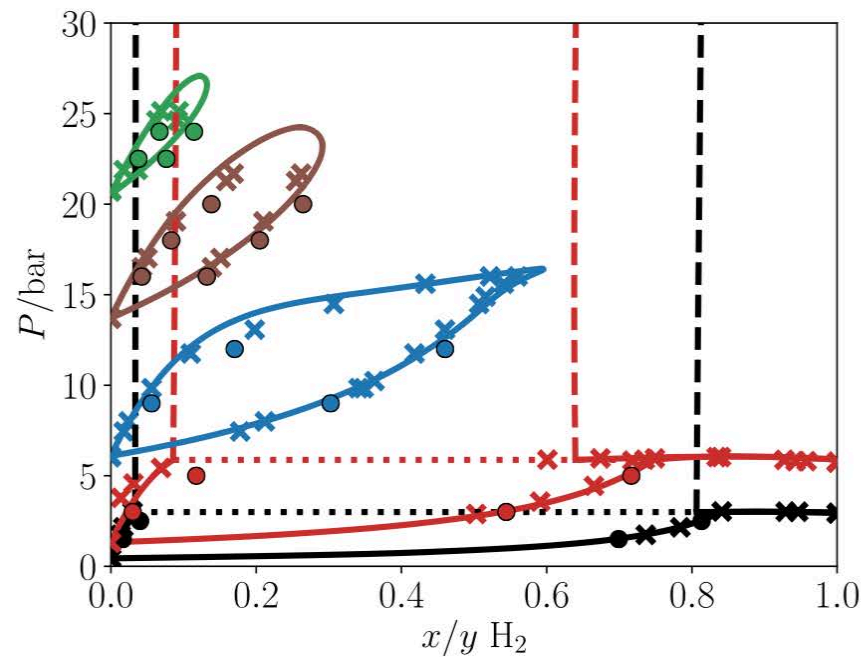
Normal boiling point prediction. AAD 1.15 %

Cryogenic fluids

Adding the Feynman-Hibbs correction



Hydrogen-Neon mixtures at 25 K, 28 K, 35 K, 40 K and 43 K

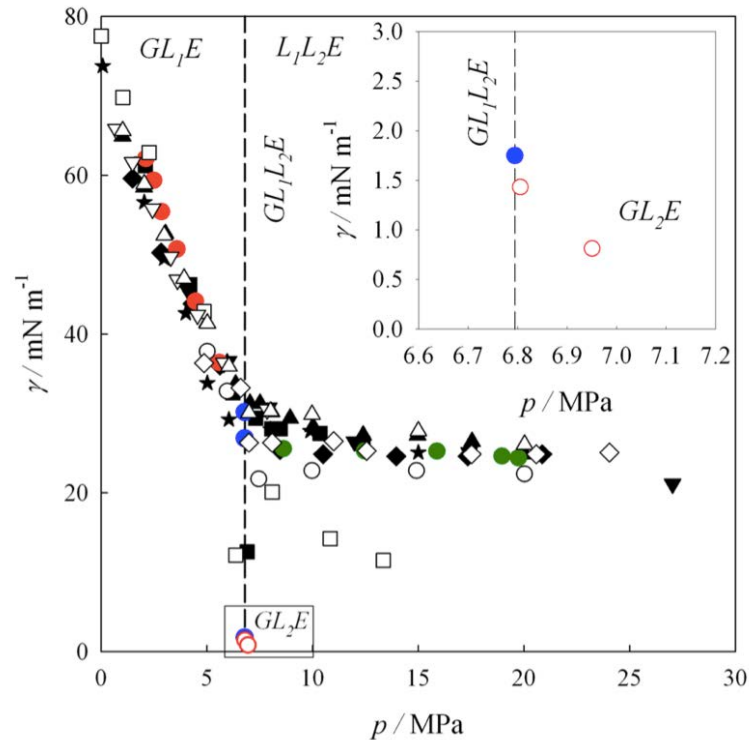


How about transport / interfacial properties?

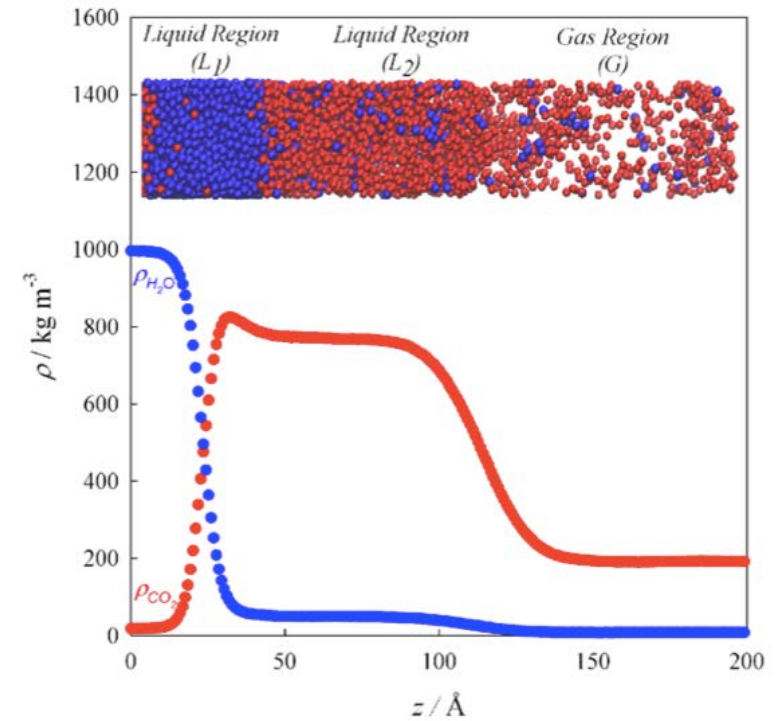
- Although only volumetric properties are employed for parametrization, the procedure is basically a very broad fit of the effective Helmholtz free energy landscape.
- This seems to be sufficient to produce robust optimised force field parameters which can be employed to explore other properties (e.g. interfacial, transport, etc..)

Resolving discrepancies in experimental data

High pressure CO₂-water interfacial tension

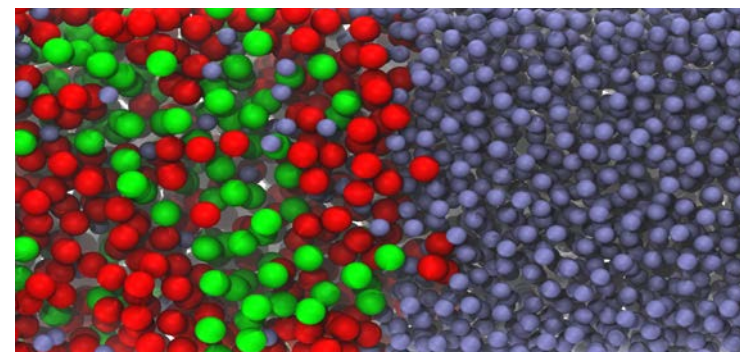
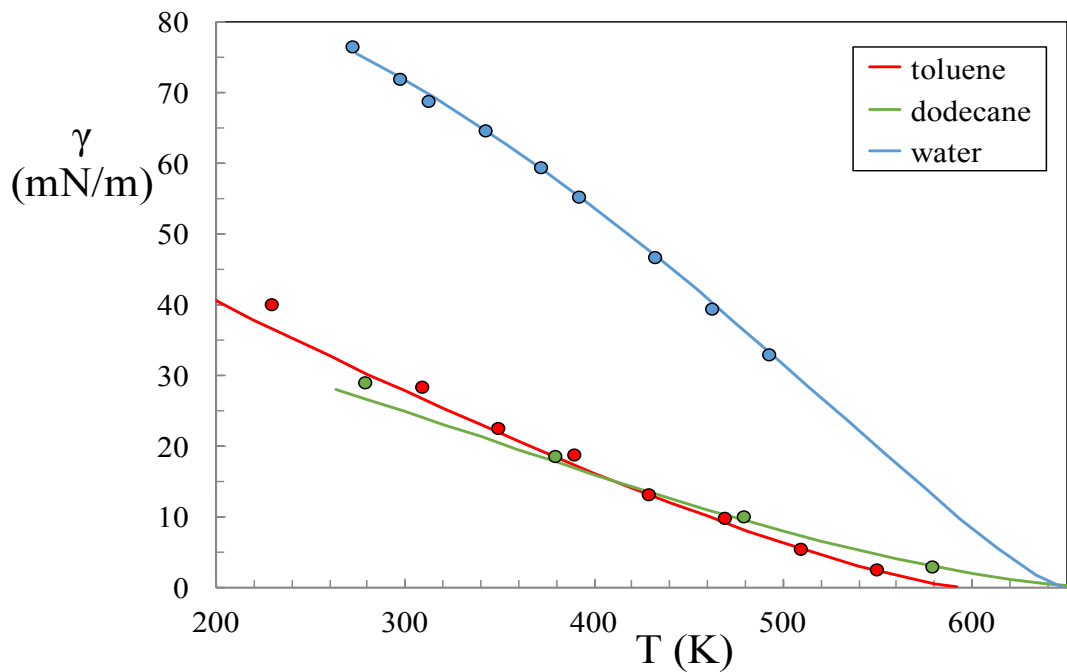


Experimental data (B&W) show extreme scatter



Large scale CG simulations demonstrate the existence of a liquid-liquid-vapor region (3 phases) which explain the experimental findings

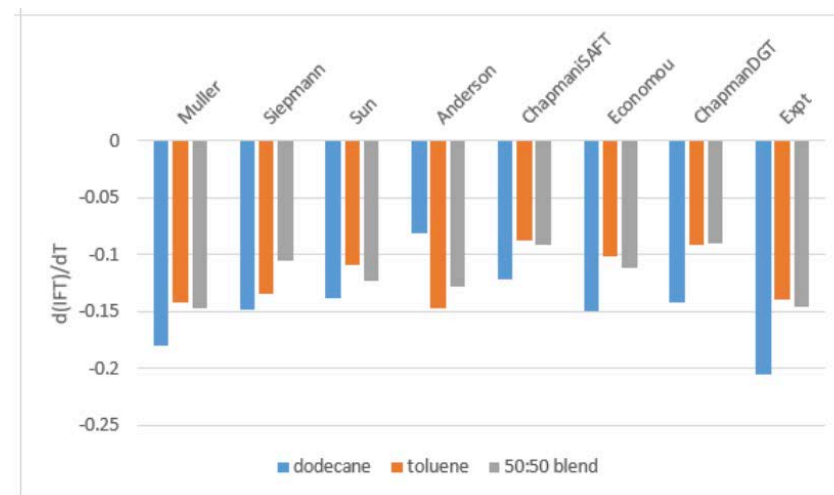
Oil/water interfacial tension



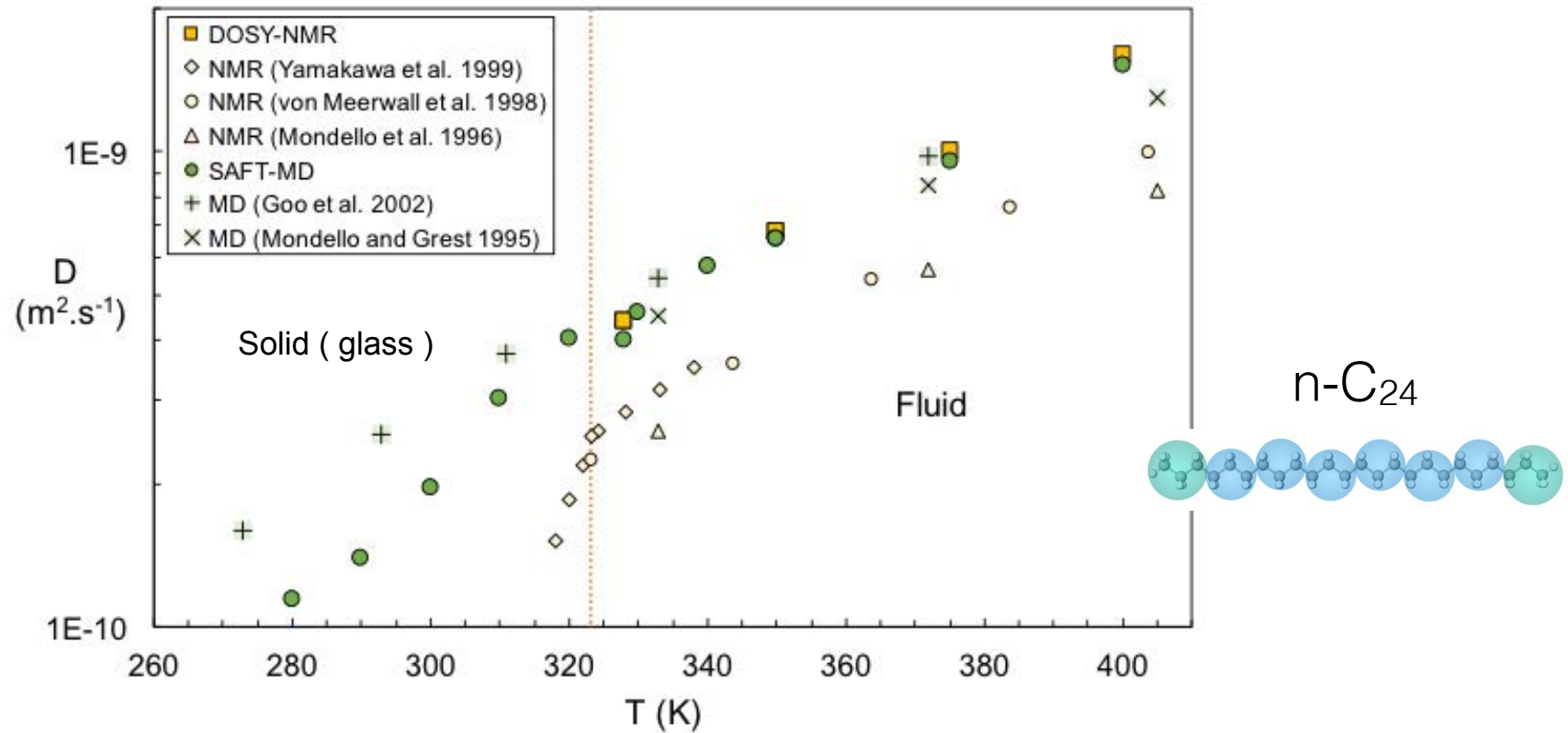
P = 250 psi
T = 110-170 C

The MOST ACCURATE modelling technique to determine mixture oil/water IFT at high P and T
AIChE 2016 – Industrial fluid property challenge

<http://fluidproperties.org/results-ninth-challenge>

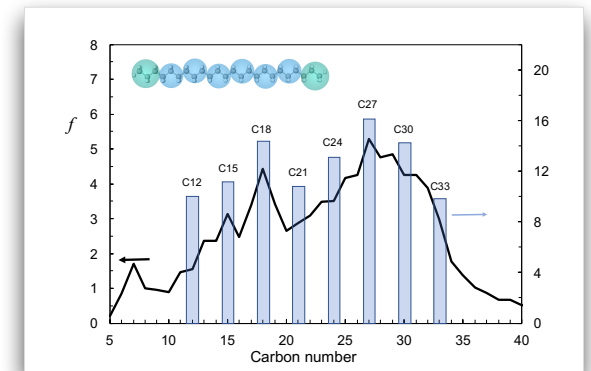
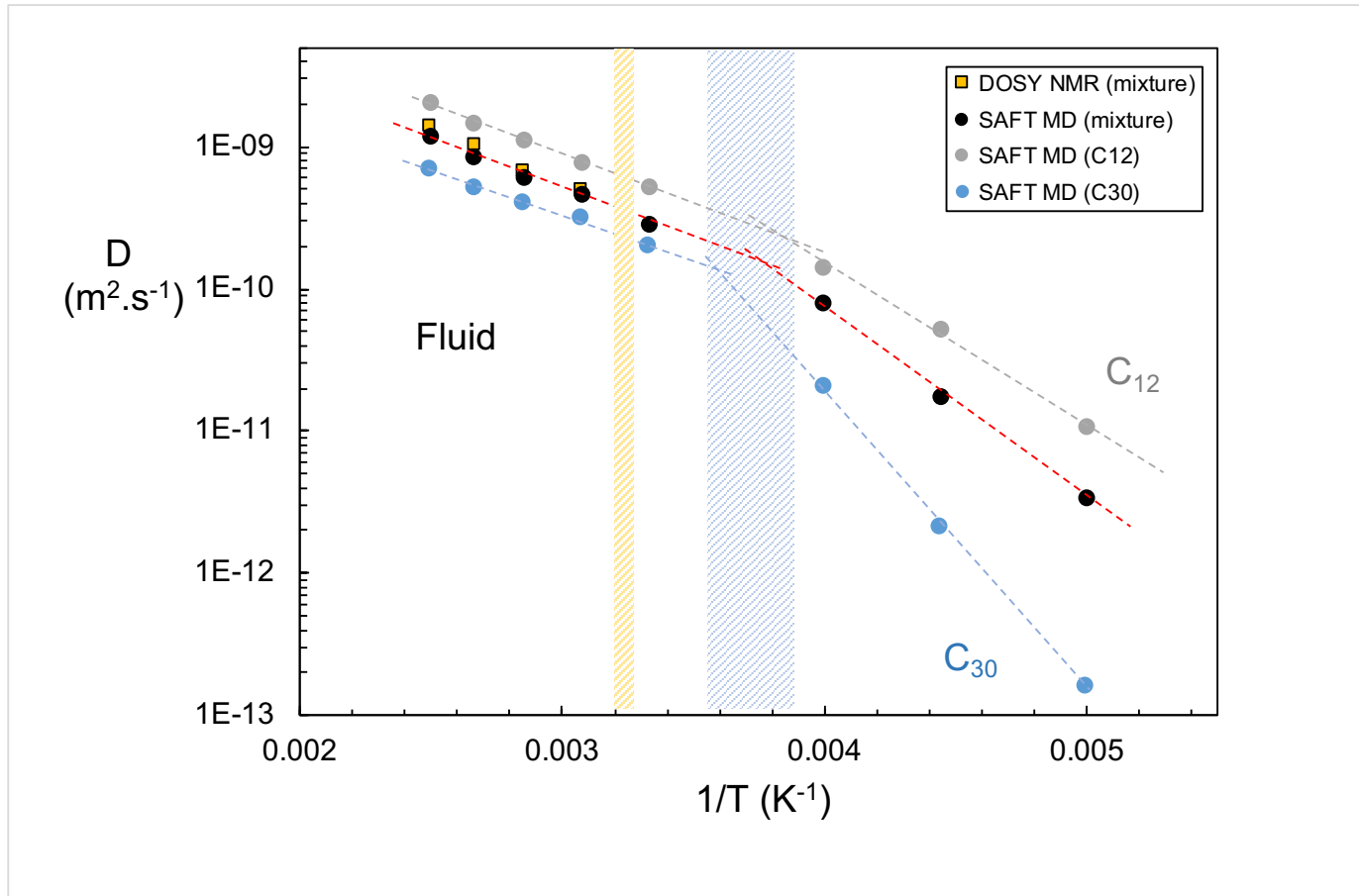


Self-diffusion coefficients



DOSY-NMR : Diffusion ordered spectroscopy nuclear magnetic resonance

Wax appearance temperature



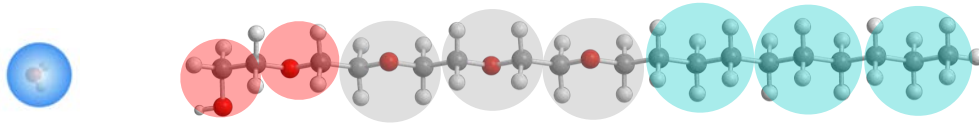
8 component alkane mixture fitted to HPLC data of a real Malaysian crude

How about complex molecules?

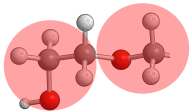
- Fitting parameters are usually done on a per-molecule basis with homogeneous molecules.
- There is no reason why one could not invoke a **group-contribution** approach.

Ethoxylated Alkyl surfactants

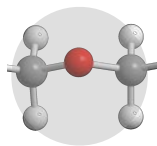
(transferring groups from smaller molecules)



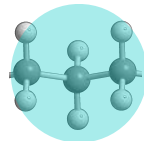
- Break down the molecule into similar moieties
- heteronuclear segment parameters estimated from thermodynamic properties (density, vapour pressure) of simple molecules



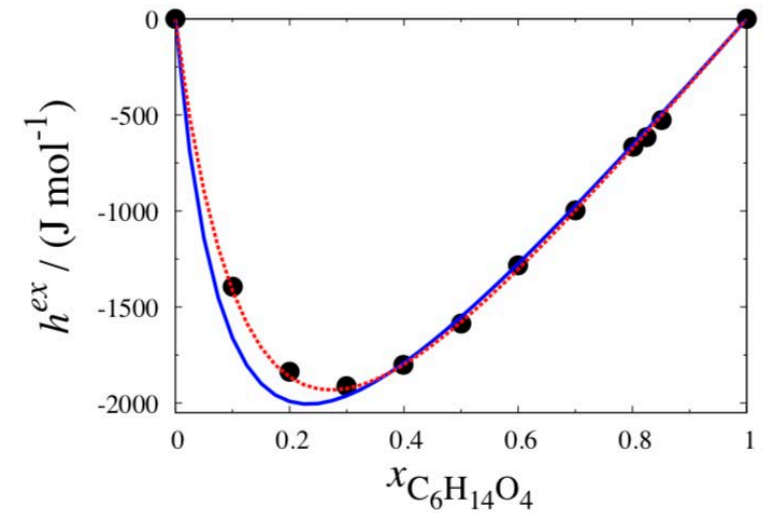
• OH-CH₂-CH₂-O-CH₂-
• triethylene glycol



• -CH₂-O-CH₂-
• ethers
• C₄H₁₀O₂ to
C₁₂H₂₆O₆

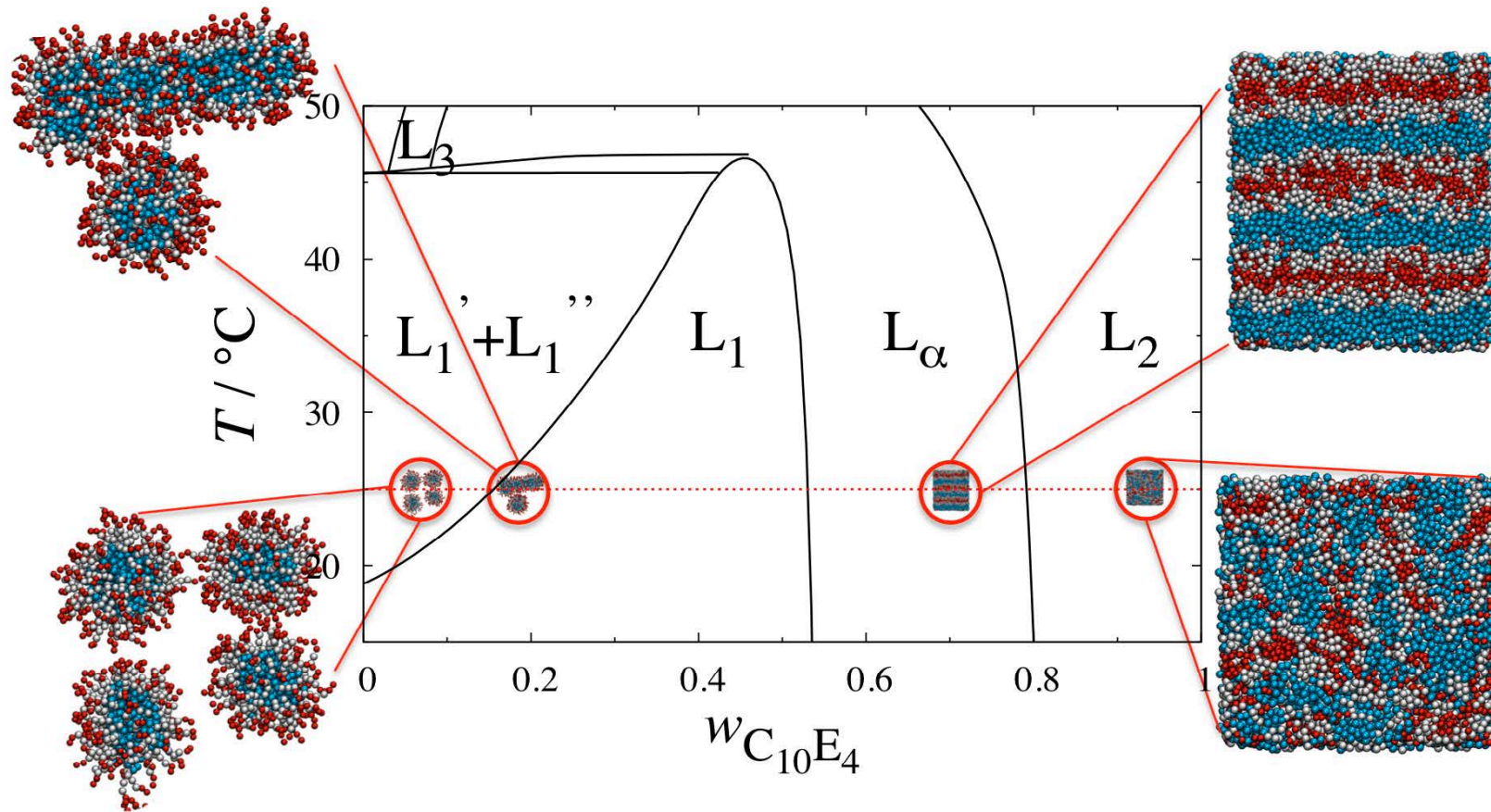


• -CH₂-CH₂-CH₂-
• alkanes
• C₃H₈ to C₂₄H₅₀



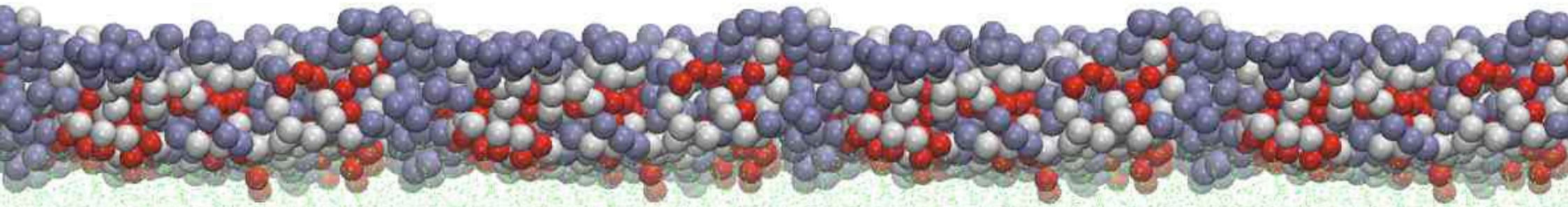
Properties of the mixture (here excess enthalpy of triethylene glycol-water) are used to estimate the unlike interaction between the surfactant and water

(C₁₀EO₄OH) in water



micellar phase L1, lamellar phase L α , and isotropic phase of inversed micelles L2

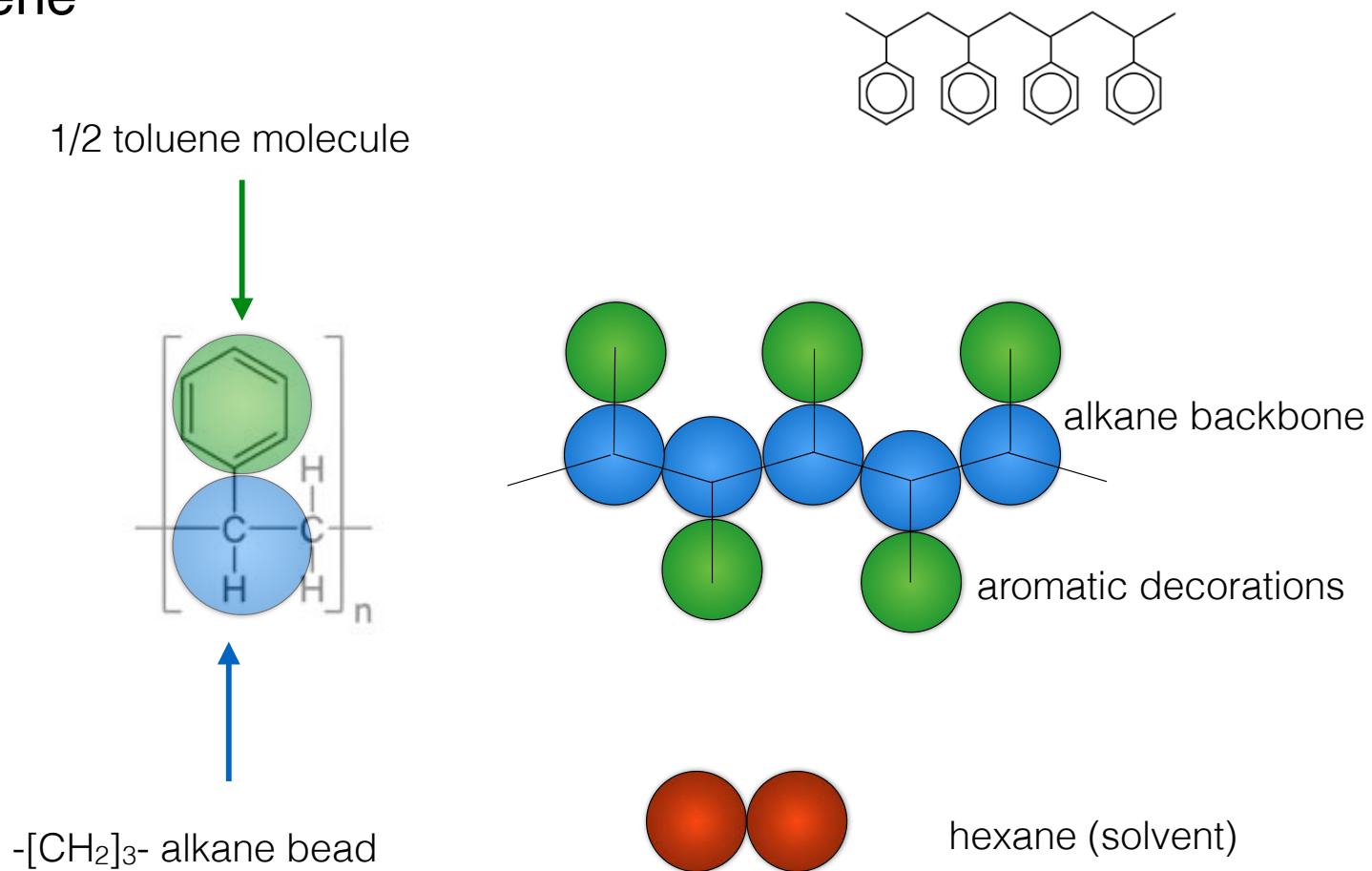
Air



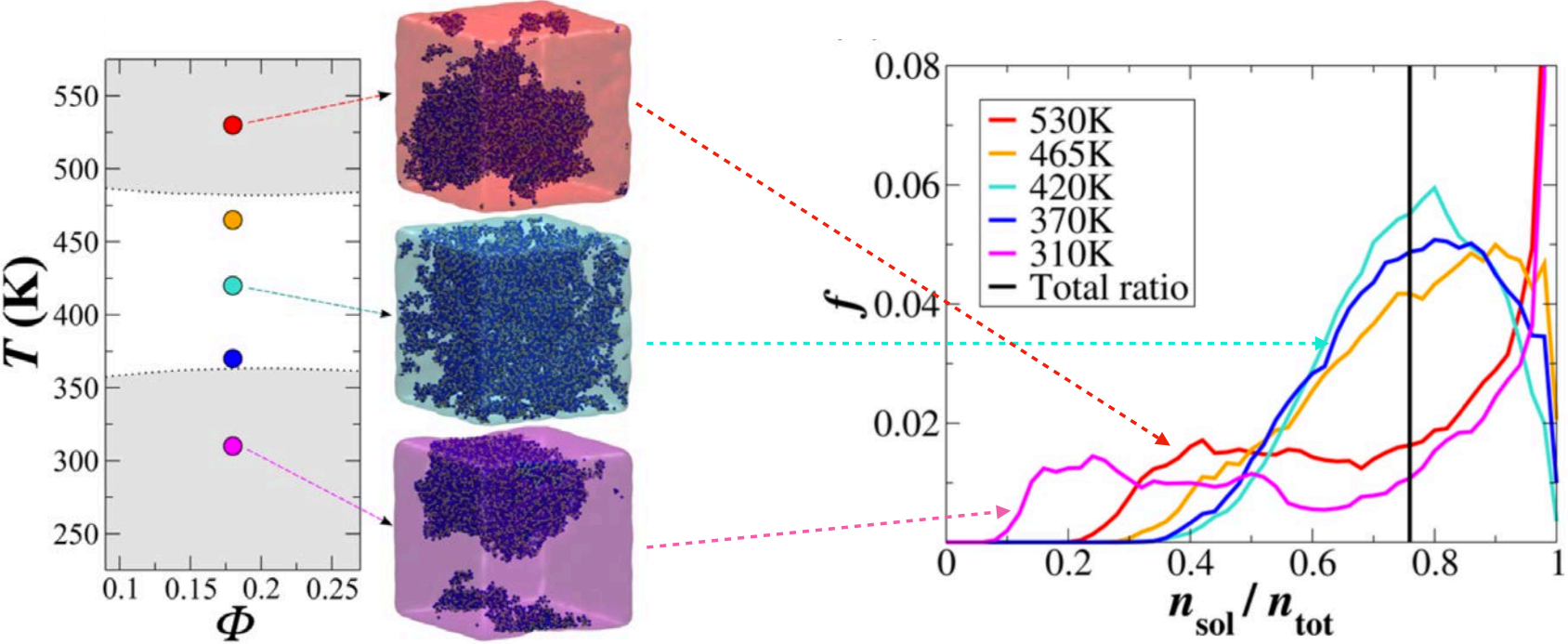
Water



Polystyrene

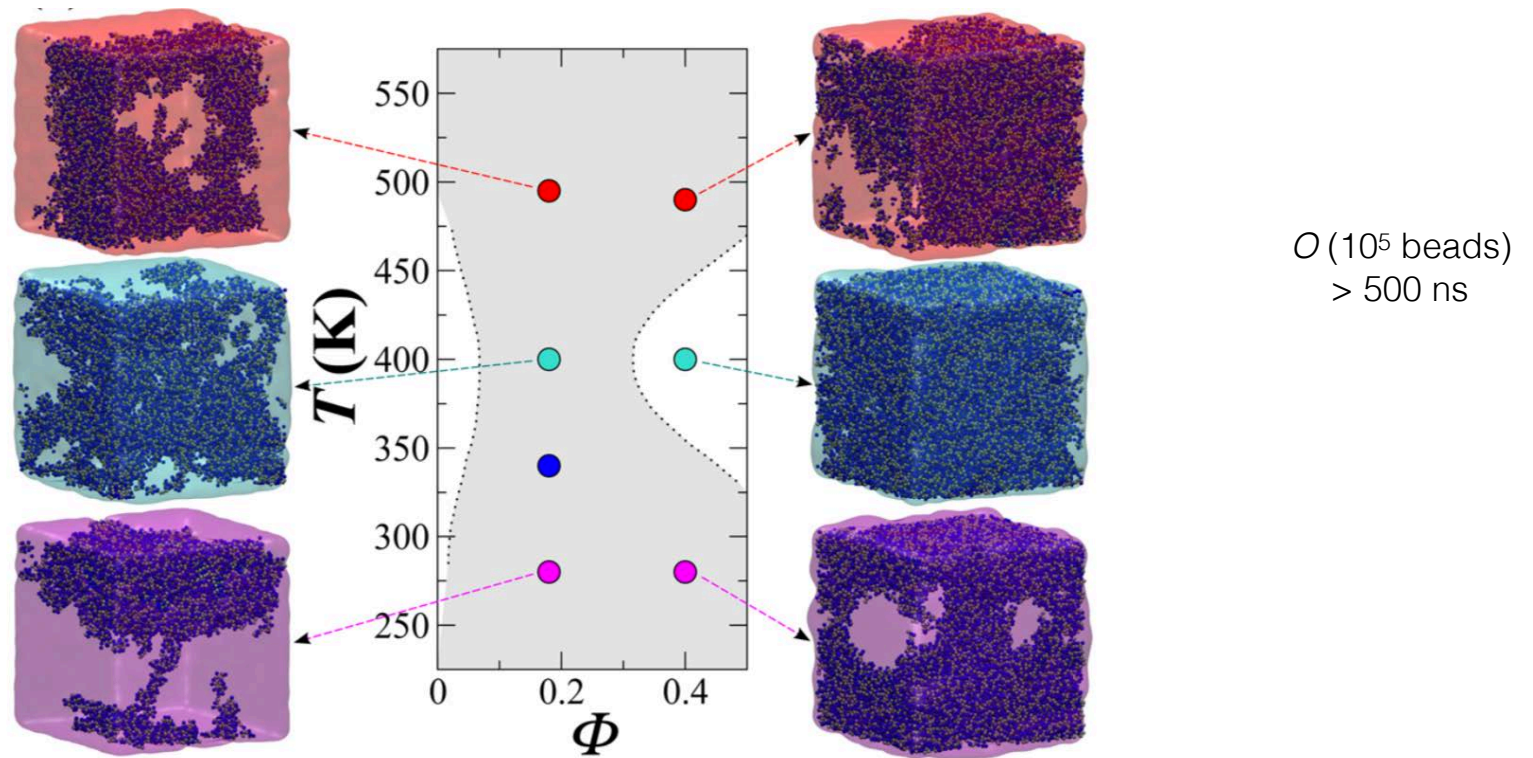


PS Mw 4800 g/mol in hexane



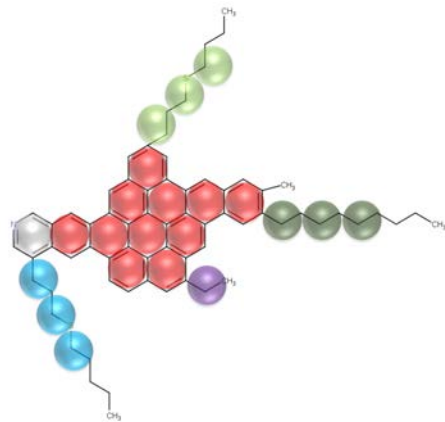
local density histograms with $(6)^3$ cells

PS Mw 4800 g/mol in heptane

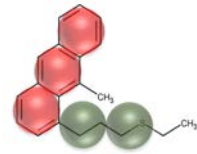


“hourglass” phase diagram is accurately predicted with the same model

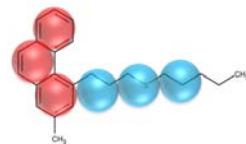
Synthetic live crude simulation (12 components)



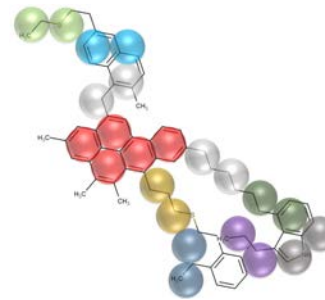
Asphaltene A18



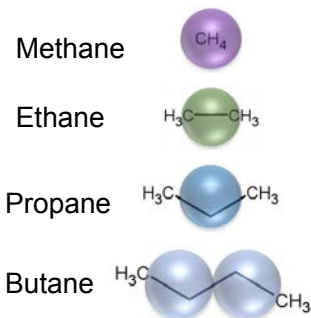
Resin A63



Resin A72



Asphaltene A81



Light ends

Heptane



Toluene



Anthracene



n-C18



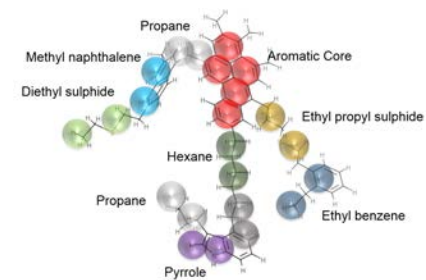
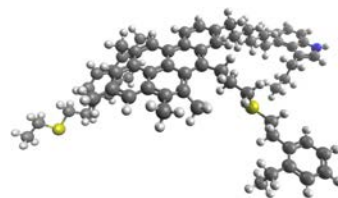
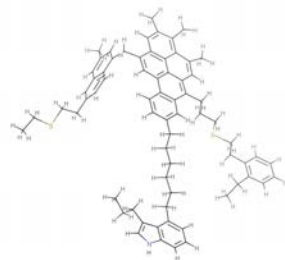
heavy ends

System size: $O(10^4)$ nm³
 138000 molecules
 2.4M atoms
 Temperature: 300K – 800K;
 Pressure range: 1 – 500 bar
 Time: 0.3 μ s
 Time step: 0.01 ps
 running on 48 cores
 Wall time: 1 month (10 ns/day)

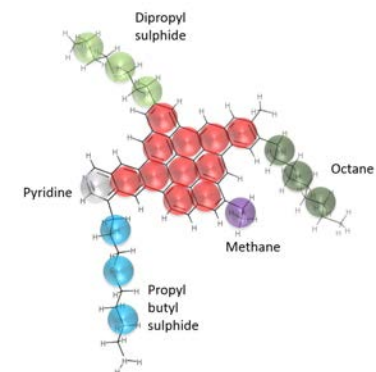
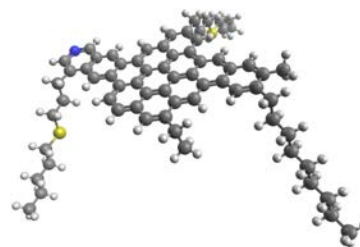
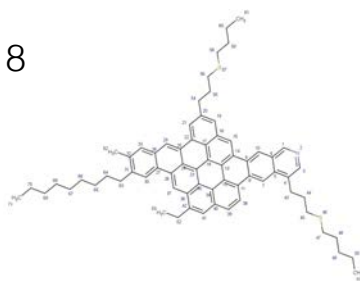
	Wt% in system
Asphaltene 18 (C68H73N1S2)	4.5%
Asphaltene 25 (C70H79N1S2)	1.2%
C1	5.8%
C2	1.9%
C3	2.8%
nC4	5.1%
Saturated n-C7	57.7%
Aromatic Toluene (C7)	3.0%
Aromatic Anthracene (C14)	3.9%
Saturated Wax (n-C18)	9.4%
Resin A (C20H22S1)	2.0%
Resin B (C23H28S1)	2.3%

Twins with different behaviour

“good” archipelago asphaltene A81
10 rings, largest core has 5
DBE 32
MW 984
 $C_{69}H_{77}N_1S_2$



“bad” continental asphaltene A18
12 rings, all in one core
DBE 33
MW 968
 $C_{68}H_{73}N_1S_2$

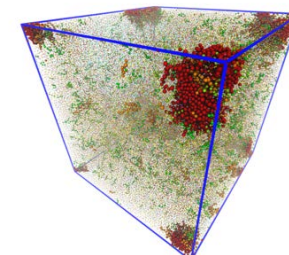


Atomistic Model
(for reference)

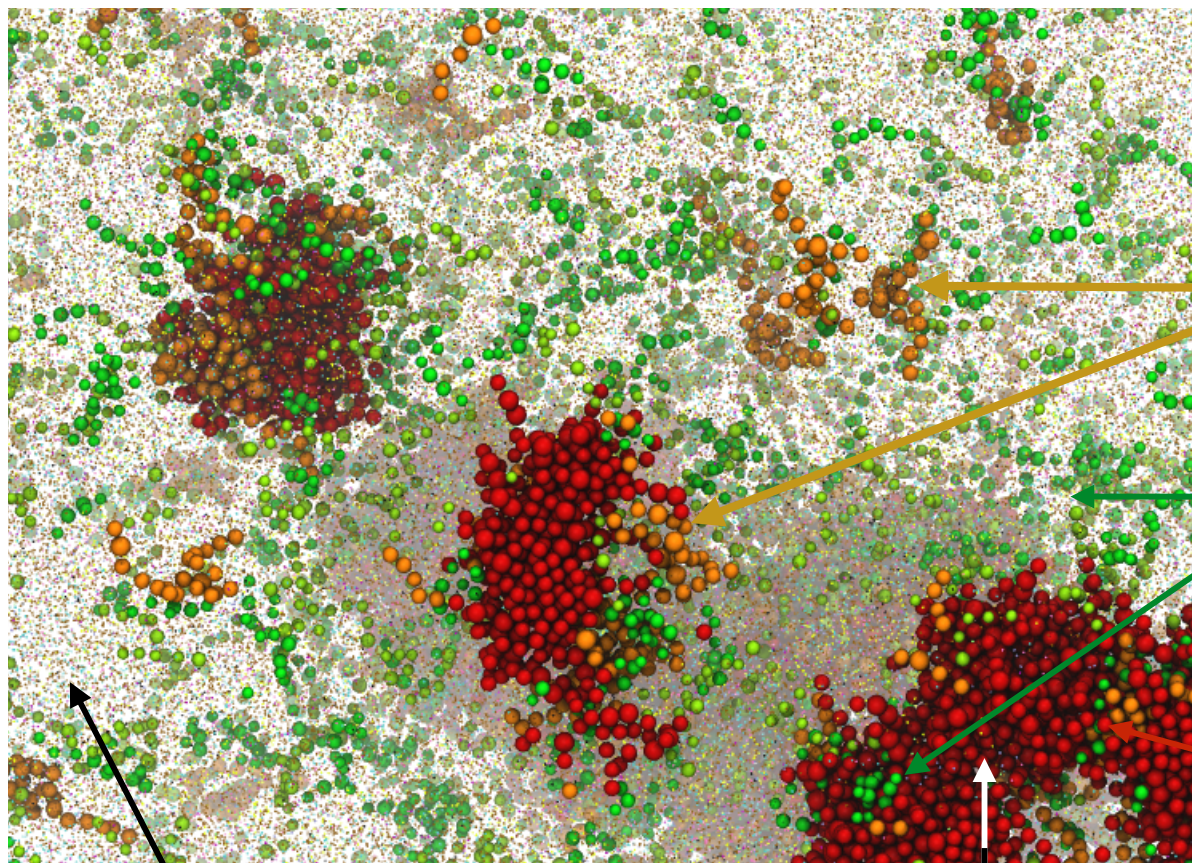
Coarse-grained model

Large scale simulations

410 K , 100 bar
7% w/w asphaltene



An equivalent all-atom simulation
would require $O(10^6)$ particles



Archipelago (good)
asphaltenes are scattered in the
bulk and within the aggregates

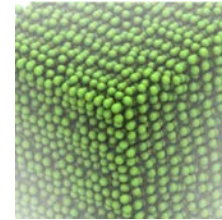
Resins are not solvating agents

Continental (bad) asphaltenes
form the core of the aggregates

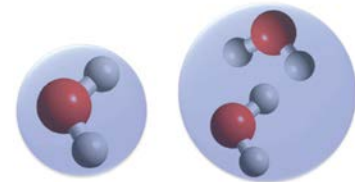
n-alkanes do not form part of the aggregates while **toluene**
does (aggregates are loose structures)

Known issues

- Premature solidification - enhanced by the spherical nature of the segments and the propensity of forming an FCC lattice.

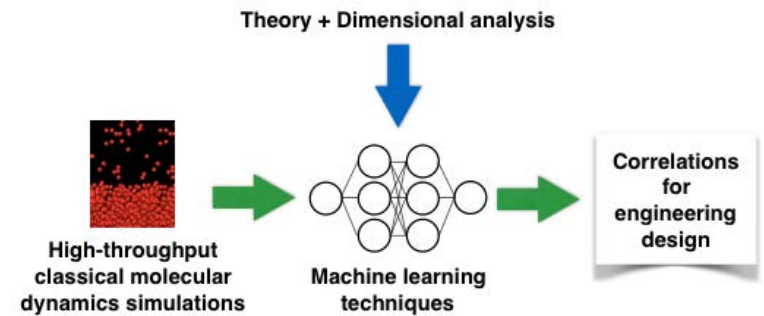
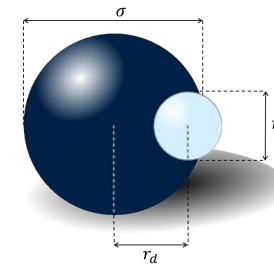
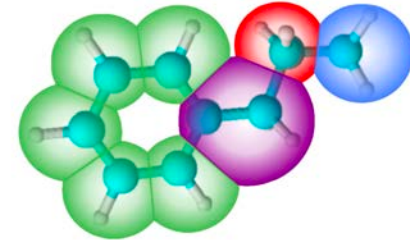


- Charges, electrostatics, etc. - It is challenging to develop the EoS for these systems, hence difficult to parametrize. One can employ “average” potentials which include charges implicitly. e.g. water



Outlook (the to-do list)

- Fused (or less coarse grained models) informed by the EoS.
- Explicit inclusion of association through molecular models
- Use of machine learning to bypass the EoS



Conclusion

- The use of a top-down approach relieves some of the fundamental issues associated with coarse-graining, in particular the “washing-out” of information when going from a finer to a more coarse scale.
- The SAFT CG approach employs the integrated information contained in the macroscopical thermophysical properties to obtain a surrogate “effective” potential which is transferrable, representative and accurate.
- The use of an analytical equation of state to make the link between the macroscopic properties and the intermolecular potential is the key to the success, but also the limitation in the universal application. Charged systems are particularly challenging.

For a list of current applications of the SAFT CG force field see <http://molecularsystemsengineering.org/saft.html>

Acknowledgements

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- ★ Carmelo Herdes
- ★ Olga Lobanova



Question and Answer Session



Professor Erich Müller

Imperial College London



Dr. Marianna Yiannourakou

Materials Design

Questions about Materials Design Webinars

Katherine Hollingsworth

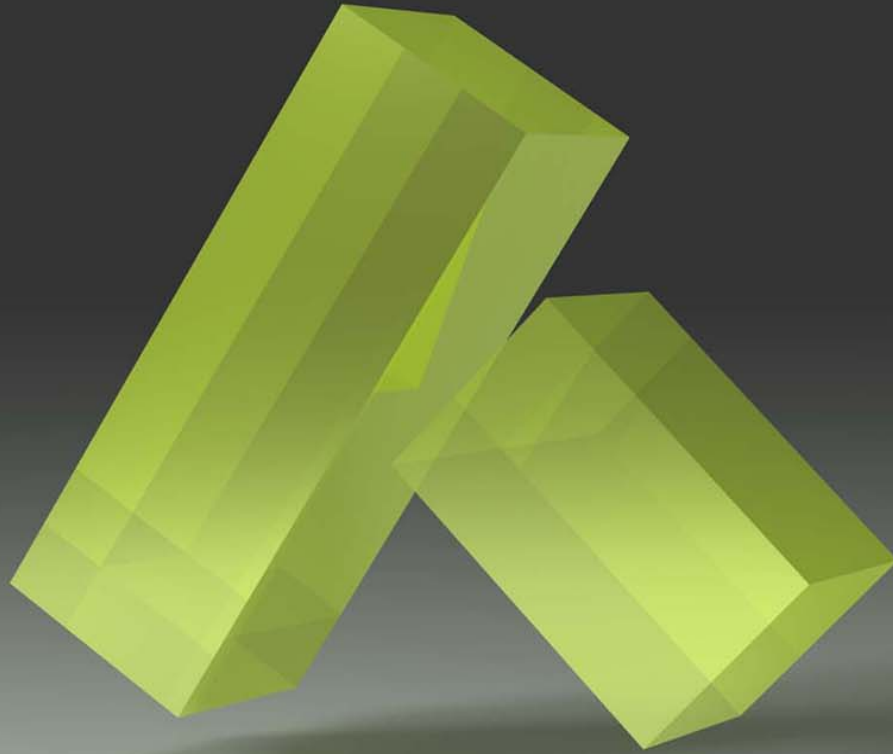
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Medea

Innovation by Simulation

