



New Horizons in Atomistic Simulation, York, January 5, 2018

Computational molecular engineering in interfacial thermodynamics

Martin Thomas Horsch



Engineering Department
American University of Iraq, Sulaimani (AUIS)

In collaboration with:

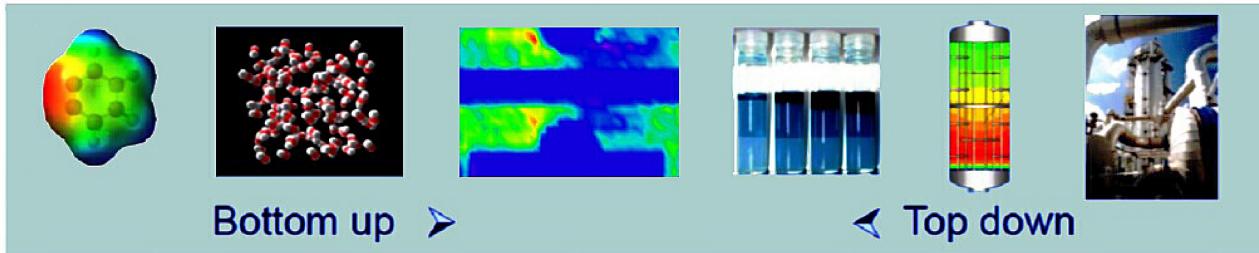
Wolfgang Eckhardt, Munich
Hans Hasse, Kaiserslautern
Peter Klein, Kaiserslautern
Philipp Neumann, Hamburg
Simon Stephan, Kaiserslautern
Katrin Stöbener, Kaiserslautern
Jadran Vrabec, Paderborn
Stephan Werth, Kaiserslautern



Computational
Molecular Engineering



Computational molecular engineering

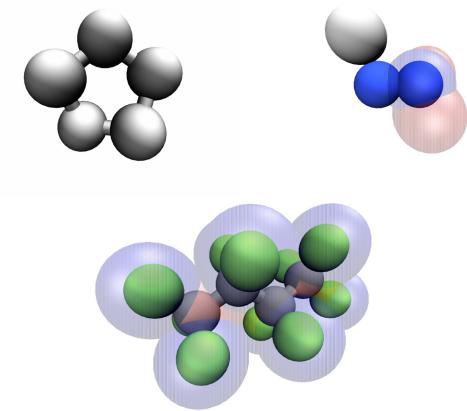


Physical Chemistry
(qualitative validity)



Chemical Engineering
(quantitative reliability)

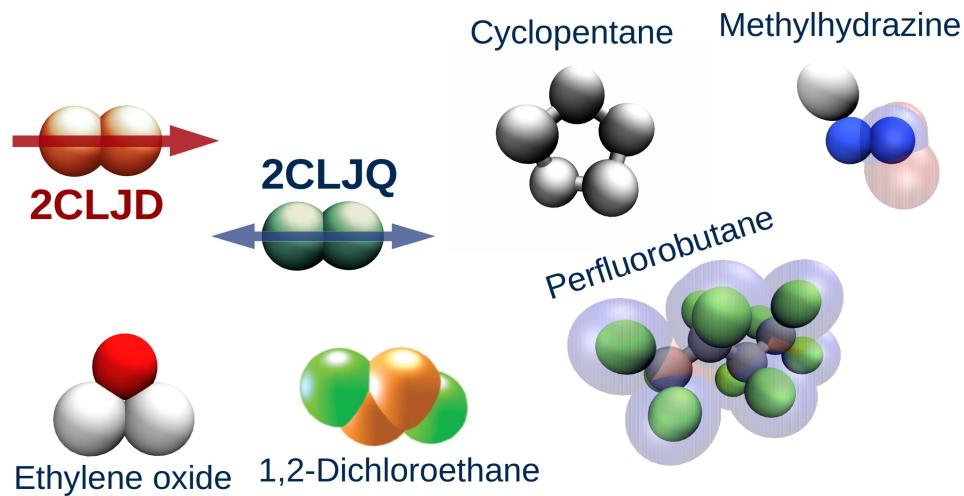
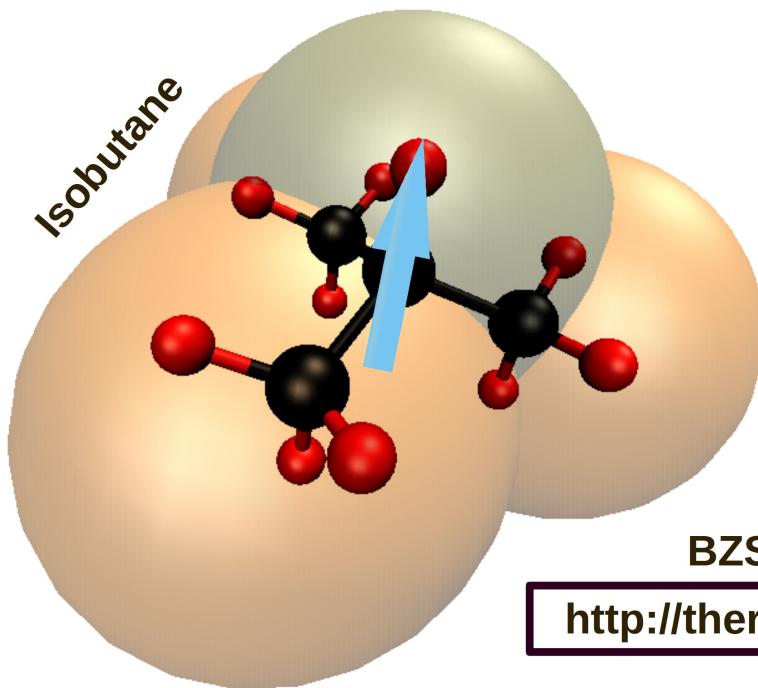
- Realistic representation of pairwise intermolecular interactions
- Short-range repulsion and dispersion as well as long-range electrostatics (multipoles)
- Models with free parameters, adjusted to thermodynamic data
- Reliable interpolation, extrapolation and prediction of fluid properties




**Computational
Molecular Engineering**



Molecular models of the Boltzmann-Zuse Society



BZS Molecular Model Database

<http://thermovm.mv.uni-kl.de/moleculeDB/>

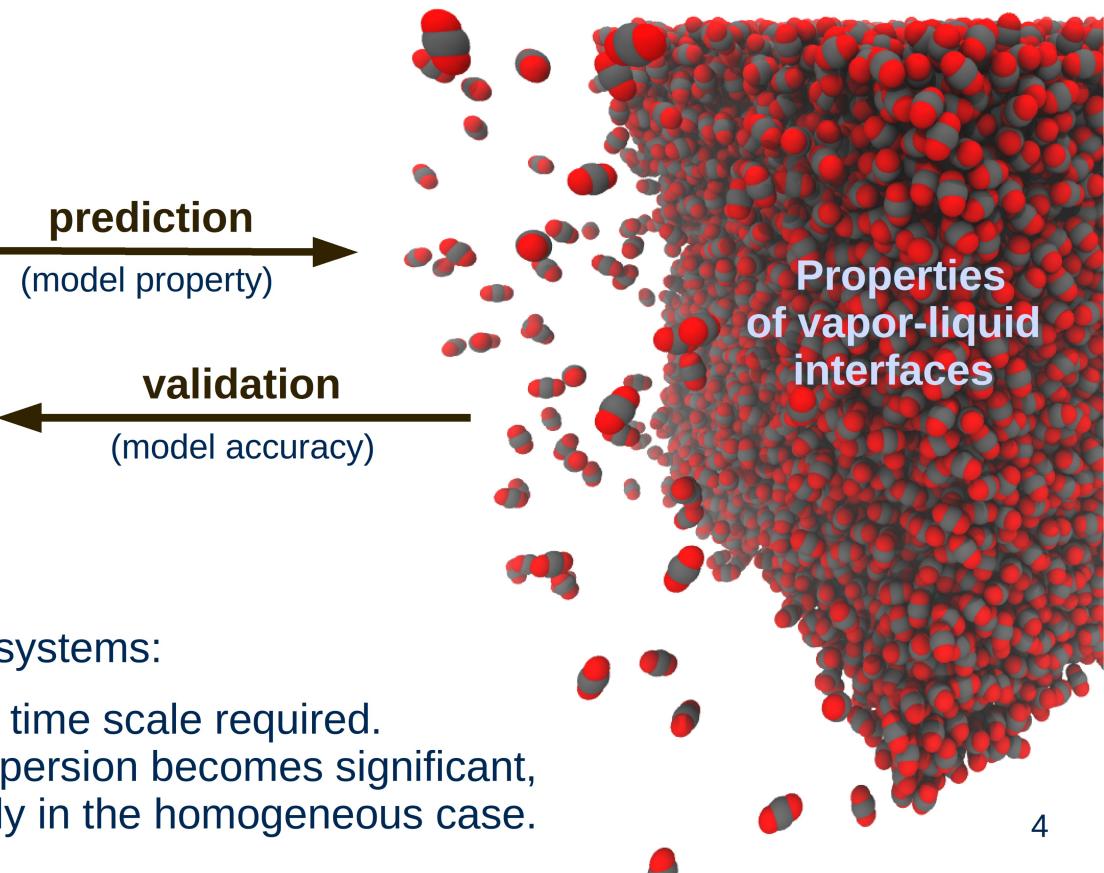
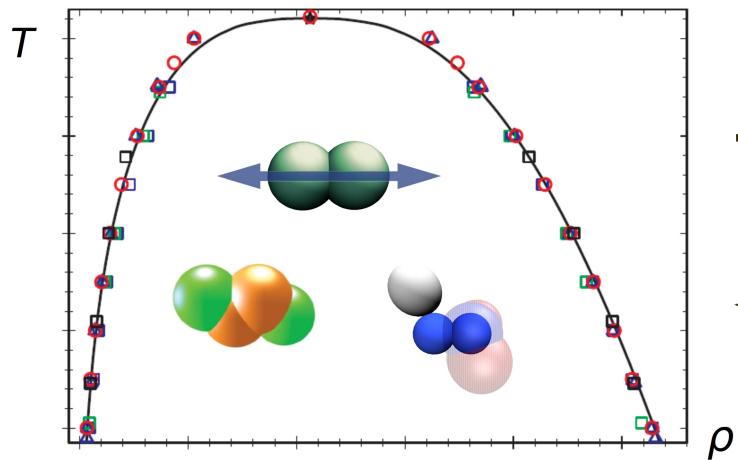
(Model parameters adjusted to vapor-liquid equilibrium data)



Computational
Molecular Engineering

Molecular simulation of interfacial properties

VLE data: Bulk fluid properties



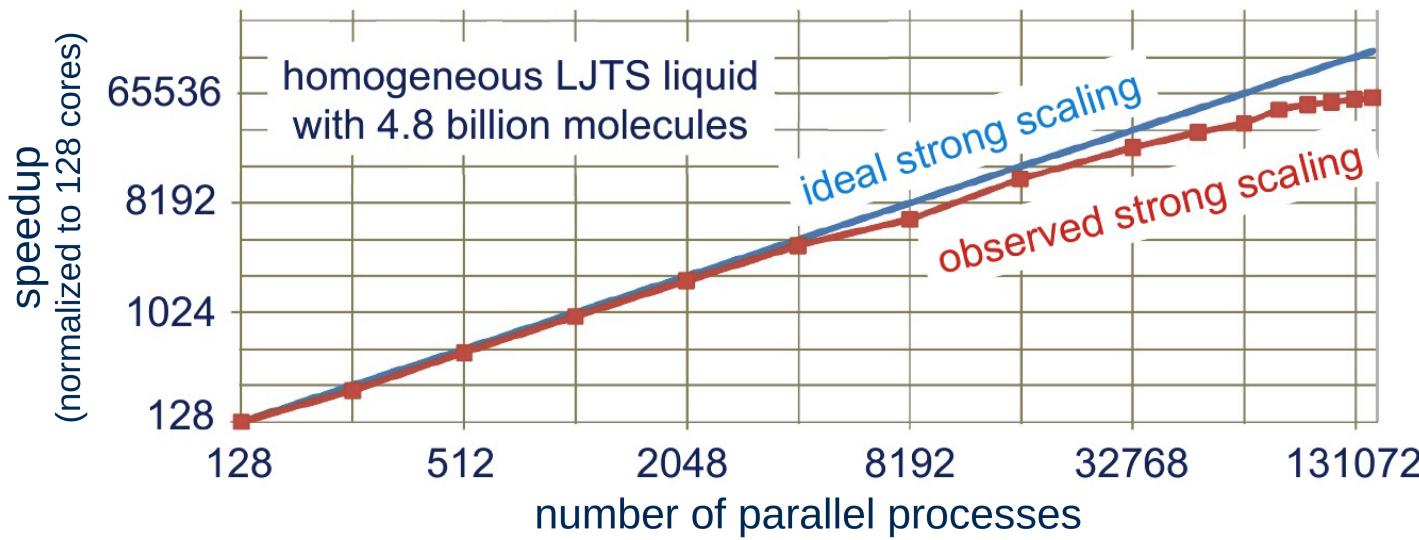
Molecular simulation of heterogeneous systems:

- Greater length scale and greater time scale required.
- Long-range contribution from dispersion becomes significant, whereas it can be handled trivially in the homogeneous case.

Massively-parallel molecular dynamics

<http://www.ls1-mardyn.de/>

(large systems 1: molecular dynamics)^{1, 2}



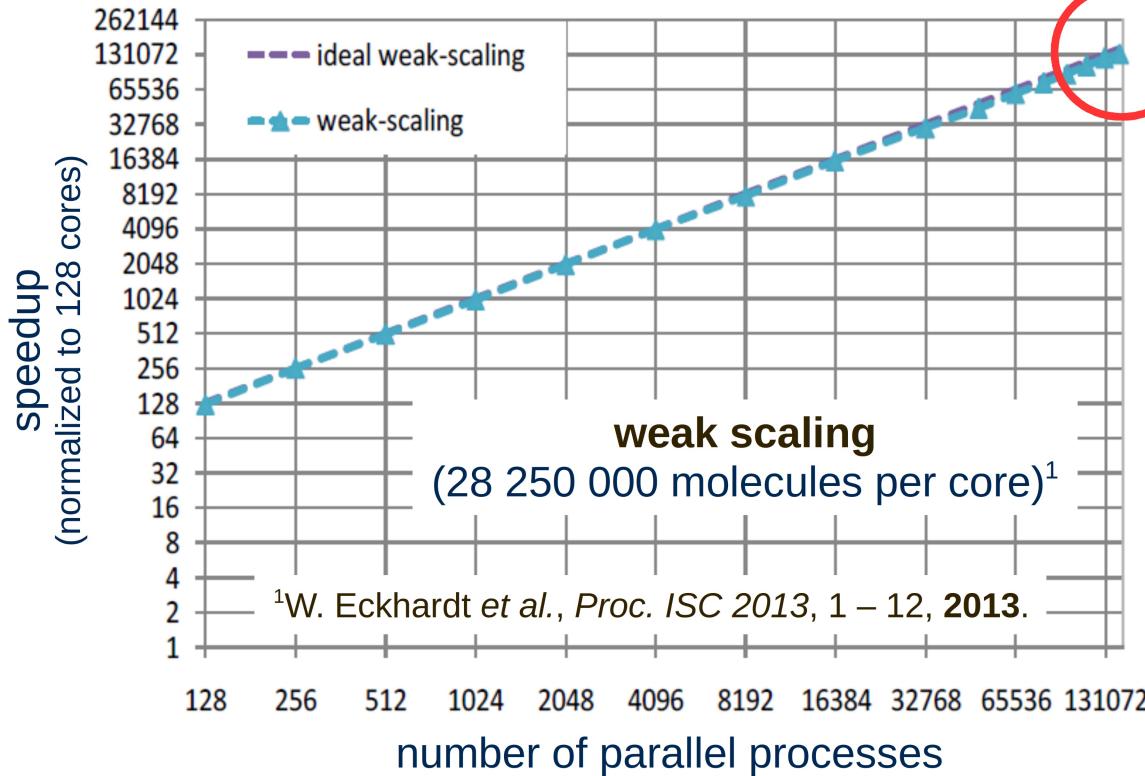
¹C. Niethammer et al., *J. Chem. Theory Comput.* 10(10), 4455 – 4464, 2014.

²W. Eckhardt et al., *Proc. ISC 2013, LNCS 7905, 1 – 12, 2013.*



Computational
Molecular Engineering

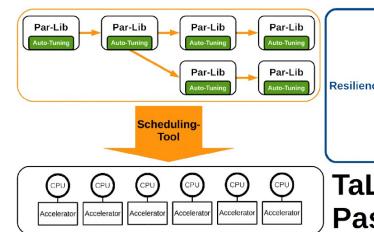
Massively-parallel molecular dynamics



$N = 4\ 125\ 000\ 000\ 000$

SuperMUC (146 000 Cores)¹

(December 2017: $N > 20\ 000\ 000\ 000\ 000$)



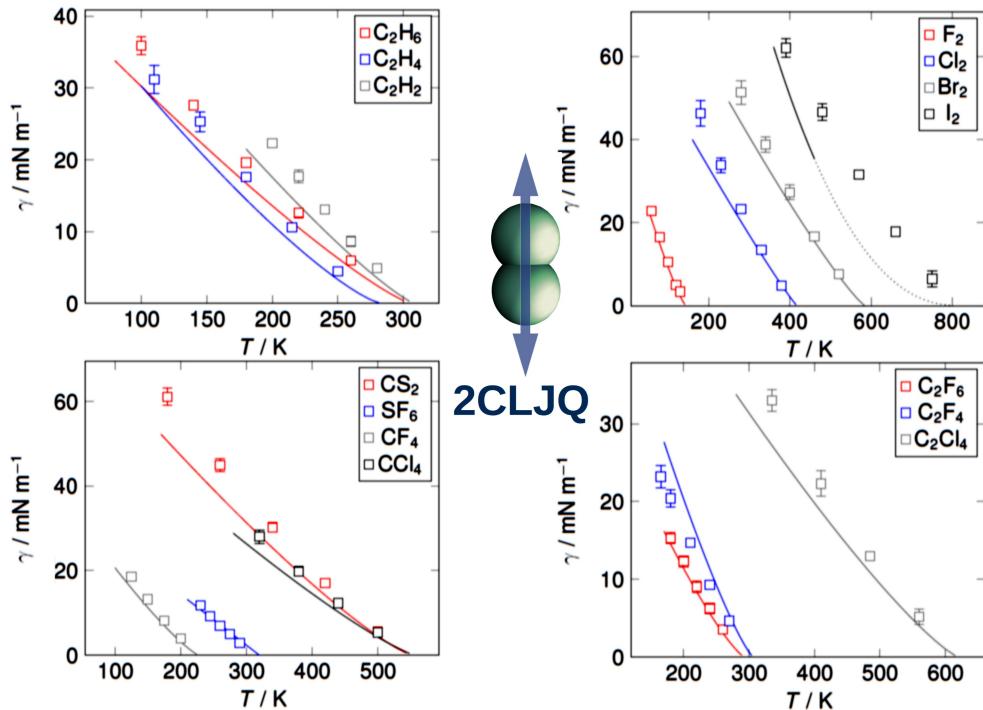
TECHNISCHE
UNIVERSITÄT
MÜNCHEN



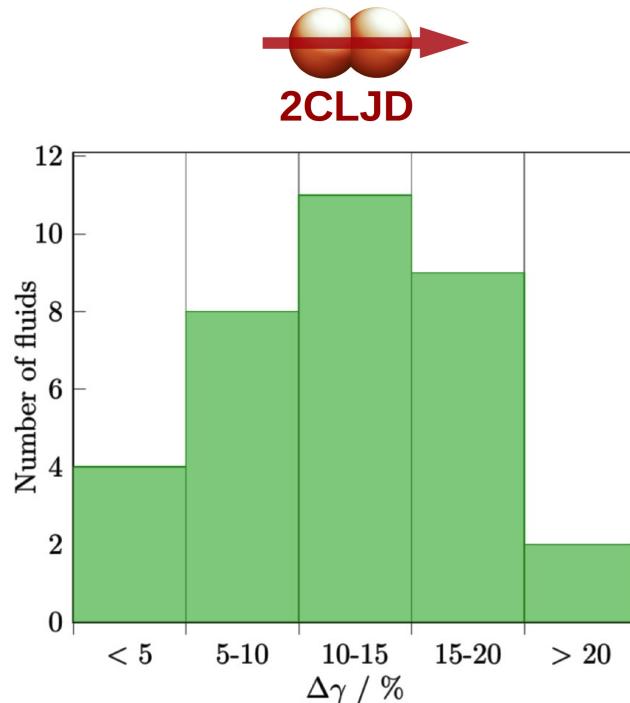
Computational
Molecular Engineering

Surface tension predicted by molecular models

2 LJ sites + quadrupole¹



2 LJ sites + dipole²



¹S. Werth et al., *Chem. Eng. Sci.* 121, 110 – 117, 2015; ²S. Werth et al., *J. Chem. Phys.* 144, 054702, 2016.



Surface tension predicted by molecular models

Non-polar: LJ

Neon (Ne), Argon (Ar)
Krypton (Kr), Xenon (Xe)
Methane (CH_4)

Dipolar: 2CLJD

Carbon monoxide (CO)

R11 (CFCI_3)

R12 (CF_2Cl_2)

R13 (CF_3Cl)

R13B1 (CBrF_3)

R22 (CHF_2Cl)

R23 (CHF_3)

R41 (CH_3F)

R123 ($\text{CHCl}_2\text{-CF}_3$)

R124 (CHFCI-CF_3)

R125 ($\text{CHF}_2\text{-CF}_3$)

R134a ($\text{CH}_2\text{F-CF}_3$)

R141b ($\text{CH}_3\text{-CFCl}_2$)

R142b ($\text{CH}_3\text{-CF}_2\text{Cl}$)

R143a ($\text{CH}_3\text{-CF}_3$)

R152a ($\text{CH}_3\text{-CHF}_2$)

R40 (CH_3Cl)

R40B1 (CH_3Br)

Methyl iodide (CH_3I)

R30B1 (CH_2BrCl)

R20 (CHCl_3)

R20B3 (CHBr_3)

R21 (CHFCl_2)

R32 (CH_2F_2)

R30 (CH_2Cl_2)

12 %

Dipolar: 2CLJD (contd.)

R30B2 (CH_2Br_2)
Methylene iodide (CH_2I_2)
R12B2 (CBr_2F_2)
R12B1 (CBrClF_2)
R10B1 (CBrCl_3)
R161 ($\text{CH}_2\text{F-CH}_3$)
R150a ($\text{CHCl}_2\text{-CH}_3$)
R140 ($\text{CHCl}_2\text{-CH}_2\text{Cl}$)
R140a ($\text{CCl}_3\text{-CH}_3$)
R130a ($\text{CH}_2\text{Cl-CCl}_3$)
R160B1 ($\text{CH}_2\text{Br-CH}_3$)
R150B2 ($\text{CHBr}_2\text{-CH}_3$)
R131b ($\text{CH}_2\text{F-CCl}_3$)
R123B1 (CHClBr-CF_3)
R112a ($\text{CCl}_3\text{-CF}_2\text{Cl}$)
R1141 (CHF=CH_2)
R1132a ($\text{CF}_2\text{=CH}_2$)
R1140 (CHCl=CH_2)
R1122 (CHCl=CF_2)
R1113 (CFCl=CF_2)
R1113B1 (CFBr=CF_2)

20 %

Quadrupolar: 2CLJQ

Fluorine (F_2)
Chlorine (Cl_2)
Bromine (Br_2)
Iodine (I_2)
Nitrogen (N_2)
Oxygen (O_2)
Carbon dioxide (CO_2)

Quadrupolar: 2CLJQ (contd.)

Carbon disulfide (CS_2)
Ethane (C_2H_6)
Ethylene (C_2H_4)
Acetylene (C_2H_2)
R116 (C_2F_6)
R1114 (C_2F_4)
R1110 (C_2Cl_4)
Propadiene ($\text{CH}_2=\text{C=CH}_2$)
Propyne ($\text{CH}_3\text{-C}\equiv\text{CH}$)
Propylene ($\text{CH}_3\text{-CH=CH}_2$)
R846 (SF_6)
R14 (CF_4)
R10 (CCl_4)
R113 ($\text{CFCl}_2\text{-CF}_2\text{Cl}$)
R114 ($\text{CF}_2\text{Cl-CF}_2\text{Cl}$)
R115 ($\text{CF}_3\text{-CF}_2\text{Cl}$)
R134 ($\text{CHF}_2\text{-CHF}_2$)
R150B2 ($\text{CH}_2\text{Br-CH}_2\text{Br}$)
R114B2 ($\text{CBrF}_2\text{-CBrF}_2$)
R1120 (CHCl=CCl_2)

22 %

Multi-site models

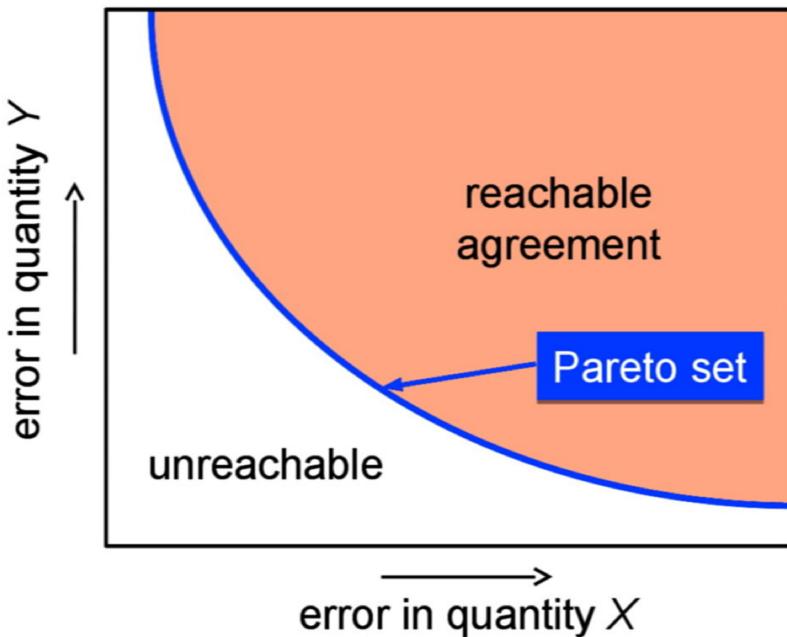
Isobutane (C_4H_{10})
Cyclohexane (C_6H_{12})
Methanol (CH_3OH)
Ethanol ($\text{C}_2\text{H}_5\text{OH}$)
Formaldehyde ($\text{CH}_2=\text{O}$)
Dimethylether ($\text{CH}_3\text{-O-CH}_3$)
Acetone ($\text{C}_3\text{H}_6\text{O}$)

Multi-site models (contd.)

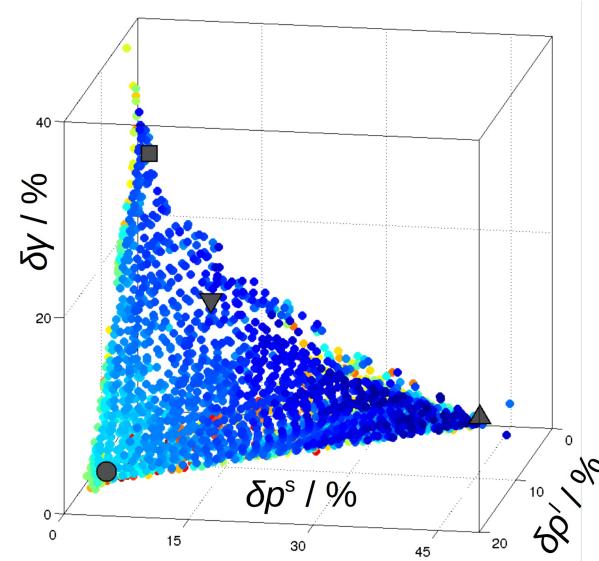
Ammonia (NH_3)
Methylamine ($\text{NH}_2\text{-CH}_3$)
Dimethylamine ($\text{CH}_3\text{-NH-CH}_3$)
R227ea ($\text{CF}_3\text{-CHF-CF}_3$)
Sulfur dioxide (SO_2)
Ethylene oxide ($\text{C}_2\text{H}_4\text{O}$)
Dimethylsulfide ($\text{CH}_3\text{-S-CH}_3$)
Hydrogen cyanide (HCN)
Acetonitrile (NC_2H_3)
Thiophene (SC_4H_4)
Nitromethane (NO_2CH_3)
Phosgene (COCl_2)
Benzene (C_6H_6)
Toluene (C_7H_8)
Chlorobenzene ($\text{C}_6\text{H}_5\text{Cl}$)
Dichlorobenzene ($\text{C}_6\text{H}_4\text{Cl}_2$)
Cyclohexanol ($\text{C}_6\text{H}_{11}\text{OH}$)
Cyclohexanone ($\text{C}_6\text{H}_{10}\text{O}$)
Cyanogen (C_2N_2)
Cyanogen chloride (CCIN)
Formic acid (CH_2O_2)
Ethylene glycol ($\text{C}_2\text{H}_6\text{O}_2$)
Water (H_2O)
Hydrazine (N_2H_4)
Methylhydrazine (CH_6N_2)
Dimethylhydrazine ($\text{C}_2\text{H}_8\text{N}_2$)
Perfluorobutane (C_4F_{10})
Ethyl acetate ($\text{C}_4\text{H}_8\text{O}_2$)
Hexamethyldisiloxane ($\text{C}_6\text{H}_{12}\text{OSi}_2$)
Octamethylcyclotetrasiloxane ($\text{C}_8\text{H}_{24}\text{O}_4\text{Si}_4$)

Multicriteria optimization

Pareto optimality criterion



Multiple objectives

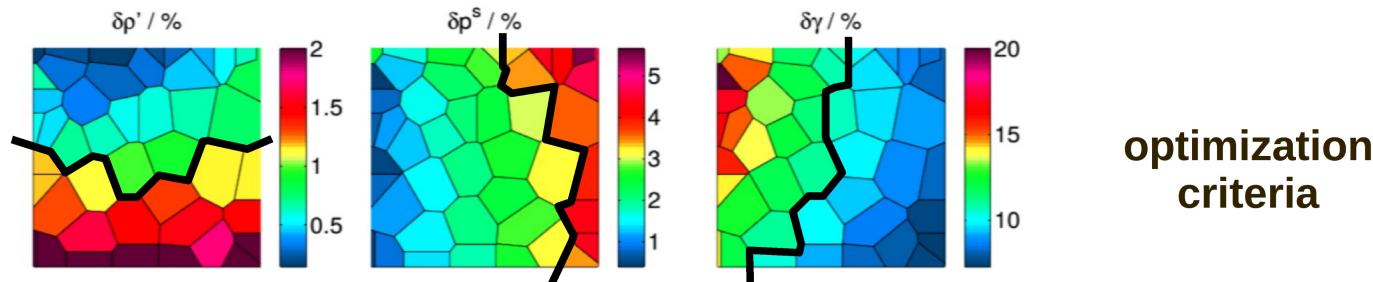


2CLJQ models for CO_2

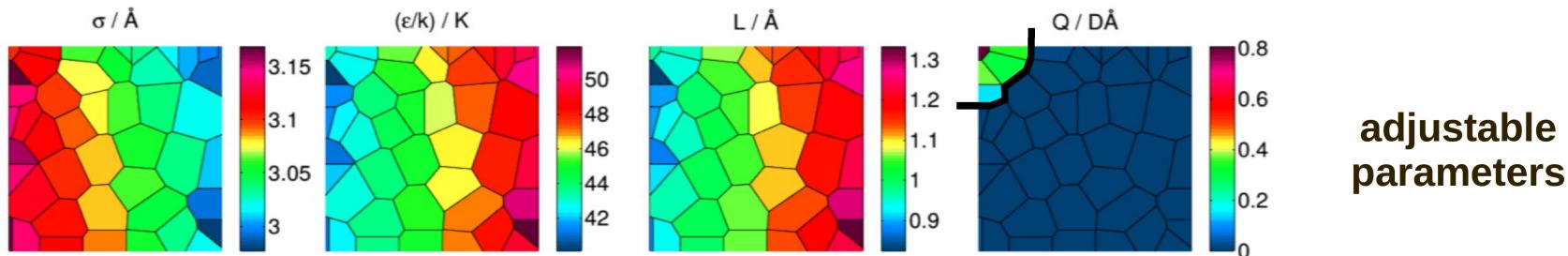
Prerequisite: Thermodynamic properties of the **model class** have been characterized.

Multicriteria optimization of molecular models

Self-organized patch plots¹ visualizing the Pareto front and the Pareto-optimal models:



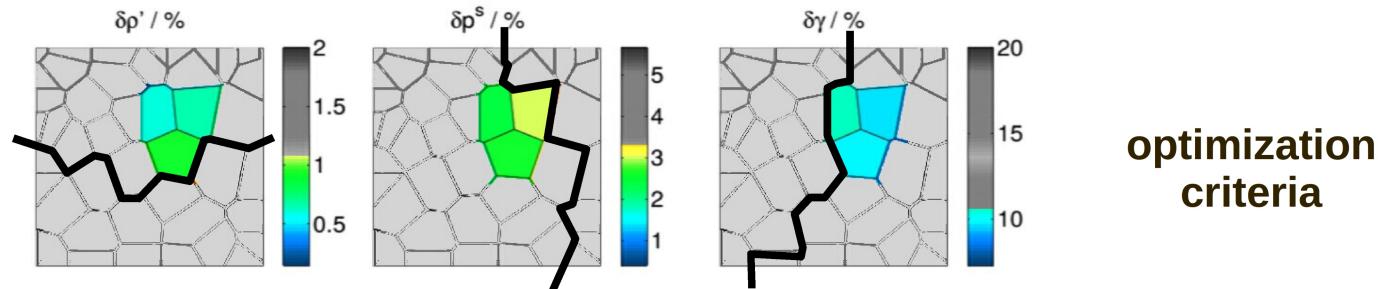
Pareto-optimal 2CLJQ models for oxygen



¹K. Stöbener et al., *Fluid Phase Equilib.* 411, 33 – 42, 2016.

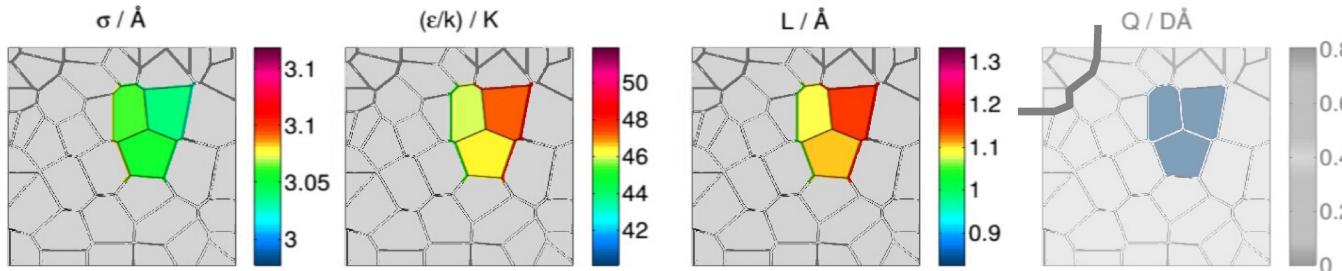
Multicriteria optimization of molecular models

Self-organized patch plots¹ visualizing the Pareto front and the Pareto-optimal models:



optimization
criteria

Pareto-optimal 2CLJ models satisfying all constraints



adjustable
parameters

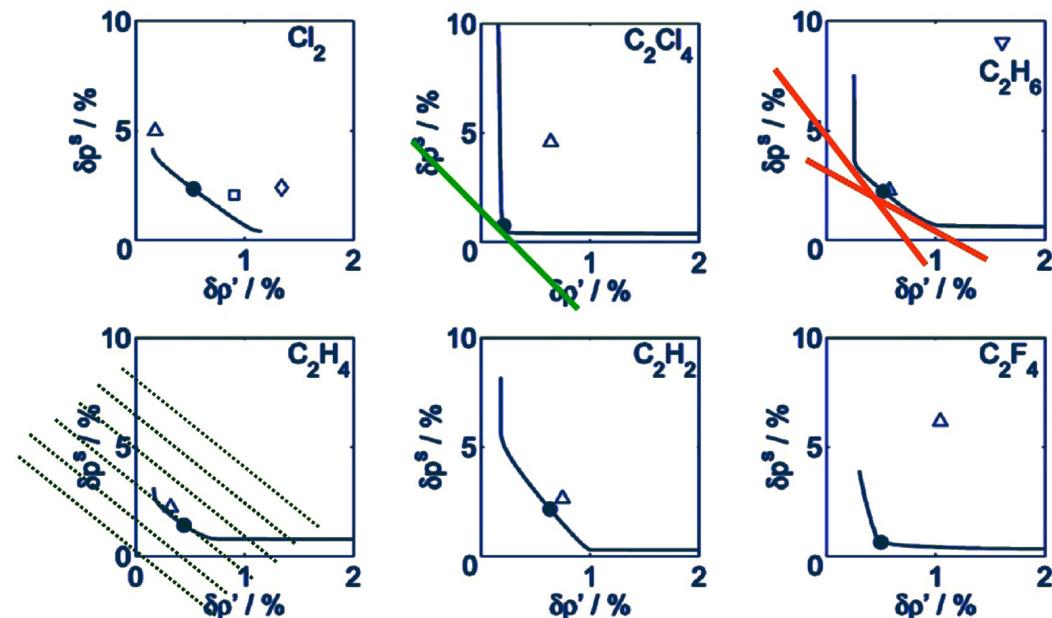
¹K. Stöbener et al., *Fluid Phase Equilib.* 411, 33 – 42, 2016.

Model resiliency with respect to priority shifts

Pareto knee:

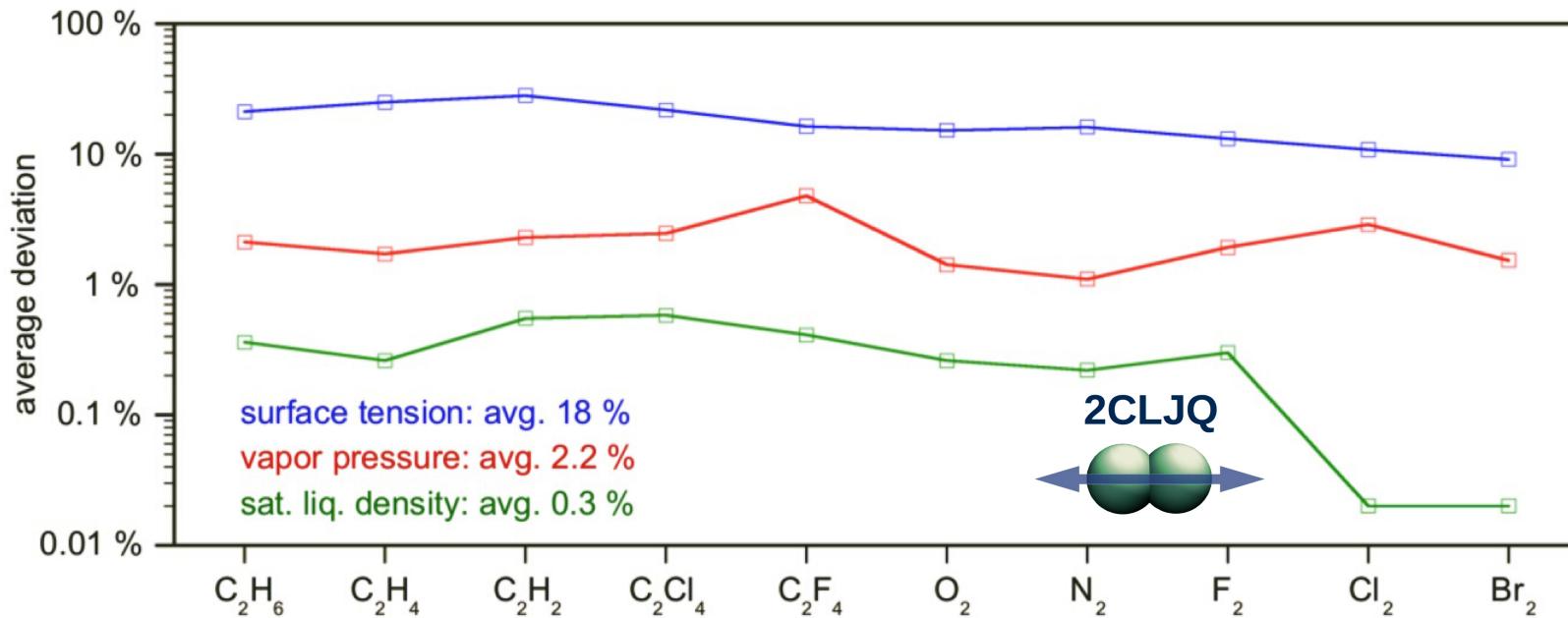
- Position where the curvature of the Pareto front becomes maximal, or
- central point of a possibly non-convex region between maximally curved parts.

The quality of models corresponding to the Pareto knee are relatively robust even if design priorities vary significantly.



Scalarizations based on linear combinations of optimization criteria are inadequate for concave regions of the Pareto front. A systematic exploration is needed to find the Pareto knee.

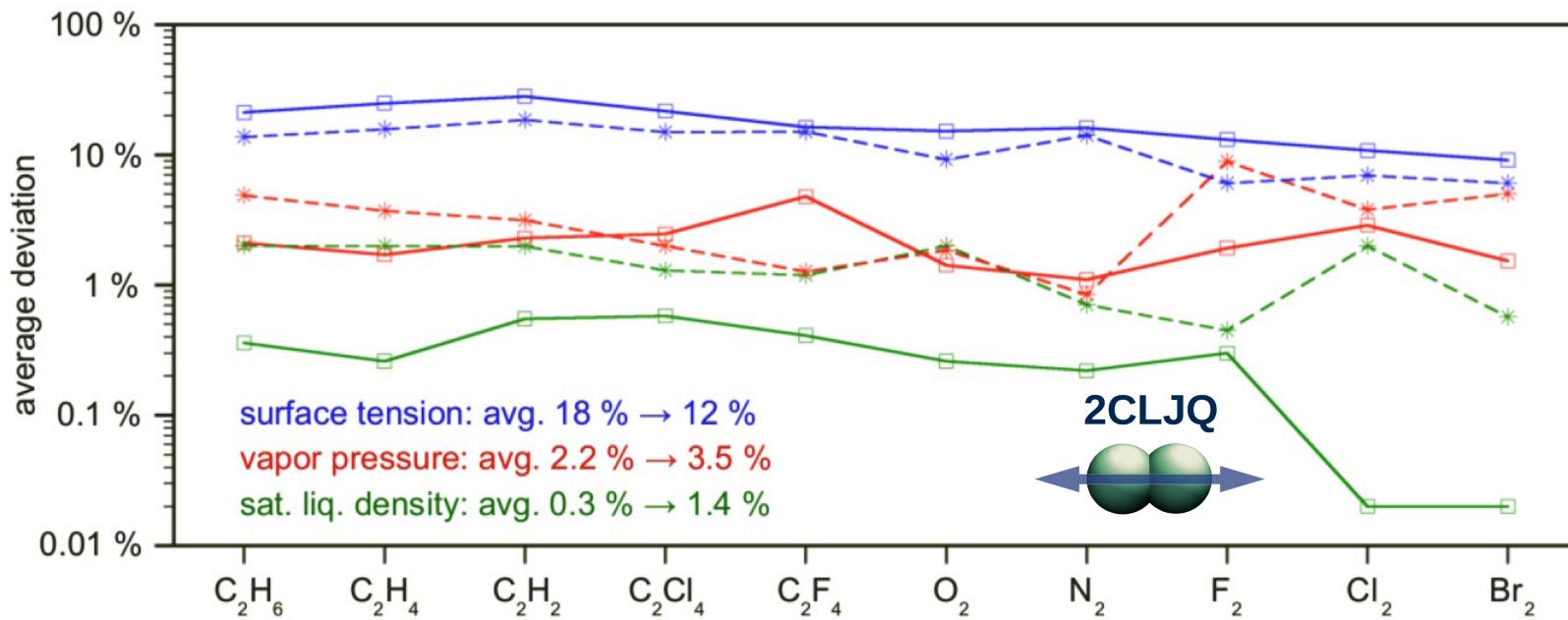
Model reparameterization for quadrupolar fluids^{1, 2}



¹J. Vrabec, J. Stoll, H. Hasse, *J. Phys. Chem. B* 105(48), 12126–12133, 2001;

²S. Werth, K. Stöbener, P. Klein, K.-H. Küfer, M. Horsch, H. Hasse, *Chem. Eng. Sci.* 121, 110–117, 2015.

Model reparameterization for quadrupolar fluids¹⁻³



¹J. Vrabec, J. Stoll, H. Hasse, *J. Phys. Chem. B* 105(48), 12126–12133, 2001;

²S. Werth, K. Stöbener, P. Klein, K.-H. Küfer, M. Horsch, H. Hasse, *Chem. Eng. Sci.* 121, 110–117, 2015;

³K. Stöbener, P. Klein, M. Horsch, K.-H. Küfer, H. Hasse, *Fluid Phase Equilib.* 411, 33–42, 2016.

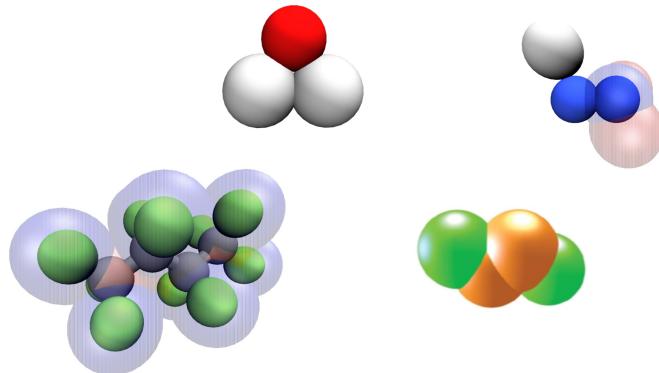


Paradigm shift in molecular modeling

Molecular model development as an “art” conducted by experts only

- Only a very limited community of people is qualified to design molecular models;
- for each fluid, an expert develops a model, which can be found in the literature;
- the way in which the model was designed and optimized is usually not disclosed.

Users need a background in molecular thermodynamics to assess the model quality, and even then, they **cannot actively contribute** to adjusting the model without repeating the expert work.



BZS Molecular Model Database

<http://thermovm.mv.uni-kl.de/moleculeDB/>



Paradigm shift in molecular modeling

Molecular model development as an “art” conducted by experts only

- Only a very limited community of people is qualified to design molecular models;
- for each fluid, an expert develops a model, which can be found in the literature;
- the way in which the model was designed and optimized is usually not disclosed.

Users need a background in molecular thermodynamics to assess the model quality, and even then, they **cannot actively contribute** to adjusting the model without repeating the expert work.

Molecular modeling as a “technology” accessible to all engineers

- Properties of the model class have been characterized (by previous expert work);
- on this basis, models are flexibly parameterized at the level of engineering practice.

Users who typically do not have a scientific background in molecular simulation methods are enabled to **actively guide** molecular modeling and simulation practice according to their needs.



Engineering Department
American University of Iraq, Sulaimani (AUIS)



Computational Molecular Engineering

In collaboration with:

Wolfgang Eckhardt
Hans Hasse
Peter Klein
Philipp Neumann

Simon Stephan
Katrin Stöbener
Jadran Vrabec
Stephan Werth

BMBF



SkaSim

