

Excitation vs. projection techniques: Towards sub-chemical accuracy for large noncovalent systems

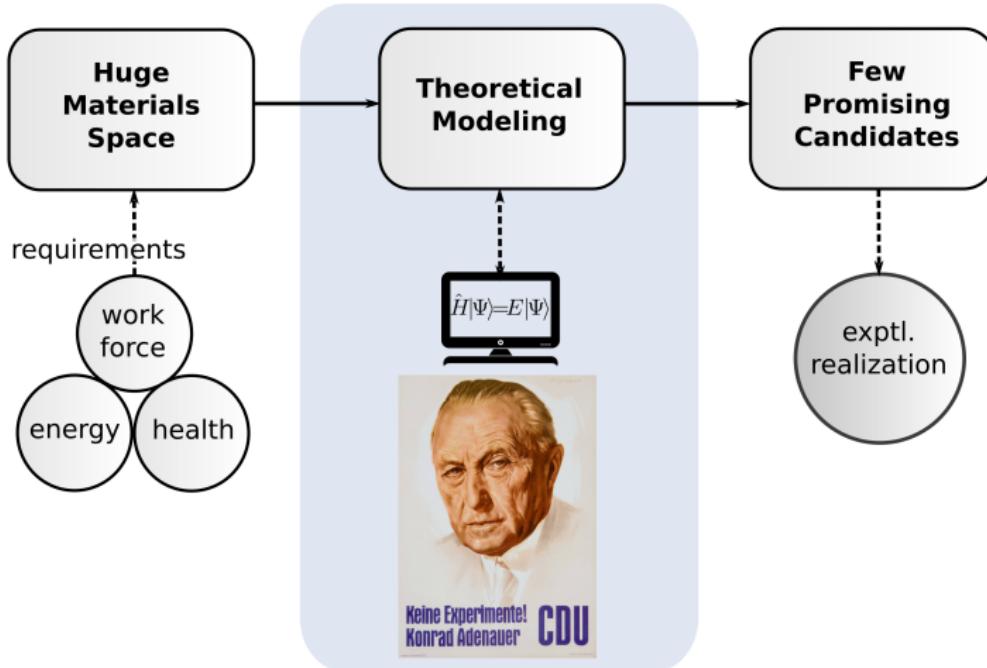
Gerit Brandenburg <g.brandenburg@ucl.ac.uk> | 8th of June 2018

SPECIAL PROBLEMS IN QUANTUM CHEMISTRY, MULLIKEN CENTER FOR THEORETICAL CHEMISTRY, BONN, GERMANY

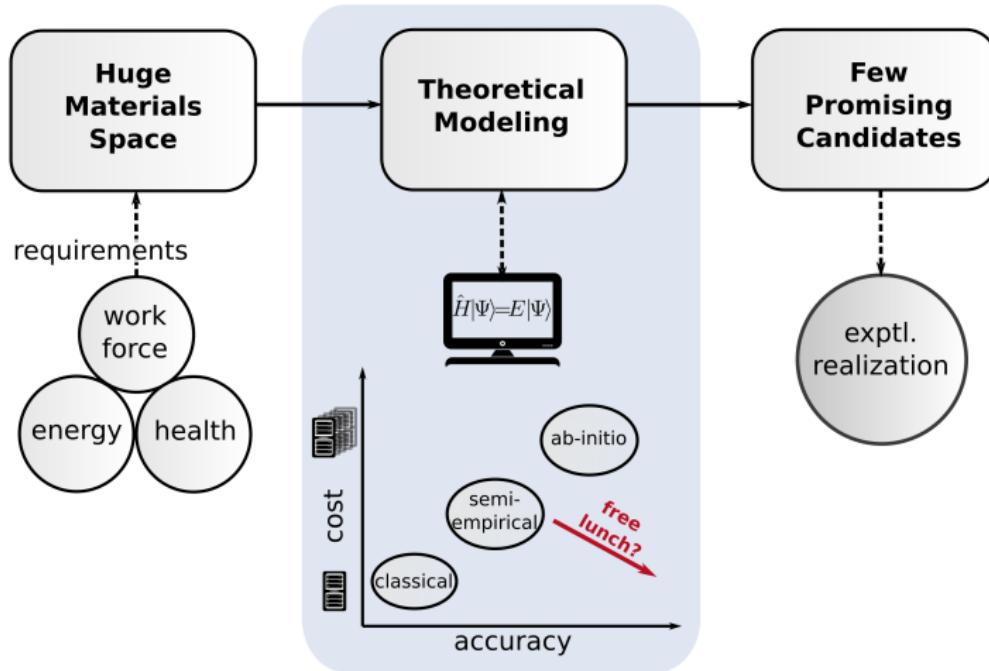
Outline of talk

- 1 Introduction
- 2 Benchmark quality methods
- 3 Quantum Monte-Carlo for molecular crystals
- 4 Quantum Monte-Carlo for water physisorption
- 5 Conclusions

Material discovery can employ computational models



Material discovery can employ computational models



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Why is benchmarking important?

- every computed or measured property needs an error bar
- possible origins: random and systematic errors
- random errors from statistics, e.g. molecular dynamics
→ straight-forward to converge
- systematic errors from intrinsic error of theoretical model
→ nontrivial to estimate

Benchmarking from economical perspective

- (1)** determine a standardized quality measure
- (2)** analyze what and where improvements are called for
- (3)** use this information to improve performance

Experimental

- final goal should be to describe experimental conditions
- possible for complex systems
- measurement uncertainty
- no separation of effects

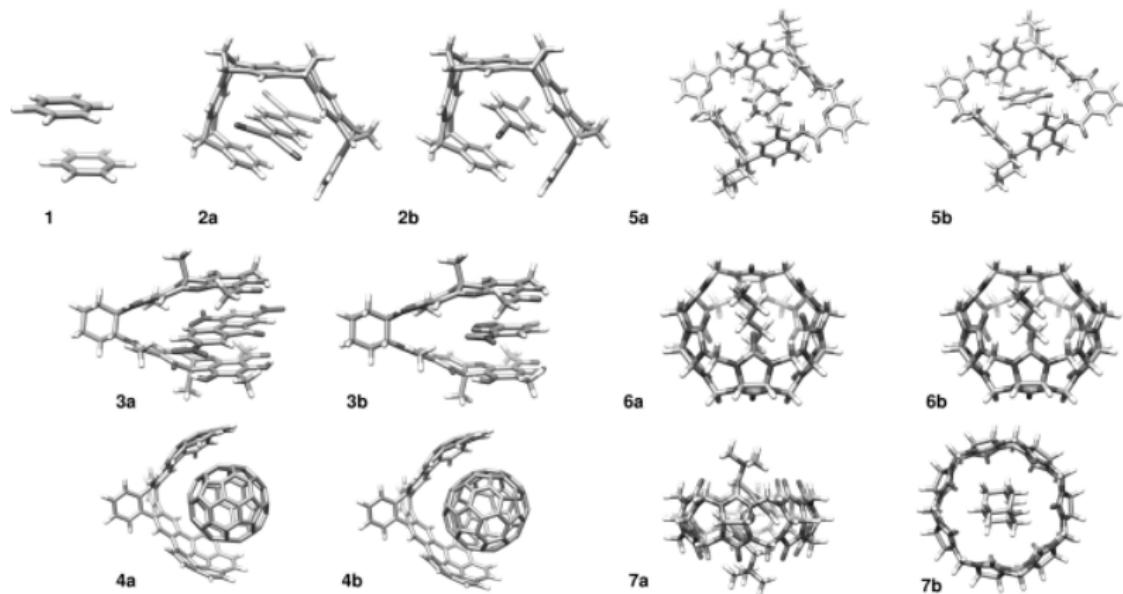
Theoretical

- straight-forward comparison
- computational cost affordable?
- only possible for 'small enough' systems
- uncertainty of reference method

Semi-experimental:

Combine measured properties with theoretical evaluation of secondary effects

S12L: Supramolecular complexes



- free energy of association measured in solution

S12L: Supramolecular complexes

- separate energy contributions

$$\Delta G_a = \Delta E + \Delta G_{\text{RRHO}}(T) + \Delta \delta G_{\text{solv}}$$

- estimate $G_{\text{RRHO}}(T)$ from harmonic frequency estimate

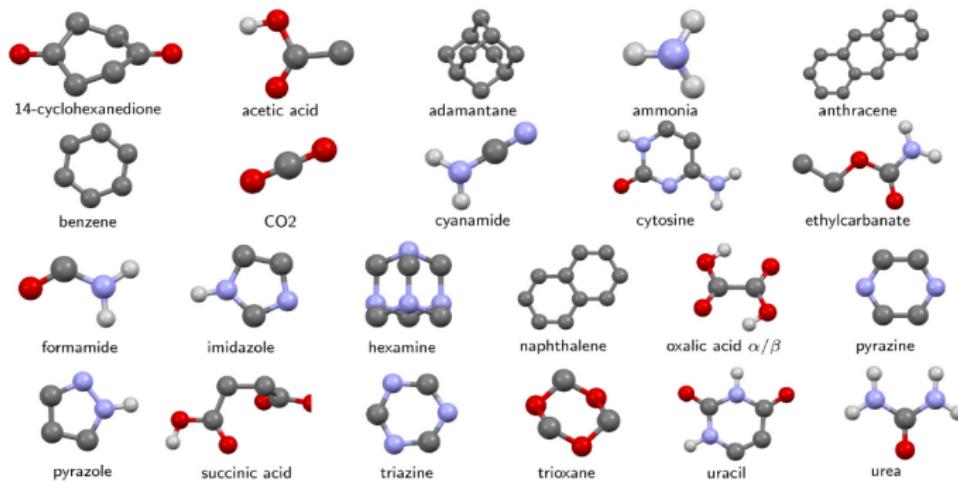
$$\Delta G_{\text{RRHO}}(T) = \sum_p \frac{\hbar\omega_p}{2} + k_B T \sum_p \left[\ln \left(1 - e^{-\frac{\hbar\omega_p}{k_B T}} \right) \right],$$

→ error of about 3%

- estimate $\Delta \delta G_{\text{solv}}$ from implicit solvation model
 - error of about 5-10%
- remove contributions yielding semi-experimental ΔE
 - error of about 5-10% $\equiv 100 \text{ meV}$

[¹] S. Grimme, *Chem. Eur. J.*, **18**, 9955 (2012)

X23: Molecular crystals

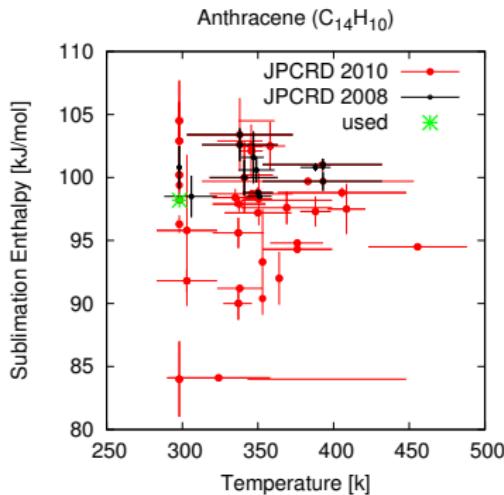
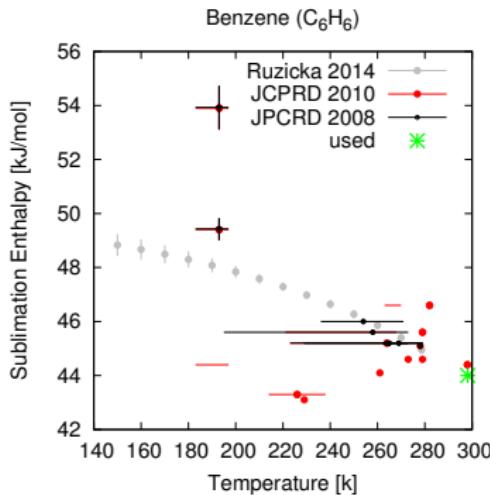


- molecular crystals of small molecules, emphasis on vdW bonding

[2] A. Otero-de-la-Roza, E. R. Johnson, *J. Chem. Phys.* **137**, 054103 (2012).

[3] A. M. Reilly, A. Tkatchenko, *J. Chem. Phys.* **139**, 024705 (2013)

Significant uncertainties in measured sublimation enthalpies



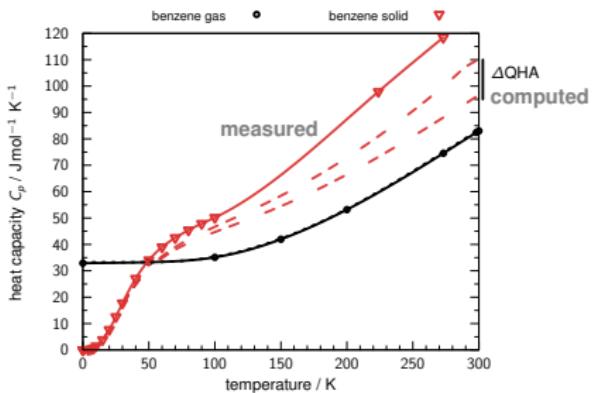
- measure H_{sub} , statistical survey yields error of $\sim 50 \text{ meV}$

[4] W. Acree, J. S. Chickos, *J. Phys. Chem. Ref. Data* **39**, 043101 (2010).

X23/ICE10: Molecular crystals

$$E_{\text{latt}} = -\Delta H_{\text{sub}}(T) + \Delta E_{\text{ZPVE}} + \int_0^T \Delta C_p(T') \, dT'$$

- measured sublimation enthalpy H_{sub}
- zero-point energy from phonon modes
- thermal contribution from heat capacities (measured or computed)



→ error of about 5% $\equiv 50 \text{ meV}$

Quantum Monte-Carlo in a nutshell: A scalable high-level method

Fixed-node diffusion Monte-Carlo

1) Enforce nodal surface of Fermions

$$\Gamma = \{\mathbf{R}; |\Psi_T\rangle = 0\}$$

2) Walkers in configuration space

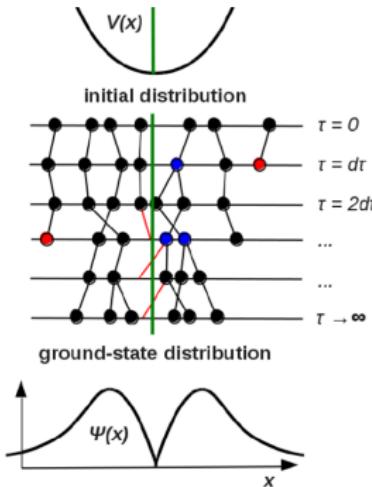
$$|\Psi_T(\mathbf{R}, \tau)\rangle = \text{hist} \left[\sum \delta(\mathbf{R} - \mathbf{R}_i(\tau)) \right]$$

3) Diffusion in imaginary time

$$\partial_\tau |\Psi_T(\mathbf{R}, \tau)\rangle = \left[\frac{1}{2} \nabla_{\mathbf{R}}^2 - (V - E_T) \right] |\Psi_T(\mathbf{R}, \tau)\rangle$$

4) Projection to exact ground state

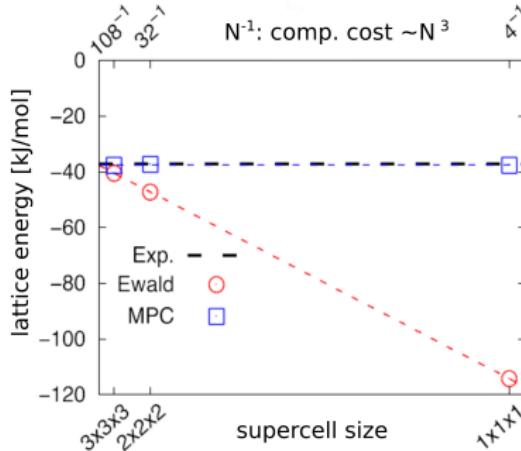
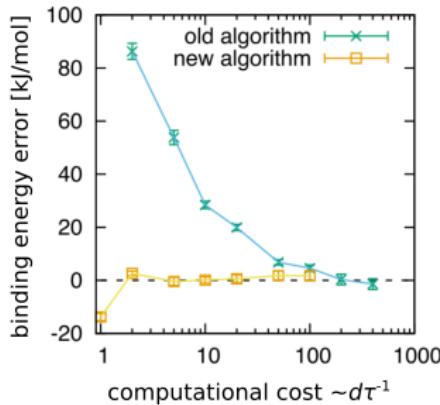
$$|\Psi_0(\mathbf{R})\rangle = \lim_{\tau \rightarrow \infty} \exp[-\tau(\hat{H} - E_T)] |\Psi_T(\mathbf{R}, \tau)\rangle$$



- exact within Γ and $d\tau \rightarrow 0$ and $\tau \rightarrow \infty$
- non-local part of pseudopotentials impact electron correlation

[5] M. Ďubecký, L. Mitas, P. Jurečka, *Chem. Rev.* **116**, 5188 (2016)

New DMC algorithm leads to substantial speed up



- new size-consistent implementation reduces $\Delta\tau$ error drastically^[6]
- Model periodic Coulomb for finite size correction^[7]

[6] A. Zen, S. Sorella, M. J. Gillan, A. Michaelides, D. Alf  , *Phys. Rev. B* **93**, 241118(R) (2016).

[7] L. M. Fraser, W. M. C. Foulkes, G. Rajagopal, R. J. Needs, S. D. Kenny, A. J. Williamson, *Phys. Rev. B* **53**, 1814 (1996).

Methods with 'benchmark quality' for noncovalent interactions



CCSD(T)

- + 'gold standard' of quantum chemistry
- + linear scaling variants
- pair-screening thresholds
- slow basis set convergence
- non-trivial to parallelize
(shared memory: 100 GB/core)

DMC

- + exact projection to ground state
- + N^3 scaling, [-] huge prefactor
- + fast basis set convergence
- + scaleable to HPCs
- stochastic error
- uncertainties from FN and non-local pseudopotentials

No benchmark quality: MP2, MP3, RPA, CCSD, CCSD(T)/cc-pVTZ

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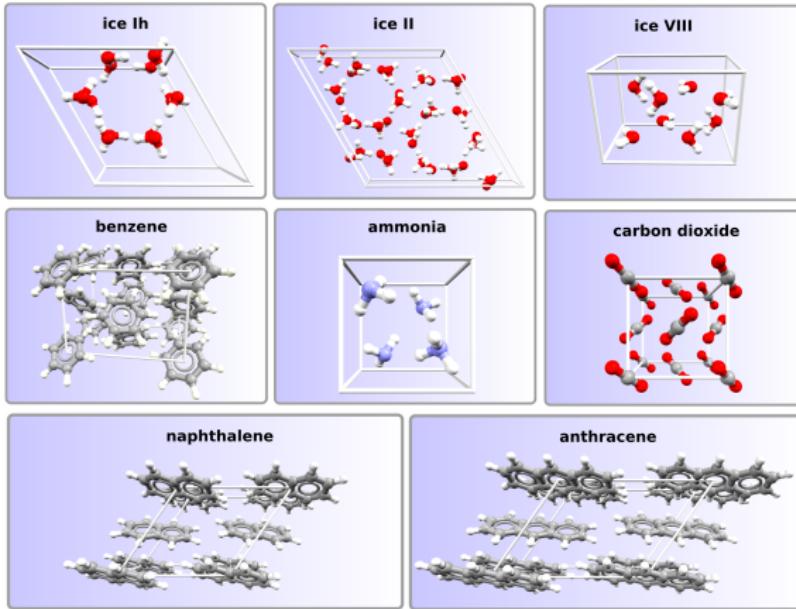
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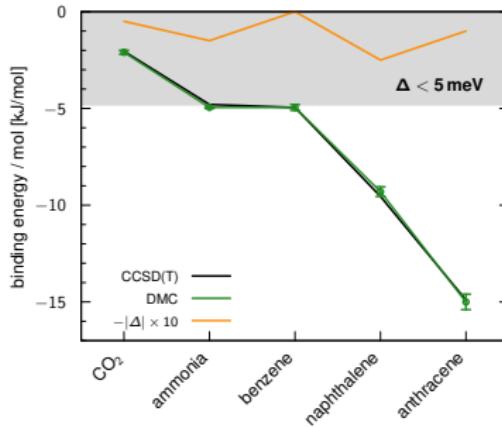
Diverse interactions in test crystals



- strong H-bonds, vdW of saturated and unsaturated molecules
- problematic for all readily applicable methods (DFA-DISP, MP2)

Close agreement between CCSD(T) and DMC on dimers

- dimers extracted from crystal structure
- CCSD(T)-F12 / cc-pV(T,Q)Z-F12 for CO₂ and NH₃
- L-DLPNO-CCSD(T) / cc-pV(T,Q)Z (tight PNO settings)^[8]
- largest deviation of 3 meV

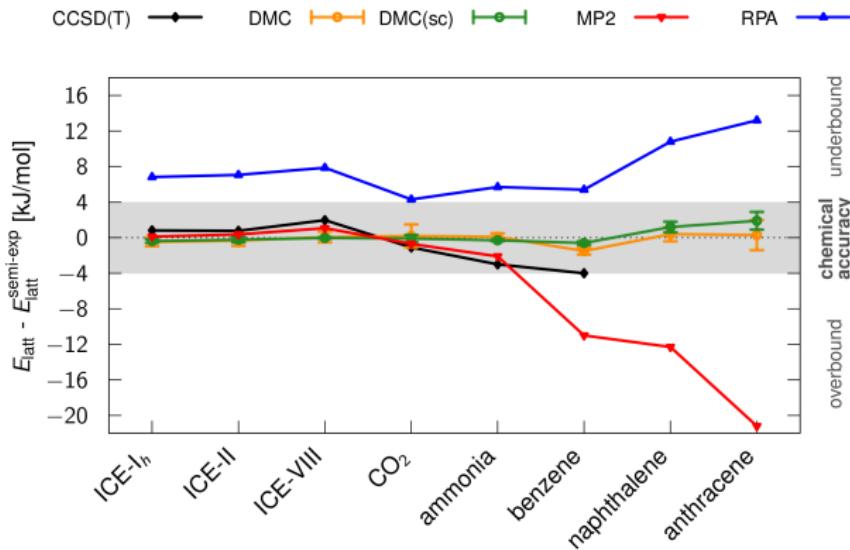


→ good agreement between different high-level methods^[9]

^[8] C. Ripplinger, B. Sandhoefer, A. Hansen, F. Neese, *J. Chem. Phys.* **139**, 134101 (2013).

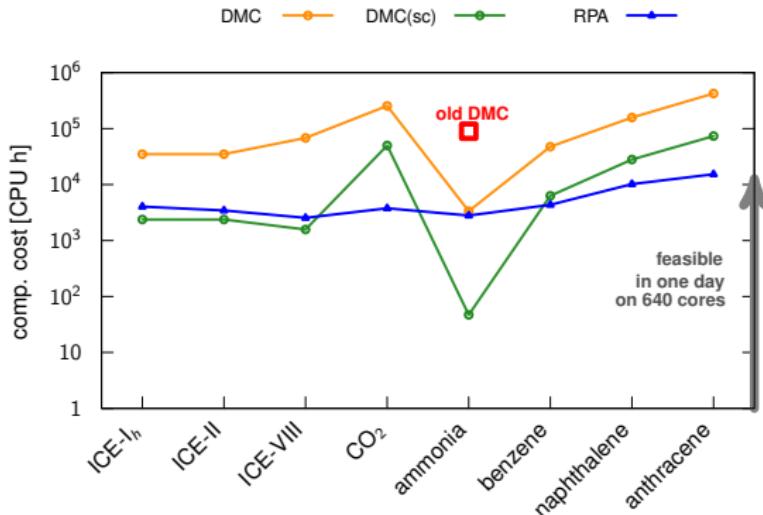
^[9] Y. S. Al-Hamdani, M. Rossi, D. Alfè, T. Tsatsoulis, B. Ramberger, JGB, A. Zen, G. Kresse, A. Grüneis, A. Tkatchenko, A. Michaelides *J. Chem. Phys.* **147**, 044710 (2017).

QMC delivers (sub-) chemical accuracy for all tested systems



- excellent agreement with experiment and CCSD(T)
- uncertainty in $H_{\text{sub}}^{\text{exp}}$ probably larger than DMC errors

QMC is feasible within one day on standard computer cluster



- up to three orders of magnitude speed-up compared to best DMC practice two years ago
- significance will extend to all classes of systems^[10]

^[10] A. Zen, JGB, J. Klimeš, A. Tkatchenko, D. Alfè, A. Michaelides, Proc. Natl. Acad. Sci. U.S.A., 115, 1724 (2018).

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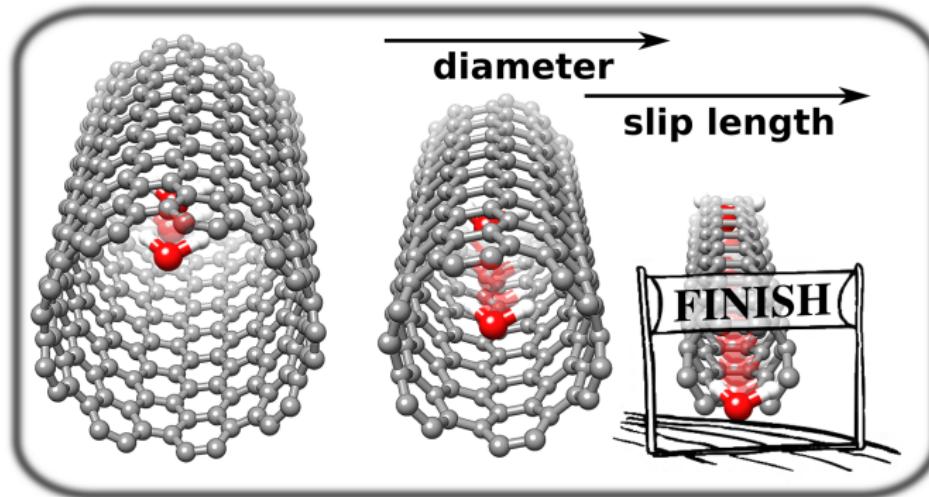
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Relevance of water graphene interaction?

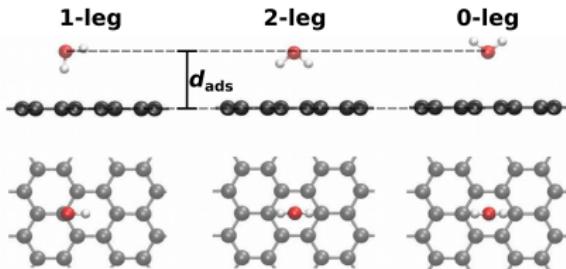


- graphite/graphene used as water filters
- water monomer adsorbed on graphene as fundamental vdW system

[11] E. Secchi, S. Marbach, A. Niguels, D. Stein, A. Siria, L. Bocquet, *Nature* **537**, 210 (2016).

[12] R. Joshi, P. Carbone, F. Wang, V. Kravets, Y. Su, I. Grigorieva, H. Wu, A. Geim, R. Nair, *Science* **343**, 752 (2014).

Severe discrepancy of literature values



DMC

−70 meV^[13]

i-CCSD(T)

−135 meV^[14]

SAPT, DFT/CC

−130 meV^[15,16]

- differences of > 60 meV not acceptable

^[13] J. Ma, A. Michaelides, D. Alfè, L. Schimka, G. Kresse, E. Wang, *Phys. Rev. B* **84**, 033402 (2011).

^[14] E. Voloshina, D. Usvyat, M. Schütz, Y. Dedkov, B. Paulus, *Phys. Chem. Chem. Phys.* **13**, 12041 (2011).

^[15] G. R. Jenness, O. Karalti, K. D. Jordan, *PCCP* **12**, 6375 (2010), ^[16] M. Rubeš, et al. *JPC C* **113**, 8412 (2009).

L-CCSD(T) setup for molecular clusters



- tight pair-thresholds and non canonical triples (T0)
- BB-CP corrected binding energies and extrapolation

$$E_{\text{corr}}[\text{CC/CSB}] = E_{\text{corr}}(\text{CC/QZ}) \times \frac{E_{\text{corr}}(\text{RPA/CBS})}{E_{\text{corr}}(\text{RPA/QZ})}$$

{non-published data}

Water@graphene binding energies



{non-published data}

Extrapolation of substrate size



{non-published data}

- convergence with substrate size very slow
→ extrapolations using benzene, coronene data unreliable

Redistribution of charge density strongly dependent on adsorption motif

{non-published data}

- convergence with substrate size very slow
→ extrapolations using benzene, coronene data unreliable

Discrepancies of literature values resolved

DMC

–70 meV^[13]

- stochastic error
- finite-size correction

i-CCSD(T)

–135 meV^[14]

- un converged basis set expansion

SAPT, DFT/CC

–130 meV^[15,16]

- unreliable extrapolation

revised DMC

–99 ± 5 meV

^[13] J. Ma, A. Michaelides, D. Alfè, L. Schimka, G. Kresse, E. Wang, *Phys. Rev. B* **84**, 033402 (2011).

^[14] E. Voloshina, D. Usvyat, M. Schütz, Y. Dedkov, B. Paulus, *Phys. Chem. Chem. Phys.* **13**, 12041 (2011).

^[15] G. R. Jenness, O. Karalti, K. D. Jordan, *PCCP* **12**, 6375 (2010), ^[16] M. Rubeš, et al. *JPC C* **113**, 8412 (2009).

Benchmarking of density functional approximations



{non-published data}

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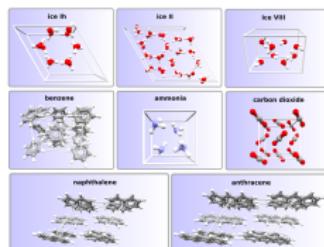
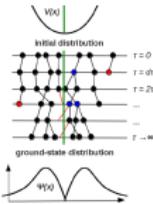
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Summary

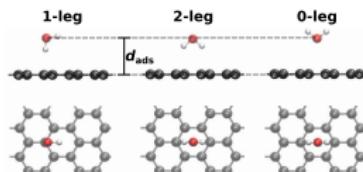
Conclusions

- DMC delivers (sub-) chemical accuracy with three orders of magnitude speed up
- chemically accurate lattice energies
- revised reference value for water@graphene



Outlook

- improved DMC for even larger systems
- use improved methodologies in CSP
- NNP training within CSP algorithms



Acknowledgements



PhD / Postdocs



Andrea Zen



Michaelides (UCL)

Principal investigator



Alf  (UCL)



Tkatchenko
(Luxembourg)



Klimes (Prague)



Eike Caldeweyer



Grimme (Bonn)



Perdew (Temple)



Kresse (Vienna)



Behler (G ttingen)

Industry



Rebecca Sure (BASF)



Luca Iuzzolino



Price (UCL)



Fernandez-Alonso
(ISIS)



Civalleri (Turin)



Maschio (Turin)



Antreas Afantitis
(NovaMechanics)



Alexander von Humboldt
Stiftung/Foundation



Acknowledgements



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Antreas Afantitis
(NovaMechanics)

Alexander von Humboldt
Stiftung/Foundation

THE
THOMAS YOUNG
CENTRE
FOR
CROSS-DISCIPLINARY
SCIENCE

Thanks

cpresso

Key references

■ QMC for molecular crystals:

JGB, A. Zen, A. Michaelides, G. Kresse, and D. Alfè, in preparation

A. Zen, JGB, J. Klimeš, A. Tkatchenko, D. Alfè, A. Michaelides,
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JGB, J. E. Bates, J. Sun, J. P. Perdew, *Phys. Rev. B*, **94**, 115144 (2016).

JGB, C. Bannwarth, A. Hansen, S. Grimme *J. Chem. Phys.*, **148**, 64104 (2018).

■ Crystal structure prediction:

S. L. Price, JGB, *Molecular Crystal Structure Prediction*, G. DiLabio, A. Otero-de-la-Roza, Eds., Elsevier Australia, 336-363 (2017).

■ website: **gerit-brandenburg.de**