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## **EDITORIAL**

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## Benchmark experiments for numerical quantum chemistry

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Driven by impressive advances in both hard- and software as well as a steady flow of new concepts, quantum chemical predictions go from strength to strength. Nonetheless, the benefits from this progress will starkly depend on the quality of data we provide or are provided with. Quantum chemical methods can replace and generate novel research workflows, but these will only be useful with a clear knowledge of their accuracy and scope. Benchmarking of electronic structure methods has been a common practice ever since the 1990s, partly as a response to the plethora of density functional methods introduced every year. With a somewhat bloated toolbox available to the computational chemist, it became more relevant to know which is the right tool, rather than to add further alternatives into the mix. This has also translated into the nagging question: "Which functional should I use?" Many authors have answered the call and have compared time and time again different DFT methods for different properties, sometimes as a main focus, others as a side task to justify the choice of theory level. Whatever the case may be, the shift clearly moved from questioning whether

one can compute it to how one should compute it.

Most of the benchmarking practices to date tend to focus on comparing relative electronic energies, chiefly between the methods in test and a reference (expensive) quantum chemical approach (e.g., the CCSD(T) gold standard). This is a straightforward computational protocol, but fails to cover the complexity involved in a large number of chemical observables. Experiment remains the ultimate adjudicator about the suitability of theoretical models and protocols. This usually requires combined solutions for the electronic structure and for the dynamics of the nuclei. To avoid misleading error compensation in theory and misleading experimental references, great care in the design of benchmark experiments is required. This includes sufficiently large databases, multi-experimental cross-validations, and the organization of blind challenges for unbiased predictions. On the theory side, it is crucial to have approaches at hand which minimize error in either electronic or nuclear dynamics problems, such that rigorous lessons can be learned for the other component of the numerical challenge. On the experimental side, gasphase experiments, often at low temperature, obtained by spectroscopic techniques with high accuracy, are the natural comparison with theoretical data. In this respect, supersonic expansions or cryogenic ion traps allow studying cold isolated molecules or weakly bound complexes with unprecedented precision. High-resolution rovibrational or rotational spectroscopy are available for small systems, while larger systems are often studied at vibrational resolution. Besides these experiments providing structures and nuclear motion information, crystallographic data (https://doi.org/10.1039/D2CP04098K), luminescence (https://doi.org/10.1039/ D2CP01623K), mass spectrometry experiments such as photon- or collision-induced dissociation or ion mobility (https://doi.org/ 10.1039/D2CP01414A), NMR spectroscopy (https://doi.org/10.1039/D2CP04092A or https://doi.org/10.1039/D2CP03992C) or X-ray scattering (https://doi.org/10.1039/ D2CP02933B) give valuable information on electronic effects. A wealth of experimental methods allow studying reactive systems in solution (https://doi.org/ 10.1039/D2CP03937K) such as electrochemistry or luminescence, and provide new areas of benchmarking.

This themed collection, Benchmark Experiments for Numerical Quantum Chemistry, of more than 40 articles (about one third being classified as hot articles) addresses different aspects of this endeavour, bringing quantum theory and experiment together at suitable meeting points, for the mutual benefit of both communities. Two perspectives in the field of non-covalent interactions address the theoretical advances in fully coupled, numerically exact rovibrational states (https://doi.org/10.1039/D2CP04005K) and

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encompass the description of quadrupole how to organize a particular blind challenge on hydrate vibrational shifts from the coupling, the proper description of conforexperimental side (https://doi.org/10.1039/ mational flexibility at an acceptable com-D2CP01119K). The latter also invites less putational cost, and the application of exact quantum approaches and machine rotational spectroscopy for chirality analearning (for the outcome, see https://doi. lysis. Describing the coupling between org/10.1039/D3CP01216F), to be put under rotation and vibration, especially large scrutiny in later rounds. For somewhat amplitude motion, is still a grand challenge more rigid molecular systems, a review (https://doi.org/10.1039/D2CP03897H). Size (https://doi.org/10.1039/D2CP04706C) and conformer-selective characterisation of demonstrates how closely the Bornthe excited-state deactivation pathways or Oppenheimer concept of molecular equiprocesses involved in host-guest interaclibrium structure and the experimentally tions or molecular recognition are used observable rotational constants can be for assessing the validity of excited-state brought together. Formic acid is reviewed descriptions or complex potential-energy as an elementary example for the vibrasurfaces commonly used by experimentalists. (https://doi.org/10.1039/D2CP04570B, tional characterization of a bistable molecule (https://doi.org/10.1039/D2CP04417J). https://doi.org/10.1039/D2CP03796C, Finally, a tutorial review addresses how https://doi.org/10.1039/D2CP03953B, reactivity scales help in structuring https://doi.org/10.1039/D2CP01414A, and overcoming challenges in kinetics https://doi.org/10.1039/D2CP03110H). benchmarking (https://doi.org/10.1039/ Several papers extend the applicability D2CP03937K). The remaining articles in range for molecular benchmarking, to the themed collection elaborate on simiradical and biradical complexes (https:// lar problems while expanding to other doi.org/10.1039/D2CP04092A, https:// areas. On the theory side, this includes doi.org/10.1039/D3CP01156A, https:// doi.org/10.1039/D2CP04101D, https:// uncertainty quantification in rolling benchmarks (https://doi.org/10.1039/ doi.org/10.1039/D2CP03889G), to non-D2CP01725C) and nuclear quantum standard electronic transitions (https://doi. effects in reaction kinetics (https://doi. org/10.1039/D3CP00160A, https://doi.org/ org/10.1039/D2CP03809A) as well as multi-10.1039/D2CP01623K) or to X-ray scatterlevel schemes for larger system sizes ing off small molecules (https://doi.org/ (https://doi.org/10.1039/D2CP05056K). In 10.1039/D2CP02933B). Other contributhe gas phase as a natural benchmarking tions revisit previous benchmarking efforts, habitat, high resolution rotational spectrosuch as for intermolecular balances (https:// scopy is certainly among the toughest doi.org/10.1039/D2CP03907A, https://doi. experiments to be met by theory and it org/10.1039/D2CP05141A), formic acid comgoes far beyond just providing rotatioplexes (https://doi.org/10.1039/D2CP03893E, nal constants (https://doi.org/10.1039/ https://doi.org/10.1039/D2CP04176F) or for D2CP05774C, https://doi.org/10.1039/ micro-hydration (https://doi.org/10.1039/ D2CP04825F, https://doi.org/10.1039/ D2CP04174J), and even for elementary dia-D2CP04067K, https://doi.org/10.1039/ tomic molecules (https://doi.org/10.1039/ D2CP03897H), often with several research D2CP03964H). New benchmark data sets

are presented and used for practical pur-

poses (https://doi.org/10.1039/D2CP04049B,

https://doi.org/10.1039/D2CP03992C,

https://doi.org/10.1039/D2CP04052B),

with data sizes up to several million (https://doi.org/10.1039/D2CP03966D). The extension of studies towards complex systems has resulted in a diversity in the problems tackled experimentally, such as the study of the crystalline phase or complex protein environments (https:// doi.org/10.1039/D2CP04098K. https://doi. org/10.1039/D2CP00184E). The studies are here extended to electrostatic properties (https://doi.org/10.1039/D2CP04052B), or to metal surface adsorption (https://doi. org/10.1039/D2CP04398J). While biomolecular docking processes (https://doi.org/ 10.1039/D2CP04671G) represent a relatively mature area of benchmarking practice, AI-based approaches are more recent (https://doi.org/10.1039/D3CP01216F). The ultimate goal must be to bring these different areas together to better assert the robustness of methods and avoid depending on error cancellation (https://doi.org/ 10.1039/D2CP04098K).

Benchmarking should be a continuous activity, keeping our theoretical models grounded to the highest standard: empirical validation. It requires an incessant review and expansion of references, a critical eye to mismatches and shortcomings plus the insight to propose new theories/approximations which effectively overcome the latter. Bringing different communities together generates common data points to everyone involved, fostering interdisciplinarity. It can also build moments of respite away from the individualistic "publish-or-perish" culture, by sharing experience and data for the greater good.

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groups joining forces (https://doi.org/

10.1039/D2CP03962A). The open questions

https://doi.org/

https://doi.org/

10.1039/D2CP04663F,

10.1039/D2CP04060C,