



Cite this: *Phys. Chem. Chem. Phys.*,
2023, 25, 6562

The application of QM/MM simulations in heterogeneous catalysis

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The QM/MM simulation method is provenly efficient for the simulation of biological systems, where an interplay of extensive environment and delicate local interactions drives a process of interest through a funnel on a complex energy landscape. Recent advances in quantum chemistry and force-field methods present opportunities for the adoption of QM/MM to simulate heterogeneous catalytic processes, and their related systems, where similar intricacies exist on the energy landscape. Herein, the fundamental theoretical considerations for performing QM/MM simulations, and the practical considerations for setting up QM/MM simulations of catalytic systems, are introduced; then, areas of heterogeneous catalysis are explored where QM/MM methods have been most fruitfully applied. The discussion includes simulations performed for adsorption processes in solvent at metallic interfaces, reaction mechanisms within zeolitic systems, nanoparticles, and defect chemistry within ionic solids. We conclude with a perspective on the current state of the field and areas where future opportunities for development and application exist.

Received 28th September 2022,
Accepted 20th December 2022

DOI: 10.1039/d2cp04537k

rsc.li/pccp

1 Introduction

QM/MM (Quantum mechanics/molecular mechanics) techniques are a key methodology for computational simulations of extended systems.¹ The core of this technique lies in the partition of the system into separate regions, with quantum mechanics applied to a central region of interest, and classical mechanics applied to the environment. The partitioning allows for favourable scaling when performing simulations of large systems, where high accuracy methods such as density functional theory (DFT) or post-Hartree Fock can calculate the energetics of salient features for the system under study; meanwhile, the electrostatics of the extended system are included through classical empirical force-fields, thereby significantly reducing the computational workload compared to optimising the electronic structure of the entire extended system.²

Both cluster and periodic QM/MM models have been extensively used for problems in structural biology;^{3,4} however, multiscale approaches are also gaining popularity in heterogeneous catalysis, which is motivated by the ability of QM/MM to calculate system observables with greater speed and accuracy compared to fully *ab initio* simulations. For example, in systems with strong concentration effects (*i.e.*, in the simulation of charge defects, or the adsorption of molecules with strong lateral interactions⁵), an aperiodic embedded-cluster QM/MM model can be used to calculate the energetics at the dilute limit,

while the influence of the wider system is abstracted using a larger classical embedding representation. In contrast, a periodic QM simulation would require an infeasibly large simulation cell to correctly model defects in the dilute limit, or compromise accuracy by allowing interactions between charge defects in uncharacteristically close proximity to one another.⁶

The QM/MM method is also ideal for calculating salient system information with high accuracy methods, while representing less chemically relevant regions with computationally cheap classical simulations. This benefit has been widely employed for simulations of periodic and embedded-cluster QM/MM models of zeolites, for example detailing fundamental reactions in the MTH process at the H-ZSM-5 zeolite.⁷ For such systems, modelling of the entire unit cell of the zeolite was avoided by limiting the QM simulation to the adsorbate and a small segment of the zeolite, which emulates chemistry sought in the full periodic simulation. The reduction of the size of the QM calculation in turn enables application of post-Hartree Fock methods or higher-level density functional theory (DFT), *e.g.* including contributions of exact exchange, allowing for greater accuracy when describing the energetics of adsorption and reactivity.

In addition to reducing the cost of conventional geometry and single point energy evaluations, QM/MM enables the evaluation of free energy changes in dynamic solvent environments. This has been achieved by representing the solvent with a classical force-field, paired with techniques such as Thermodynamic Integration (TI)^{8,9} and Free Energy Perturbation

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(FEP).^{10,11} The schemes use QM/MM to calculate the free energy of adsorption in the aqueous environment, which provides a realistic comparison to experimental studies in aqueous solvent. Multiscale solvation schemes drastically reduce the cost of evaluating the dynamic solvent environment, which would otherwise be infeasible with a full *ab initio* simulation. These approaches have a promising future in describing liquid-phase catalytic processes at surfaces and nanoparticles.

In this review, the application of QM/MM methods to a range of challenges in heterogeneous catalysis are presented, demonstrating the potential of the approach. To aid our discussion, the fundamental theory of QM/MM and important considerations for configuring simulations are first introduced. Then, applications of QM/MM are presented in domains of solvent simulations, zeolite studies, metallic surfaces and nanoparticles, and the simulation of defects in ionic solids. To finalise our discussion, emergent and potentially impactful methodologies are discussed, as well as developments necessary to improve the uptake and ease of use of the QM/MM methodology in heterogeneous catalysis.

2 Basics of QM/MM

2.1 Subtractive and additive coupling schemes

As outlined, QM/MM calculations require partitioning of a model such that different methods can be applied to each sub-partition (Fig. 1). The expression for the QM/MM total energy term is typically presented *via* two dominant schemes; additive and subtractive.

Subtractive schemes are a conceptually simple method wherein three separate calculations are performed for the central region of interest (1) embedded in an extended representation of the environment (2),

$$E_{\text{QM/MM}}^{\text{sub}} = E_{\text{QM}}^1 + (E_{\text{MM}}^{12} - E_{\text{MM}}^1), \quad (1)$$

where $E_{\text{QM/MM}}^{\text{sub}}$ is the total energy for the QM/MM system, E_{QM}^1 is the QM energy for region 1, E_{MM}^{12} is the total MM energy of the whole system, *i.e.* regions 1 and 2, and E_{MM}^1 is the MM energy of region 1 calculated with MM. Examples of this scheme include the implementation into the ONIOM ('Our own *n*-layered integrated molecular orbital and molecular mechanics') framework.^{12,13}

In contrast, additive schemes calculate the inner and outer regions on the level of QM and MM, respectively, and the interaction between the two regions is treated through an additional coupling term

$$E_{\text{QM/MM}}^{\text{add}} = E_{\text{QM}}^1 + E_{\text{MM}}^2 + E_{\text{couple}}^{12}, \quad (2)$$

where E_{couple}^{12} is a coupling term describing the interactions between the two regions. In effect, this requires only two calculations, while the coupling scheme allows for the selection of different terms from the MM region to include within the Hamiltonian of the QM region.^{2,14}

Although the two approaches are in principle mathematically equivalent under conditions when the QM and MM potential energy landscape are commensurate,¹⁴ in reality challenges occur due to the difficulty in reaching equivalence in the landscapes. The challenges primarily concern the treatment of the QM/MM boundary, which we will discuss in depth in Section 2.3.

2.2 Embedding schemes

Embedding methods can be further categorised in their treatment of coupling between the QM and MM regions (*i.e.*, mechanical, electrostatic and polarisable). Mechanical coupling simply considers interactions between the internal and external region through MM structural parameters. Although

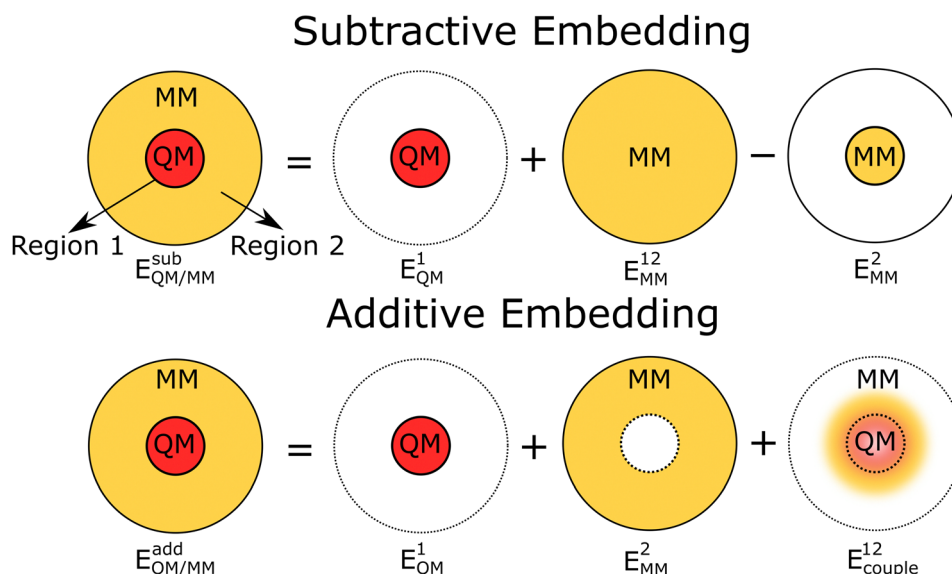


Fig. 1 A simple diagrammatic representation of the subtractive and additive QM/MM schemes. The former is carried out using three separate calculations, corresponding to a QM calculation of region 1, a MM calculation for the combined region 1 & 2, and another MM calculation for the inner region (eqn (1)). The additive scheme is carried out using two calculations, of the inner and outer regions using QM and MM, respectively, with the coupling interactions between the QM and MM regions introduced by a separate coupling term (eqn (2)).



conceptually simple, mechanical coupling introduces significant inaccuracies unless a specific force-field is generated that commensurately describes the interactions between the QM and MM regions. These inaccuracies are exacerbated when significant changes in electron density or geometry occur, which requires updated force-field terms to correctly describe the altered structure.⁴

In contrast, electrostatic embedding includes the MM charges within the one-electron Hamiltonian, introducing the influence of the embedding charges into the self-consistency cycle; polarisable schemes then introduce coupling of the MM region with the charge density of the QM region, thereby dynamically altering the dipoles of atomic centers in the MM region depending on the quantum mechanical charge. Within the additive QUASI scheme, this is implemented through the polarisable shell model for the MM atoms.^{2,15,16} Polarisable embedding has also been implemented to the subtractive ONIOM scheme under the ONIOM(QM:MM)-PE framework, which is employed to calculate the interaction between the classical charges and the dielectric continuum of an implicit solvent environment.¹⁷

QM/MM calculations are feasible with both open and periodic boundary conditions.^{18–22} Embedded-cluster QM/MM models commonly have open boundary conditions, which allows for the facile calculation of charge without periodic interactions; charged models under periodic boundary conditions interact with their periodic replicas, which at high concentrations can distort the desired energetics energy from the dilute limit.²³ The open boundary conditions also provide direct access to an absolute energy reference (*i.e.*, the vacuum level), allowing alignment of the band edges of any materials;²⁴ in contrast, the reference energy level of simulations of 3D periodic systems is ill-defined due to the arbitrary choice of the average background potential (*i.e.*, where $\mathbf{G} \rightarrow 0$).^{25,26} It is noted that 2D periodic systems have access to a reference energy level, where methods such as the Coulomb cut-off may yield a definite vacuum energy level;²⁷ however, periodic 2D models do not accurately represent charged systems by default.^{28,29} Discussion of modern approaches for correcting the periodic representation of charged systems are summarised by Walsh.³⁰ It should also be noted that QM/MM has also been formulated for periodic boundary conditions, such as the formulation of QM-Pot.²⁰ Alternative implementations of periodic QM/MM *via* the Particle Mesh Ewald (PME) method are demonstrably useful for representing the long range electrostatics of the dynamic solvent environment without introducing artificial cut-offs.^{18,19}

Beyond the atomistic QM and MM representations, long range polarisation of an extended aperiodic bulk system may be treated through a simple additive term known as the Jost correction.^{31,32}

$$E_{\text{Jost}} = -\frac{Q^2}{2R} \left(1 - \frac{1}{\epsilon}\right), \quad (3)$$

where Q is the charge of the QM region, R is the radius of the cluster and ϵ is the bulk dielectric constant of the material. The correction holds well assuming polarisation occurs *via* long range electrostatics with isotropic dielectric permittivity.

A notable drawback in the QM/MM methodology is the confinement of electronic states within the QM region. Although the localised states of the valence electrons are well-represented, the diffuse conduction states are shifted significantly upwards in energy.³³ As a consequence, studies that require the conduction band as an energy reference (such as those in Section 3.4) often elect to use experimental values of the conduction band minimum (CBM) for the bulk.^{34,35}

2.3 QM/MM boundary and link atoms

At the boundary of the QM/MM interface, the electron density of the QM atoms can spill over into the positively charged MM region. This is especially problematic for covalent compounds, where atoms with incomplete valences must be terminated to prevent nonphysical radicals forming at the boundary.

A common scheme for completing the valence shell of boundary atoms is the link atom approach,³⁶ which involves placing a capping QM atom along the bond length between the boundary QM and MM atoms, usually in the form of a hydrogen atom; however, double-counting of the energetic terms associated with the link atom must be accounted.

The subtractive scheme of eqn (1) naturally cancels the discrepancies introduced by the link atom; both the QM and MM calculations for the inner region 1 contain energy contributions for the H atom, which are subtracted from one another. Provided the force-field correctly emulates the potential energy surface for the QM/MM region boundaries, the difference in energies calculated for the classical link atom in E_{MM}^1 and the original boundary atom in E_{MM}^{12} may be treated as an extrapolative correction to the quantum mechanical energy from the artificial link to the correct boundary atom.^{12,14} However, an improperly parameterised force-field – which incorrectly describes the interaction across the boundary – may lead to discontinuities across the potential energy surface.⁴

The additive scheme in its strict form (eqn (2)) provides no such correction for link atoms; however, eqn (2) may be adapted to include elements of the subtractive expression,²

$$E_{\text{QM/MM}}^{\text{add,link}} = E_{\text{QM}}^1 + E_{\text{MM}}^2 + E_{\text{couple}}^{12} - E_{\text{MM}}^{\text{link}}, \quad (4)$$

where $E_{\text{MM}}^{\text{link}}$ is the MM energy of the isolated link atom. As with the subtractive case, a well-parameterised force-field is necessary.² Alternatively, the MM energy contributions (stretching, angle distortion, torsion, *etc.*) between the MM and QM atom may simply be deleted to eliminate the double-counting in the energetic coupling term. As a result, the energy of the boundary MM atom is approximated by using the distortion energy introduced by the link atom alone.²

Ionic species at the boundary may be represented using Gaussian-based effective core potentials (ECPs) or numerical pseudopotentials (PSPs), which emulate the presence of a boundary QM atom without introducing further QM species. ECPs are widely available from the literature, though typically optimised for molecular systems, due to the prevalence of Gaussian basis sets for such calculations;³⁷ PSPs, in contrast, can be generated and tested for solid state systems using widely available frameworks.³⁸ For PSPs, the local potential represents



the long-range electrostatic features of the atom, whilst the nonlocal expressions, such as those in the Kleinman–Bylander formalism, effectively project the orbitals of the QM region away from the pseudopotentials, thereby preventing charge spillover. Boundary ECPs and PSPs have been most commonly applied for ionic solids, but extensions to covalent systems have been implemented; known as the pseudobond approach, frozen, specially parameterised hybridised basis functions may be placed on the ECPs to represent the bonding interaction with the missing atom.³⁹ This has been readily used for proteins and organic systems, where a single parameterised C(sp³) hybridised basis function is placed on the boundary carbon atoms.⁴⁰

2.4 Force-field parameterisation

The QM/MM approach can potentially utilise the same diversity of force-fields applied in solely classical simulations, provided an appropriate interface exists between the classical potential and QM Hamiltonian; however, the selected parameters should ideally reproduce both the structure and potential energy surface of the associated quantum calculation. The consistency of the QM and MM structural landscapes is necessary to avoid lattice parameters or bond lengths in the MM region that deviate significantly from those of the underlying QM method, as these would introduce an artificial strain on the QM region and may distort the region of interest from its analogous full QM ground state.

In bulk and condensed systems, the energetic discontinuity between the two regions can lead to significant displacement of QM atoms from their equilibrium positions.⁴¹ These errors occur even if the selected force-field matches the structural properties of the corresponding QM method, resulting in significant errors in QM/MM dynamical simulations.^{42,43} The discrepancy can be alleviated by introducing a buffer region,⁴¹ where the forces on the boundary QM atoms are calculated using terms from an equivalent MM atom (see also: Section 2.5). Alternatively, MM atoms at the QM/MM boundary can be

replaced by pseudopotentials.³¹ However, parameterising force-fields to match the energetics of the QM and MM regions more reliably reduces the energetic discontinuity without introducing additional corrective potentials.^{42–44}

There are numerous strategies for selecting force-fields to satisfy the above conditions. Many studies transfer an optimised force-field designed to minimise the error of observables relative to experiment; however, this approach may lead to energetic and structural discontinuity between the classical and quantum regions, especially if the QM method deviates from the empirical structure. Therefore, it is often desirable to generate bespoke force-fields for the structure under study, which match the structural properties or bonding energies of the QM method (Fig. 2). To achieve such a parameterisation, software such as GULP⁴⁵ and Paramfit⁴⁶ for AMBER force-fields represent two approaches: the former minimising the errors of the least squares fit with respect to calculated observables; the latter using a genetic algorithm to match the energetics and forces of a classical force-field with corresponding QM simulations for a set of molecules. An alternative technique is demonstrated by Zimmerman, Li *et al.*,^{47,48} where successive QM/MM simulations are used to optimise an adsorption energy by iteratively improving the Lennard-Jones parameters. The iterative approach was applied to calculate the adsorption energy of a set of small organic molecules to a range of zeolites, reducing the mean absolute error (MAE) with respect to experiment from 6.7 kcal mol^{−1}, calculated with the default CHARMM force-field, to 1.4 kcal mol^{−1}. The reduced error demonstrates optimisation of the force-fields is a key component for ensuring accuracy in QM/MM simulations.

In the case of polarisable embedding, the dielectric response of the force-field must also align with calculated QM properties. However, in some scenarios (particularly for two-body shell potentials of TiO₂), optimising the static dielectric constants may be incompatible with the objective of acquiring compatible equilibrium lattice constants.⁴⁹ Although some studies have generated accurate force-fields for both the high frequency

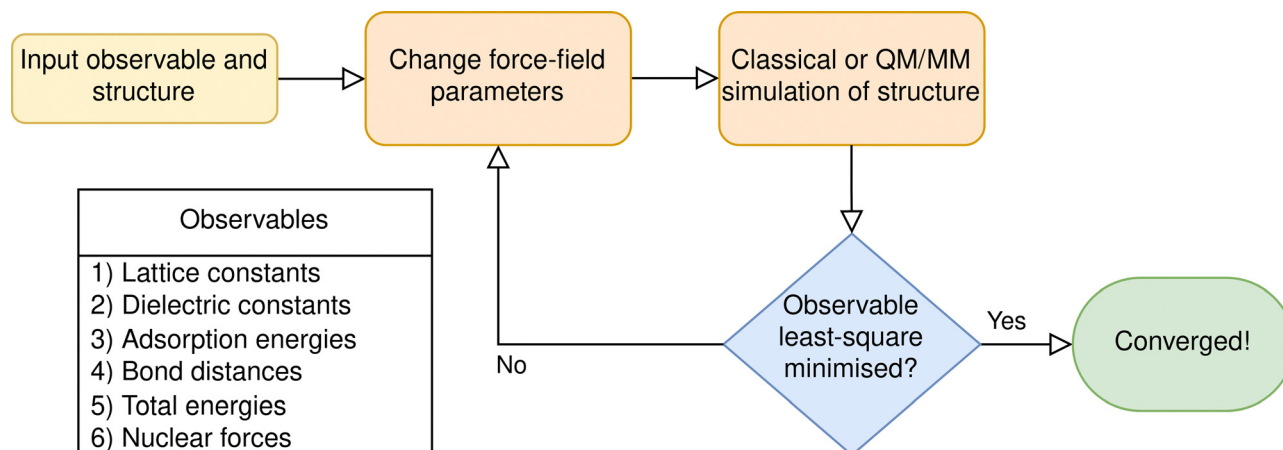
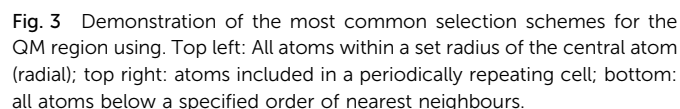


Fig. 2 General schematic for parameterising the classical force-fields for use in QM/MM simulations. Approaches may be differentiated by: (a) their choice of input parameters, (b) the driver used to calculate the observable with the updated parameters (*i.e.*, a MM classical simulation, or a QM/MM simulation), or (c) the mechanism used to minimise the least-square error function, (*i.e.*, local minimisation techniques, such as BFGS, or global optimisation). A non-exhaustive list of observables is provided.



As an alternative, the recent popularisation of machine-learning methods has opened a new avenue for force-field construction to support QM/MM simulations.^{53,54} Zhang *et al.* highlight that machine-learned potentials (MLPs) may, in principle, naturally eliminate the energetic and force discontinuities describing the ML-MM and QM regions on a matched potential energy surface. Furthermore, hybrid-DFT QM/MM has been used to as a reference to optimise standalone MLPs for free-energy simulations of enzymes in the solution phase; accuracies of ~ 1 kcal mol⁻¹ were obtained with respect to the hybrid QM/MM simulations.⁵⁴ Another approach, known as Δ -ML,⁵⁵ looks to generate either correction potentials or energetic/force terms^{56,57} that correct lower accuracy but inexpensive semi-empirical QM/MM methods to a reference, higher order functional. The Δ -ML approach can significantly reduce the cost of the QM calculation, making it more feasible to perform dynamic simulations and obtain free energies at QM accuracy. We note the above implementations have largely been applied to biochemical systems, but the underlying principles should in future be applied to challenges in heterogeneous catalysis.

Alternatively, one can expand the QM region *via* the number of nearest neighbours from a central atom. This approach is popular within zeolite studies, where target properties, such as adsorption energy of reactants, are observed to converge at five nearest neighbours.^{7,58} Furthermore, as zeolites have a well understood topology defined in terms of rings and cages, graph theory approaches can be used to select atoms associated with a substructure of interest.⁵⁹



In protein studies, it has been remarked the QM region size converges asymptotically with respect to an extremely large QM cluster, especially for mechanical embedding.⁶² Even with electrostatic embedding, however, QM regions of several hundreds, or even thousands of atoms, may be necessary to replicate the correct partial charges for negatively charged systems⁶³ and NMR chemical shifts⁶⁴ compared to the full QM simulation. Recent schemes such as the charge shift analysis (CSA) and Fukui shift analysis (FSA) have been employed to analyse which fragments significantly influence the partial charges of the active site.⁶⁵ Using an embedded system with a very large QM region as a comparison, fragments are systematically placed within the MM region and the effect on the charge of the QM region is calculated; fragments producing a charge shift of less

than $\|0.04\text{--}0.05\| e$ are placed within the MM region. Compared to the simple radius convergence strategy, this technique can significantly improve the efficiency of QM/MM simulations by eliminating hundreds of atoms from the QM region; however, studies for ionic solids demonstrate that even small clusters are relatively well converged in terms of NMR shieldings,⁶⁶ though studies including larger QM regions are needed to determine whether the slow convergence found in proteins precludes accuracy below 0.5 ppm.

Partitioning of the QM and MM regions in dynamics simulations requires more consideration. This is especially relevant for bi-phasic simulations with significant chemical interactions between the solute and solvent, meaning the first solvent shell must be included in the QM region.^{67,68} However, with a static definition of QM and MM atoms, exchange of QM with MM solvent molecules over time exposes the QM region of interest to short-range classic forces, decreasing the accuracy of the simulation. To account for this discrepancy, adaptive QM/MM schemes have been implemented to dynamically switch atoms between quantum and classical definitions based on their proximity to the quantum region.⁶⁹ The sudden change in the physics of individual atoms as they cross the boundary can cause a significant discontinuity in the potential energy surface, motivating the development of schemes such as Permuted Adaptive Partition (PAP),⁷⁰ ONIOM-XS,⁷¹ and Size-Consistent Multipartitioning (SCMP).⁷² These schemes introduce a buffer region between the QM and MM regions, which smoothly

interpolate energetics and forces of molecules as they switch definition between the two regions. Recently, adaptive partitioning schemes have been extended to covalent solids, potentially expanding their applicability for heterogeneous catalytic applications.⁷³ Machine-learned methods such as BurNN have also been implemented to provide corrective potentials for QM/MM buffer region.⁷⁴ The corrective approach introduces a Δ -learning correction, yielding a potential that describes mutual polarisation between the buffer and inner QM region quantum mechanically.

Although the adaptive partitioning schemes above are widely adopted, Watanabe *et al.* discovered artefacts remain in both the energetics and statistical ensembles of dynamical QM/MM simulations with SCMP adaptive partitioning;⁴³ this was attributed a physical mismatch in structure and physics between the QM calculation and the selected force-fields.⁴² Force-field solutions are explored in Section 2.4.

2.6 QM/MM codes

QM/MM has been implemented internally in a variety of quantum chemistry and classical molecular mechanics packages, as well as in specialised wrapper software that interfaces these packages. The software implementations vary in their use of subtractive and additive schemes, as well as their implementation of mechanical/electrostatic/polarisable embedding approaches. The implementations may be further distinguished in their use

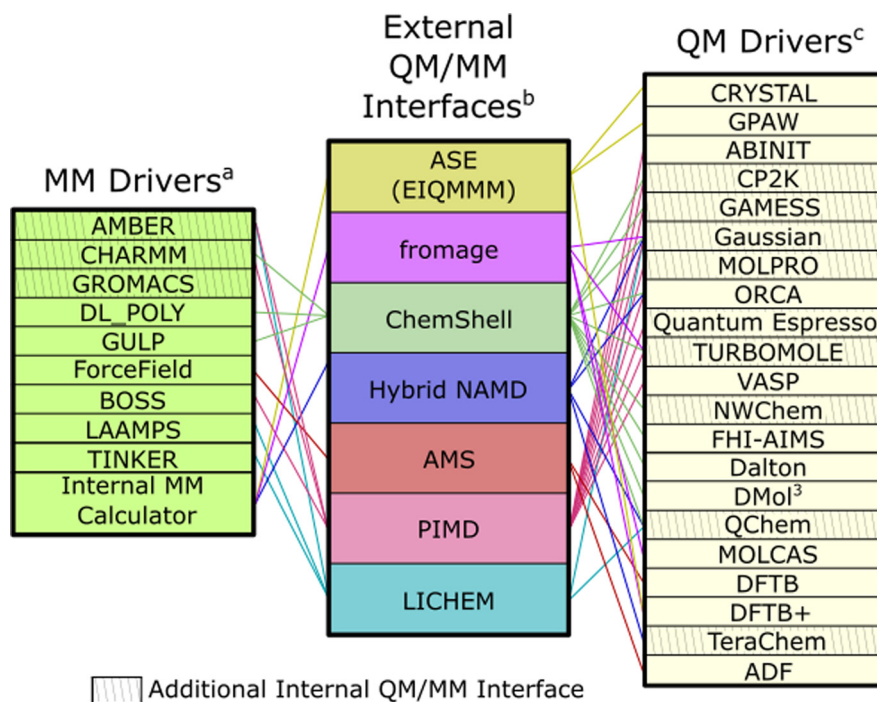


Fig. 4 A network showcasing some of the QM/MM external interfaces and connections to their respective QM and MM drivers. Cross-hatched codes have additional internal interfaces that perform the operations associated with embedding internally, and interface with a specific set of codes. Ordering of entries is chosen to reduce intersection between connections. ^aAMBER,⁸¹ CHARMM,⁸² GROMACS,⁸³ DL_POLY,⁸⁴ GULP,⁴⁵ ForceField,⁸⁵ BOSS,⁸⁶ LAAMPS,⁸⁷ TINKER.⁸⁸ ^bASE,⁷⁶ fromage,⁷⁹ ChemShell,⁷⁷ Hybrid NAMD,⁸⁹ AMS,⁹⁰ PIMD,⁹¹ LICHEM.⁷⁸ ^cCRYSTAL,⁹² GPAW,⁹³ ABINIT,⁹⁴ CP2K,^{75,95} GAMESS,⁹⁶ Gaussian,⁹⁷ MOLPRO,⁹⁸ ORCA,⁹⁹ Quantum Espresso,^{100,101} TURBOMOL,¹⁰² VASP,¹⁰³ NWChem,¹⁰⁴ FHI-AIMS,¹⁰⁵ Dalton,¹⁰⁶ DMol³,¹⁰⁷ QChem,¹⁰⁸ MOLCAS,¹⁰⁹ DFTB,¹¹⁰ DFTB+,¹¹⁰ TeraChem,^{111,112} ADF.⁹⁰

When implemented directly in existing QM or MM packages, direct interfaces to external software are typically realised to evaluate the quantum or classical energetic terms, though the calculation of the MM energy has also been implemented within QM software (with limitations to the range of MM formalisms available⁷⁵). Alternative, more flexible implementations of QM/MM have been realised in wrapper packages that have multiple external interfaces to QM and MM energy calculators (commonly referred to as drivers), with examples including ASE,⁷⁶ ChemShell,⁷⁷ and LICHEM⁷⁸ (Fig. 4). These codes connect a range of QM and MM packages, which is advantageous for broadening functionality; external drivers remain under separate management, but can then provide energetics, forces and densities when required. Management of geometry optimisations, transition state searches or molecular dynamics are performed at the wrapper level. Furthermore, wrappers may be developed to target specific systems or observables; for example, fromage⁷⁹ is centered around calculation of excited states within molecular crystals, whereas PIMD⁸⁰ focuses on simulating the dynamics of condensed systems and enhanced sampling techniques. In principle, the wrapper platforms are designed to be extensible, meaning they may be expanded to a greater variety of driver codes; however, not all software packages support all QM/MM formalisms, with some QM and/or MM package modification necessary to access more advanced features, such as polarisable embedding or capping ECPs.

3.1 QM/MM solvent at the interface

Single configuration adsorption and transition state searches in vacuum are the mainstay of most DFT studies in heterogeneous catalysis; however, solvent has a significant influence on the energetics of adsorption and reaction kinetics in heterogeneous reactive systems.^{113–117} These effects arise from a variety of phenomena including the stabilisation of charged transition states with polar molecules, the introduction of an energy penalty (and entropic gain) for displacing solvent from the catalytic surface,¹¹⁴ and the active participation of the solvent in the reaction mechanism as a co-catalyst or reagent.¹¹⁶

Simulating the dynamic solvent environment necessary for liquid-phase catalysis is a challenging problem with a solely quantum mechanical approach. Firstly, many solvent molecules must be simulated, which drastically increases the computational expense of the simulation. More importantly, the dynamic nature of the solvent environment requires configurational sampling through Monte Carlo or molecular dynamic (MD) approaches. The large number of energy and force evaluations required to obtain equilibrated observables further increases the computational work load. Although *ab initio* MD simulations of surface/water interfaces have become feasible with advances in parallel computing,¹¹⁸ kinetic studies require the evaluation of a large quantity of snapshots to represent the evolution of reaction

Faheem *et al.* formulated such a model using the eSMS (explicit surface-metal-solvent) approach,¹¹ primarily for modelling organic reactions at periodic metallic surfaces. The eSMS-FEP technique includes two forms of embedding to calculate the total energy of the surface/adsorbate complex: A subtractive QM-in-QM scheme is utilised to represent the energetics of the fully periodic metallic surface; and an additive coupling based on the polarisable embedding electrostatic cluster method (PEECM)¹¹⁹ to calculate the contributions from solvent. The system is calculated using a metallic cluster (with size converged with respect to energetics) and adsorbate, which is in turn embedded in an extended system of classical point charges representing both the metallic surface and the solvent environment. The total energy expression is thus:

where $A_{\text{QM/MM}}^{\text{eSMS}}$ is the free energy of the solvated system, $E_{\text{QM}}^{\text{vac,periodic}}$ is the energy of the periodic slab and adsorbate in vacuum, $E_{\text{QM}}^{\text{vac,cluster}}$ is the total energy of the isolated cluster in vacuum, and $A_{\text{QM/MM}}^{\text{solv,cluster}}$ is the free energy of solvation for the embedded cluster. The technique ensures that the long range electrostatics of the DFT metal atoms are included in the total energy evaluation.

Calculations of the free energy changes of activation and reaction were performed with a free energy perturbation (FEP) approach.¹⁰ The coupling between the degrees of freedom of the adsorbate and the solvent requires sampling over a huge configurational space; therefore, a potential of mean force (PMF)¹²⁰ approach can be used to decouple the degrees of freedom of the solvent system by treating the force of the water on the fixed QM system as a mean-field. In effect, the PMF approach integrates out the degrees of freedom associated with the water shell for a given snapshot, meaning integration of the free energy over the reaction pathway can be performed solely using the evolution of the adsorbate reaction coordinates. Although the PMF technique drastically reduces the dimensionality of the FEP calculation, it effectively excludes the impact of solvent on the entropy of adsorbate itself, which introduces a small error. This topic will be discussed further at the end of this section.

The eSMS and FEP techniques have been used primarily to investigate the cleavage of O–H and C–H bonds for various polyols on transition metal surfaces. Initial studies focused on comparing eSMS to its implicit solvent counterpart, iSMS (implicit solvent-metal-surface).¹²¹ Overall, the eSMS schemes predict that the presence of solvent stabilises the free energy of activation ($\Delta G_{\text{react}}^{\ddagger}$) for the cleavage of the O–H and C–H bond by 21.2 kJ mol^{−1} and 3.9 kJ mol^{−1}, respectively; in contrast, the iSMS model predicts a quantitatively lower stabilisation of 3.9 kJ mol^{−1} and 1.0 kJ mol^{−1}, respectively. The findings

emphasise the benefits from an explicit atomistic representation of the solvent environment to capture the energetic changes of reaction; however, the authors also demonstrate that care must be taken when selecting the force-field for the metal and solvent. In particular, the Spohr–Heinzinger and metallic Lennard-Jones potential displayed differences in the structure of water molecules at surface, with the solvation shell with the latter being denser than the former. Although the overall qualitative findings were unchanged, a variation of up to 19.2 kJ mol^{-1} was observed for the free energy changes of solvation.¹²¹

Subsequent studies with the eSMS-FEP approach investigated the impact of the solvent environment on the catalytic activity of different metallic species.¹²² Using the model C–H and O–H cleavage reaction, the overall free energy change and the kinetic barriers are drastically altered by the elemental composition of the metallic surface. In the most extreme case, the activation energy of the O–H bond cleavage is predicted to be $\sim 53.0 \text{ kJ mol}^{-1}$ lower for Cu (111), and only $\sim 11.6 \text{ kJ mol}^{-1}$ lower for Au (111), when compared to vacuum.

The differences in activation energy are primarily rationalised by the degree of charge transfer between the metallic surface and the transition state moieties, which varies with electronic structure for the metallic surface; the interfacial water molecules both reduce the surface dipole of the periodic slab model,¹²³ and they are involved in transfer of charge density back into the metallic surface.¹¹⁸ Overall, the degree of transition state stabilisation correlates strongly with the change in charge transfer for both C–H and O–H cleavage. A secondary factor was the change in the number of hydrogen bonds around the transition state, especially for O–H cleavage, due to the disruption of existing hydrogen bonds caused by the loss of hydrogen. These contributing factors were compiled into a linear descriptor, where the authors further introduced ΔG_{gas} as a variable, which corrects for the greater solvent-induced changes in activation energies for cleavage reactions with large barriers in the gas phase. Fig. 5 shows the resulting strong linear correlation for a descriptor consisting of ΔG_{gas} and the charge transfer between the cleaved fragments, providing strong evidence that these factors strongly influence the change in reaction kinetics for the solvated system. Overall, these studies demonstrate that multiscale atomistic representations of the solvent environment can support the identification of specific solvent effects on reaction energy for different catalytic surfaces.

Addressing the structural dependence of water on the selected classical potentials, Steinmann *et al.* and Clabaut *et al.*^{124,125} developed a set of bespoke force-fields (GAL17 and GAL19) that introduce specific parameters for interaction with periodic noble metal surfaces. After parameterisation against an ensemble of 826 water configurations obtained through DFT, the overall force-field predicts an average error for water adsorption energies of $0.4 \text{ kcal mol}^{-1}$. The force-fields were then utilised within a QM/MM study to model the adsorption of phenol and benzene to the Pt (111) surface,⁹ introducing the solvHybrid method (Fig. 6).

The SolvHybrid technique differs from the eSMS-FEP approach of Saleheen and Faheem in several key ways. Similar

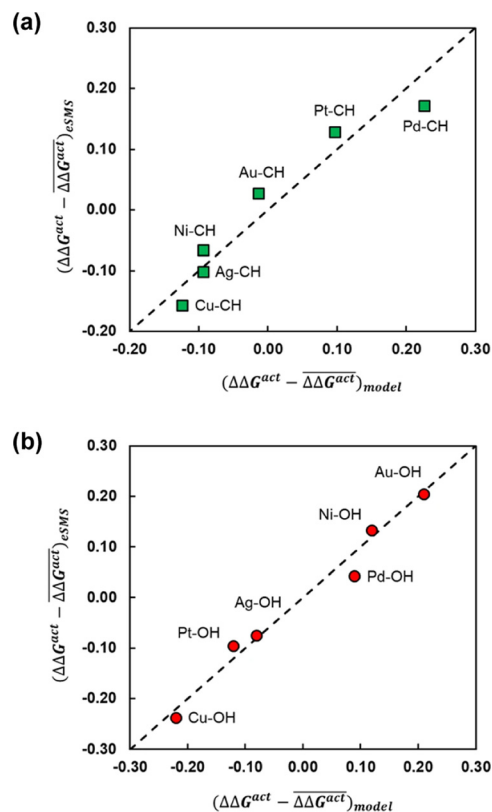


Fig. 5 The effect of solvent on the free energy of activation with respect to vacuum ($\Delta\Delta G^{\text{act}} = \Delta G^{\text{act}}_{\text{sol}} - \Delta G^{\text{act}}_{\text{vac}}$) for the bond cleavage of (a) O–H and (b) C–H in ethylene glycol over noble metal surfaces. The figures compare the errors with respect to the mean $\Delta\Delta G^{\text{act}}$ for a parameterised descriptor $((\Delta\Delta G^{\text{act}} - \overline{\Delta\Delta G^{\text{act}}})_{\text{model}})$ and values calculated with the eSMS method $((\Delta\Delta G^{\text{act}} - \overline{\Delta\Delta G^{\text{act}}})_{\text{eSMS}})$, where bars indicate the mean. The descriptor includes the degree of charge transfer and the gas phase activation barrier of reaction. (a) is performed with a linear fit, while (b) uses a quadratic fit. All energies in eV. Reprinted from ref. 122 under a Creative Commons license (CC BY 4.0).

to the MM-FEP scheme of Steinmann *et al.*,²² the energy of the QM region is computed with a fully periodic simulation, as opposed to the subtractive cluster eSMS scheme. This gives greater flexibility to vary the adsorbate coverage, as the simulation inherits the periodicity of the unit cell, whereas the cluster model implicitly assumes adsorption is taken to the dilute limit. However, the water environment must have the same periodicity as the metallic surface, meaning the classical charges of water are represented with the Particle-Mesh-Ewald (PME) method of CHARMM^{82,126} as opposed to the PEECM approach. A further important contrast between the two approaches is the use of mechanical as opposed to an electrostatic embedding. Finally, SolvHybrid calculates free energy changes through thermodynamic integration (TI) as opposed to the FEP method.

Adopting the nomenclature of their publication, the overall free energy of adsorption in solvent is then represented as

$$\begin{aligned}\Delta_a G^{\text{solv}} &= \Delta_a G^{\text{vac}} + \Delta_a \Delta_h G \\ &= \Delta_a G^{\text{vac}}_{\text{QM}} + \delta \Delta_h G_{\text{MM}} - \Delta_h G_{\text{MM}}(M),\end{aligned}\quad (6)$$



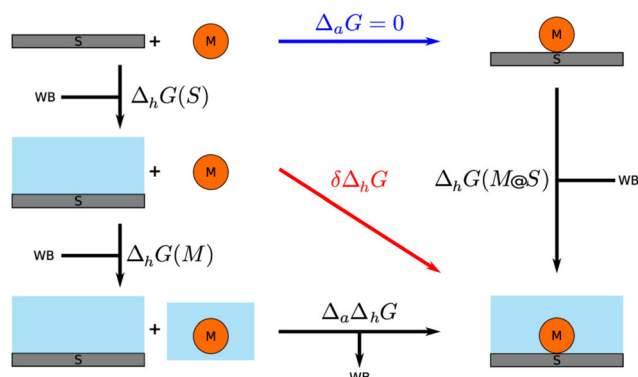


Fig. 6 The thermodynamic cycle for the MM-FEP scheme implemented in the SolvHybrid framework. Reprinted with permission from ref. 9. Copyright 2020 American Chemical Society.

where Δ_h and Δ_a are the free energy changes of hydration and adsorption, such that the terms on the right-hand side of the first line correspond to the free energy of adsorption in vacuum and the free energy change of hydration for the adsorbed complex, respectively. The calculation of $\Delta_a \Delta_h G$ is obtained indirectly through the free energy of desolvation for the adsorbate, M, represented through MM ($-\Delta_h G_{MM}(M)$), and the free energy difference of transferring the molecule (M) from the gas phase to the solvated surface-adsorbate complex, which is calculated through thermodynamic integration ($\delta \Delta_h G_{MM}$). Meanwhile, the entropy of adsorption is calculated only in the gas phase.¹²⁷

Considering the adsorption of benzene and phenol, a decrease in the adsorption energy with respect to vacuum was correctly identified (Fig. 7). Compared to experimental results obtained through cyclic voltammetry ($-9 \text{ kJ mol}^{-1}/-14 \text{ kJ mol}^{-1}$ ^{114,128}), the QM/MM approach obtained a free energy change of solvation of -40 kJ mol^{-1} at saturation coverage. This represents the correct qualitative decrease in the adsorption free energy in solvent relative to vacuum, which

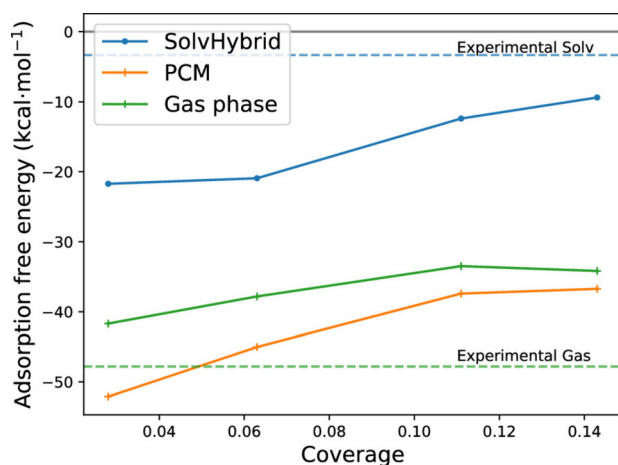
is calculated to be -118 kJ mol^{-1} using experimentally derived values for the enthalpy of adsorption,¹²⁹ and the computationally derived entropy of adsorption *via* TI¹²⁷ at 300 K at a coverage of 1/16. The discrepancy of $\sim 30 \text{ kJ mol}^{-1}$ may be rationalised through the accumulated errors in the choice of density functional, the free energy change of desolvating the metallic surface, and utilising the gas phase adsorption entropy in place of the aqueous entropy. Furthermore, as with gas phase studies of coverage dependent adsorption energies, the use of a single periodic configuration of phenol as opposed to an ensemble of different orientations can introduce a further error of up to approximately 10 kJ mol^{-1} .⁵

Lim *et al.* demonstrate an alternative implementation to the TI and FEP approaches in the DFT-CES/2PT method (DFT in Classical Explicit Solvents – Two Phase Thermodynamic mode).¹³⁰ The calculation of free energies of solvation in DFT-CES/2PT differs significantly from the previously discussed approaches in two ways: (1) the electrostatic embedding of the solvent environment is included in the one-electron Hamiltonian of the QM environment through a mean-field potential of the water environment, which determines the reorganisation energy of the QM region in the MM environment (E_{reorg}); and (2) ΔG_{solv} is determined through vibrational density of state functions, which is decomposed into gas-like diffusive modes and solid-like vibrational modes.^{131,132} Overall, the total free energy of solvation is expressed as a sum of the previous terms,

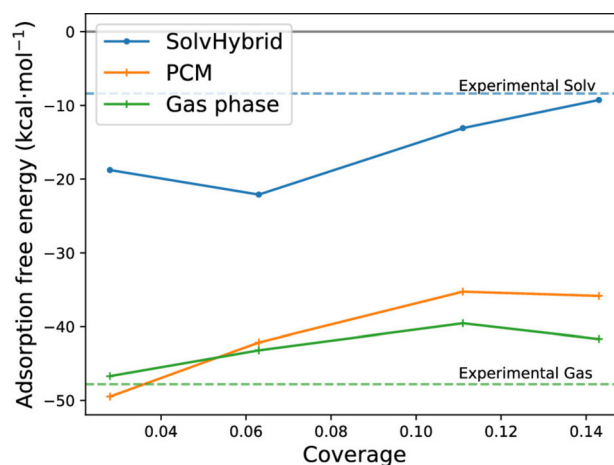
$$\Delta G_{\text{solv}}^{\text{tot}} = E_{\text{reorg}} + \Delta G_{\text{solv}}, \quad (7)$$

where the ΔG_{solv} may be further decomposed into its electrostatic, dispersion and cavitation free energy change components.

When applied to a small test set of molecules, $\Delta G_{\text{solv}}^{\text{tot}}$ yielded a MAE of 5.51 kJ mol^{-1} with respect to experiment, which compared favourably to the MAE of the highly parameterised SM8 implicit solvent model (3.68 kJ mol^{-1}).¹³⁰ DFT-CES/2PT



(a) Benzene



(b) Phenol

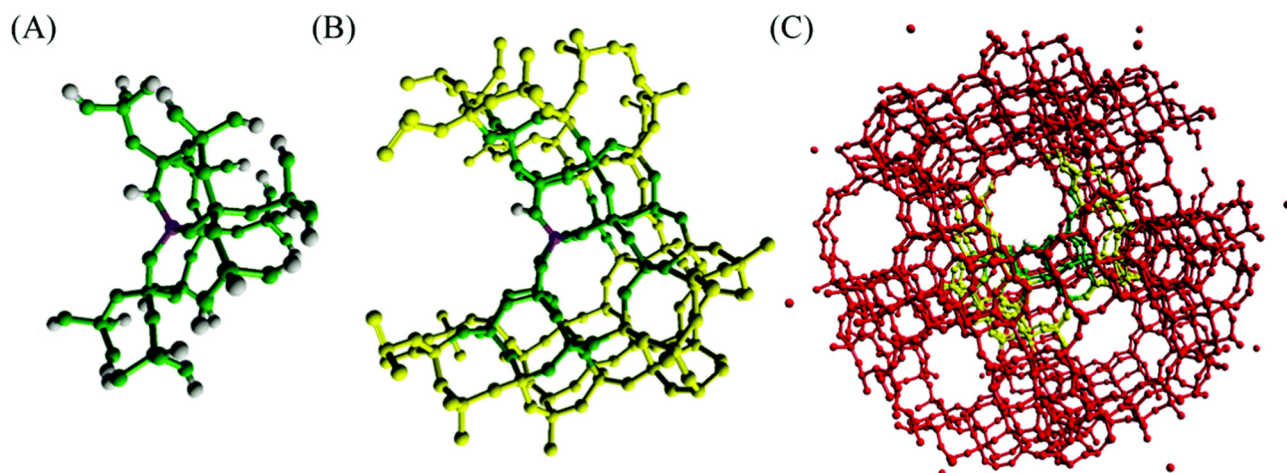
Fig. 7 $\Delta G_{\text{ads}}^{\text{solv}}$ calculated for benzene and phenol on the Pt (111) surface across a range of coverages in the aqueous environment, performed using the MM-FEP scheme. Reprinted with permission from ref. 9. Copyright 2020 American Chemical Society.

introduces an error by excluding the influence of solvent on the entropy of the adsorbate/surface complex. The approach of Clabaut⁹ addresses this challenge somewhat by calculating the overall free energy of solvation for the unadsorbed molecule, and the gas phase entropy of adsorption to approximate the entropy in solvent. Although the QM/MM approaches capture the majority of the entropy changes – which originate from the energetically favourable displacement of solvent from the catalyst surface^{113,128} – the effect of solvent on the entropy of the adsorbed compound warrants further investigation.

3.2 Reactions and heteroatom substitution in zeolites

Zeolites are a class of aluminosilicate materials that have gained much interest due to a range of applications in industry, such as molecular sieves and fluid catalytic cracking of petrochemicals, and are potential catalysts for reactions important for green and sustainable chemistry. Zeolites are crystalline frameworks composed of interlinking SiO_4 and AlO_4^- tetrahedra (T-sites), which form a porous 3D network of cavities and channels with dimensions of 3–20 Å.^{140,141} The existence of AlO_4^- tetrahedra within zeolite frameworks creates a need for a cation to compensate the negative charge on the alumina, and when this compensating cation is a proton, Brønsted acidity is induced within the framework.^{142,143} Furthermore, doping the zeolite framework with heteroatoms induces Lewis acidity,¹⁴⁴ *i.e.*, the ability of a species to donate electron density.¹⁴⁵ In combination, zeolites can behave independently as Brønsted- or Lewis-acid catalysts, or as a bifunctional mixture.^{146–149} The strong Brønsted- and Lewis-acid properties place zeolite catalysts at the forefront of materials for green and sustainable energy; therefore, advancing simulation techniques for the study of zeolite catalysts is of the utmost importance.

QM/MM methods have become a popular approach to model catalytic zeolitic systems due to the local nature of the active site (Fig. 8).^{150–155} QM calculations on small cluster models of the active site have long been used to model zeolite active sites; however, discrepancies from experiment in structures and



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energies can arise due to the small size of the cluster models relative to the extended frameworks. Phenomena such as steric constraints, which are characteristic of adsorption in zeolites, are not well described due to insufficient incorporation of the zeolite pore in the models. Periodic calculations (typically with DFT for reactive processes) are a reliable alternative method for modelling zeolites, but these calculations are generally restricted to small unit cells and/or low-level DFT functionals, such as the generalised gradient approximation (GGA), as the expensive nature of running higher-level exchange–correlation (XC) functionals or larger unit cells creates a computational bottleneck. In this regard, QM/MM methods allow calculations on large models, facilitating the capture of long-range interactions and steric constraints of the zeolite pore whilst using a small QM region around the active site, subsequently allowing for the use of higher-order XC functionals (such as hybrids) or application of correlated methods. As a consequence, QM/MM methods, through embedded-cluster or periodic models (ONIOM, QM-pot) can be applied to investigate a range of zeolite catalysts with applications in heterogeneous catalysis.

Brønsted-acid zeolite catalysts have been successfully employed for the conversion of methanol to hydrocarbons (MTH).^{156,157} As methanol production may be conducted through the reduction of CO₂, or derived from biomass, recent interest has been elevated by the potential as a green route to hydrocarbon synthesis. Embedded-cluster QM/MM investigations by O'Malley *et al.* considered the deprotonation, methanol adsorption, and methoxylation energies in H-ZSM-5 and H-Y zeolites, investigating the influence of framework topology and active site location on sorbate behaviour.¹⁵⁸ The study used a mixture of GGA (PW91) and hybrid (B3LYP and B97-2) DFT, and found that the latter predicted higher deprotonation energies for acid sites in H-ZSM-5 and H-Y. The result was attributed to the superior localisation of electrons when using the hybrid-DFT approach; however, each functional preserved the hierarchy of deprotonation energy, as H-Y < H-ZSM-5 (sinusoidal) < H-ZSM-5 (intersection) < H-ZSM-5 (straight), with the site type given in parenthesis for H-ZSM-5, showing consistency in the trends irrespective of modelling approach. Furthermore, the results for H-ZSM-5 (intersection) using the GGA functional PW91 was 122 kJ mol^{−1} lower (1093.4 kJ mol^{−1}) than those calculated *via ab initio* molecular dynamics simulations on MFI structures using the same functional.¹⁵⁹ The authors attribute this discrepancy to the sorbate–sorbate interactions or charge compensation, which occurs in periodic systems in contrast to the more complete representation of polarisation afforded in embedded-cluster QM/MM methods. The final result highlights that, even at the GGA level of theory, aperiodic embedded-cluster QM/MM approaches can offer advantages over periodic modelling approaches.

The MTH process was investigated further with the embedded-cluster QM/MM method by Nastase *et al.*^{7,58,160} Initially,⁷ the adsorption of multiple methanol molecules in H-ZSM-5 and H-Y zeolites was investigated, concluding that the arrangement of hydrogen bonds between methanol and the framework significantly influence the adsorption strength. By investigating the

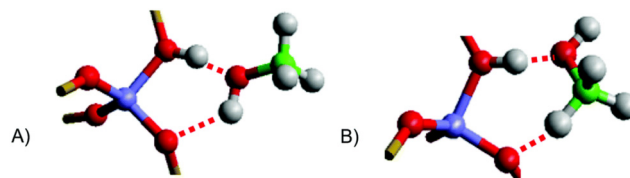


Fig. 9 Representations of methanol adsorption configurations in H-ZSM-5. (A) “end-on” coordination (B) “side-on” coordination. Reproduced from ref. 7 under a Creative Commons license (CC BY 3.0).

different orientations of methanol on the Brønsted-acid sites (Fig. 9), the importance of dispersion in the calculations was identified. Values calculated using the dispersion-corrected hybrid B97-D and second order Møller–Plesset perturbation theory (MP2), which accurately accounts for dispersion effects, are in good agreement (Table 1); quantitatively the results also align to previous dispersion-corrected studies with PBE-D (−115 kJ mol^{−1}), and to experiment (−110 kJ mol^{−1}).^{161,162} The dispersion corrected adsorption energies qualitatively correlate with other hybrid-DFT methods, such as B97-3, but quantitatively disagree by ~50 kJ mol^{−1}, which highlights the need to account for dispersion and the localised nature of electrons. The validation achieved with hybrid-DFT and MP2 are made accessible by QM/MM, owing to the size of the QM region in their models (74 atoms), which is much lower than the number in a periodic DFT calculations (288 atoms).

Nastase *et al.* applied QM/MM methods to study subsequent intermediates formed during the initial stages of the MTH process, namely on dimethyl ether (DME) and carbene species.^{58,163} DME adsorption on Brønsted-acid sites of H-Y and H-ZSM-5 resulted in pronounced proton transfer over the intersectional site in H-ZSM-5, which correlates with greater Brønsted-acidity, and potential superior catalytic performance compared to H-Y. Here, QM/MM was used to perform transition state calculations with hybrid XC functionals, which again is challenging with periodic DFT, and the active site was observed as remaining deprotonated during the reaction for DME production, which contrasts with previous reports. The discrepancy was attributed to the difficulty with GGA-type modelling to localise the electron density, reinforcing the importance of employing higher-level approaches.

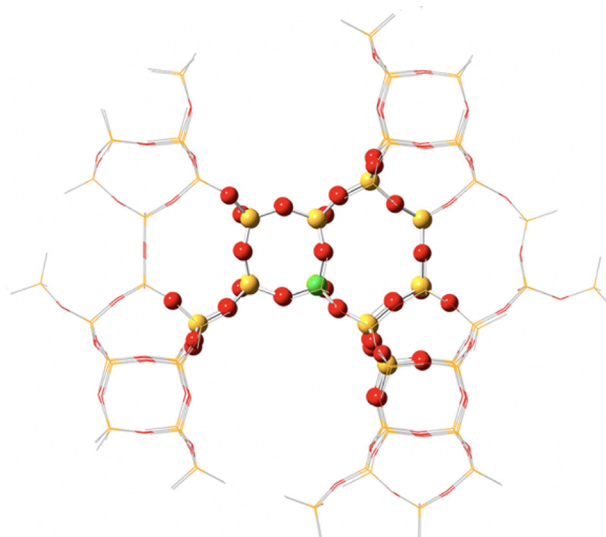
As an alternative embedded-cluster QM/MM models, periodic QM/MM approaches have also been developed and applied extensively towards the modelling of Brønsted-acidity. The

Table 1 Adsorption energies for “end-on” (EO) and “side-on” (SO) methanol with different DFT functionals and correlated methods. Demonstrated for different sites substituted with an Al center. Reproduced from ref. 7. All energies reported in kJ mol^{−1}

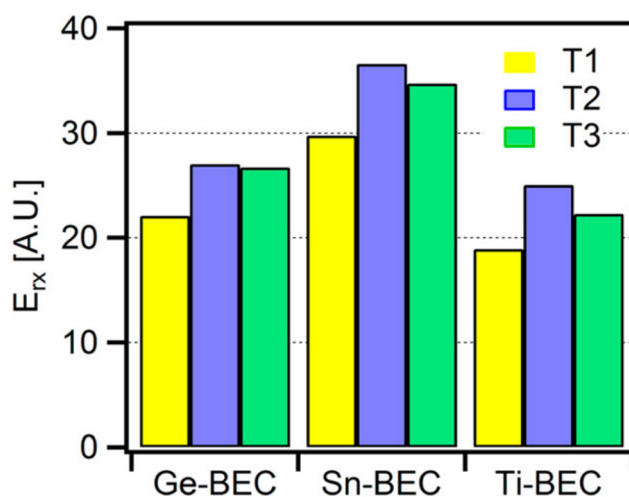
Site	B97-3		B97-D		MP2	
	EO	SO	EO	SO	EO	SO
H-Y	−70	−65	−106	−100	−102	−96
H-ZSM-5 [T12]	−81	−78	−124	−120	−117	−113
H-ZSM-5 [T4]	−82	−80	−126	−115	−121	−112
H-ZSM-5 [T1]	−81	−85	−115	−114	−107	−113



Through QM/MM methods applied *via* the ONIOM approach, Montejo-Valencia *et al.* examined the Lewis-acidity and hydrothermal stability of Ti, Sn, and Ge doped beta frameworks (Fig. 10).¹⁶⁸ A hybrid ω -B97XD XC functional was made feasible *via* QM/MM, and allowed comparison of relative energetics in polymorph A (BEA) and polymorph C (BEC) of the beta



framework. The comparison of polymorphs is necessary as superior catalytic activity and selectivity were suggested for BEC in some reactions.¹⁶⁹ Considering 3 T-sites in each framework (T1–T3), T1 was the most stable site for Ti, Sn, and Ge substitution in the BEC framework (Fig. 11); in contrast, for BEA, both T1 and T2 were the most stable sites for Ti, with energies of 14.84 and 14.89 kcal mol^{−1}, respectively. Calculations on hydrolysed Ti, Ge, and Sn (with one water molecule coordinated), combined with correlation of the Lewis-acidity and the energy of the lowest unoccupied molecular orbital (LUMO), showed Ti-BEC as having the lowest LUMO energies, and thus greatest potential Lewis-acidity, of all heteroatoms for that framework.



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Taking further advantage of the reduced computational workloads enabled by QM/MM, studies may be centered around accurate, high throughput workflows that can validate the lowest energy state for a large ensemble of systems. Such studies are of particular importance in zeolites, where the silicate framework contains a large number of silicon (T) sites which may be occupied by Al. Furthermore, the Brønsted acid protons may occupy any of the neighbouring oxygens.

In order to determine the most stable structures, the workflow of Aljama *et al.* permuted over all possible combinations of Al substitution, Pd cation placements, and Brønsted acid positions. The approach filters the less stable candidates with cheaper GGA methods, and then applies more accurate hybrid simulations to the most stable structures, in a six step workflow: (1) generating all possible combinations of Al substituted T-sites, including double-substitutions within three and four nearest neighbours of one another, and including iterations over the possible positions of the Brønsted protons; (2) selecting the optimum QM region by including the rings containing the relevant substituted T-sites; (3) exchanging one/two Brønsted acid proton(s) for $\text{Pd}^+/\text{Pd}^{2+}$ to gauge stable configurations through single-point energy evaluations; (4) taking the more stable positions and performing geometry optimisations with

and solvent. The size of such systems, along with large time-scales and an interest in high-throughput screening, makes it necessary to focus efforts into development and employment of multiscale models and methods. Similar approaches are being used for simulation of nanoparticles for biomedical applications, in particular for drug delivery, so here we will also consider such studies.

Depending on the locality of the electronic states, the typical schemes for partitioning a nanoparticle system with QM/MM approaches are:

- A MM treatment of a solvent environment around a nanoparticle is equivalent to that considered in Section 3.1, though with reduced dimensionality of the catalyst model. An example of explicit solvent representation within a QM/MM simulation is shown for dopamine molecules bound to the TiO₂ nanoparticle in the water solvent environment (Siani *et al.*¹⁷⁴), where a generalized Amber force-field (GAFF)¹⁷⁵ was used for water, and SCC-DFTB theory for dopamine molecules and nanoparticle. Similar explicit solvation was considered in electrochemical CO₂ reduction to ethylene on Au nanoparticles in a water environment, which was simulated with the recently developed ReQM method by Naserifar *et al.*¹⁷⁶ The ReQM method implements a subtractive QM/MM approach and is based on a special polarisable force-field, RexPoN.¹⁷⁶ RexPoN is a reactive force-field that has terms for non-bonded (van der Waals) interactions, for valence bonds, and supports dynamic polarization and charge distribution *via* the PQEq model.¹⁷⁷

A QM/MM scheme tailored to bulk metals and surfaces has been developed by Zhang, Lu, and Curtin,¹⁷⁹ and subsequently applied to nanoparticle systems. That scheme separates the simulated system in 3 nested regions denoted I, II, and III. The total energy is estimated as follows:

where eqn (8) resembles both additive (eqn (2)) and subtractive (eqn (1)) schemes. Similar to the additive scheme, eqn (8) has a

For accurate simulations of nanoparticle-based catalysts, one takes into account a nanoparticle, reacting species, support,

purely quantum region I that does not need a MM representation; and, similar to the subtractive scheme, eqn (8) has no explicit coupling term. Considering the union of regions I + II as region 1 in Fig. 1, and region III as the region 2 in the same figure, one can rewrite the additive eqn (2) in terms of eqn (8) as:

$$E_{\text{total}}[\text{I} + \text{II} + \text{III}] = E_{\text{QM}}[\text{I} + \text{II}] + E_{\text{MM}}[\text{III}] + E_{\text{couple}}^{12} \quad (9)$$

Here the E_{couple}^{12} term can be derived from eqn (8) and (9) as follows:

$$E_{\text{couple}}^{12} = E_{\text{MM}}[\text{II} + \text{III}] - E_{\text{MM}}[\text{II}] - E_{\text{MM}}[\text{III}]. \quad (10)$$

Therefore, the Zhang, Lu, and Curtin scheme is essentially an additive QM/MM method with a purely classical coupling term. Region I would contain the catalytic active site or an adsorption complex of interest, whilst regions II and III contain solid metal. The solid metal regions can be described with *e.g.* a classical EAM-like potential,¹⁸⁰ and no special treatment of their borders is necessary for evaluation of $E_{\text{MM}}[\text{II} + \text{III}]$ and $E_{\text{MM}}[\text{II}]$.

For calculation of $E_{\text{QM}}[\text{I} + \text{II}]$, the constrained DFT (cDFT) method is employed. A constraining potential is imposed in region II, near the border with region III, which penalizes deviation of the density in region II from the so-called “target electronic density” during subsequent DFT calculations. The target density is defined as a function of metal nuclei coordinates fitted to the bulk density. Compared against the widely used frozen-density embedding¹⁸¹ method, the penalizing potential approach is favoured as it does not require an orbital-free approximation of the kinetic energy functional, which itself is not always available.

Application of the Zhang, Lu, and Curtin approach revealed how surface strain in core-shell metal nanoparticles can alter their catalytic activity towards oxygen reduction reactions, including studies of Pt-Cu,¹⁸² Ni-Pd, Ag-Pd,¹⁸³ and Fe-Pt¹⁸⁴ nanoparticles. In these papers, EAM potentials were used for the whole nanoparticles, and the DFT with constraining potential was used for the adsorption complexes of interest. The QM/MM approach allowed simulation of nanoparticles with curvature that is inaccessible in standard periodic approaches, whilst also accommodating large nanoparticle sizes of up to 270 000 atoms.¹⁸³

For metal nanoparticles, the ONIOM^{13,36} method has been efficiently applied to simulate interactions of large organic molecules with nanoclusters.¹⁸⁵ The novelty of the ONIOM formulation is its focus on the possibility of using more than two layers of theory, for example QM1/QM2/MM 3-layer ONIOM. Due to absence of the coupling Hamiltonian term $H_{\text{MM-QM}}$ in the subtractive scheme, it is simple to use different levels of theory and stack as many of them as necessary; however, the important shortcoming of the subtractive scheme is the need of a classical force-field for the QM region. For organic molecules, the requirement of a force-field is not typically problematic, and the development of the Universal

Force Field (UFF) paved the way to simulation of metal-organic systems. Previously, Ananikov *et al.*¹⁸⁵ reviewed numerous ONIOM studies of influence of ligand structure and molecular environment on activity and selectivity of metal-organic catalysts and surface-deposited nanoclusters, and the reader is directed to this work for further detail. Some recent applications of the ONIOM technique include simulation of DOPA molecules adsorption on gold nanoparticles¹⁸⁶ and amino acid adsorption on bimetallic surfaces in water solvent.¹⁸⁷ In both cases, QM was used for the adsorbed molecules and the first 2 or 3 layers of the metal support, with the UFF used for the rest of the support metal. Geometry of the adsorption complexes, adsorption energies, electron density distribution were computed.¹⁸⁷ Some of simulated systems were compared with pure periodic DFT calculations¹⁸⁶ and experiments, and overall good agreement was observed.¹⁸⁶ These results again show the potential for multiscale QM/MM methods for study of nanoparticulate catalytic systems.

3.4 Defects in metal oxides

The engineering of metal oxide materials is an important field within heterogeneous catalysis, particularly for the development of economical and sustainable processes.¹⁸⁸ Materials such as TiO₂ have been highlighted as powerful photocatalysts, with reactions such as water splitting for hydrogen generation at room temperature labelled as the “holy grail” of catalytic chemistry;¹⁸⁹ however, there are many gaps in our understanding of these processes, including the identity of charge carriers and acceptors and the characterisation of excess charge mediated reaction steps.¹⁸⁹ Similarly, the transformation of CO₂ to longer chain hydrocarbons in carbon capture utilisation (CCU) processes^{190,191} can be catalysed by rock salt oxides such as MgO and MnO; however, the low sorbent capacity of the unmodified MgO surface motivates studies into the rational design of catalysts with higher thermal stabilities and CO₂ adsorption energies.¹⁹¹ All these catalysts are desirable due to their wide availability and low manufacturing costs.¹⁹²

Accurate modelling of these materials requires high-level electronic structure methods, due to the impact of the frontier orbital energy levels on the reaction chemistry. As we have explored elsewhere in this review, the reduction of the QM system allows the use of higher-level DFT, such as hybrid functionals, which can correctly localise defects and represent the energetics of defect formation.³⁵ In this context, QM/MM provides a tool for explaining experimental observations^{193,194} and guiding the design of future catalysts.¹⁹⁵

The embedded-cluster approach provides further advantages regarding the accuracy of the defect formation energies, as they can be calculated at the dilute limit, due to the use of open boundary conditions and an extended structure of MM charges;¹⁹⁶ furthermore, the atomistic representation of the extended system incorporates long-range electrostatics, accounting for polarisation in both the QM and MM *via* polarisable embedding schemes. Berger *et al.* note, however, that challenges persist for highly polarisable systems with high dielectric constants (*e.g.*, TiO₂). In such systems, the MM potential can introduce significant



Using the Kröger-Vink nomenclature, the formation of the oxygen defect, V_O , from a generic ionic oxide surface, MO , is expressed as,

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MM and *ab initio* results increase as a function of defect charge, with greater magnitude than the neutral defect ($F^{\cdot}$: 0.50 eV, F^{2+} : 0.59 eV) and the periodic *ab initio* results consistently more positive. Some of these discrepancies can be justified by differences between density functionals (B3LYP vs. HSE06), as shown by the variation between HSE06 and PBE0 in periodic DFT ($F^{\cdot}$: 0.01 eV, F^+ : 0.20 eV, F^{2+} : 0.43 eV). Fundamentally, however, energies of charged defects in periodic models of surfaces are ill-defined, and require non-trivial corrections, which may be the major cause of the differences; an embedded-cluster QM/MM simulation carried out using the PBE0 density functional, matching the periodic work, shows that cluster models give lower defect formation energy for the same functional ($F^{\cdot}$: 6.26 eV, F^+ : 2.54 eV, F^{2+} : -0.30 eV), again with increasing magnitude of difference between aperiodic and periodic models as the charge increases.

Differences in ΔE_F are of similar magnitude for the more polarisable TiO_2 rutile (110) surface,^{15,206} ranging from 0.27 to 0.68 eV, though here the relative defect stability with each method changes as a function of charge. Again the caveats with respect to charge-corrections for 2D periodic *ab initio* simulations apply; furthermore, Berger *et al.* report errors in their embedded-cluster QM/MM calculations of ± 0.3 eV, due to the use of an analytical correction that accounts for the poor representation of the static dielectric constant within the force-field.¹⁵ Irrespective, both periodic and QM/MM simulations again show defects with higher charge are more stable, and the formation of the F^{2+} vacancy is most favourable. These observations open opportunities with dopant engineering, as p-type doping would lower the Fermi level and thus may increase the occurrence of vacancies at the surface,²⁰⁷ which allows for the tuning of catalytic properties for the ionic solid surface. However, it is important to note that charged defects are mutually repulsive and more sensitive to concentration effect compared to the neutral defects.²⁰² As such, in spite of the high thermodynamic stability of F^{2+} defects in the dilute limit, it is expected that their formation would be disfavoured at higher concentrations.

Overall, embedded-cluster and periodic studies show agreement for neutral defects when taken to the dilute limit, and with appropriate schemes used to correct the spurious interactions; however, the agreement weakens for charged defects, which is likely due to the ill-defined nature of the total energy for charged periodic slab models. The QM/MM embedding approach is advantageous due to the ability to perform the calculations with a single structure, whereas periodic work may use extrapolative schemes, requiring multiple simulations of increasing size. Of course, the accuracy of an embedding approach is dependent on the polarisation response of the MM region, for which a force-field must be parameterised to the *ab initio* parameters in order to emulate the dielectric properties.¹⁵ The requirement of an accurate force-field is undoubtedly a hurdle for adoption, requiring specialist knowledge, and also presents challenges when target observables are incommensurate during fitting (*e.g.* static dielectric and lattice constants⁴⁹).

In addition to calculating the oxygen vacancy formation energies, QM/MM embedded-cluster methods have been applied to the surface adsorption of molecules involved with heterogeneous catalysis. For example, the adsorption of CO_2 on MgO (100) with QM/MM, using the B3LYP density functional, reveals both charged and uncharged oxygen vacancies radically alter the energetics of adsorption.¹⁹⁴ Considering the parallel and perpendicular orientation of CO_2 with respect to the surface (Fig. 13 and Table 4), Downing *et al.* demonstrated that both the neutral and singly-charged vacancy stabilise both bonding modes compared to the pristine surface. The vertical binding mode becomes more favourable compared to the perpendicular, with increased binding strength when there is greater charge transfer between the MgO and the bonding O of CO_2 . Relaxation of the perpendicular CO_2 eliminates the oxygen vacancy, restoring the pristine Mg (100) surface and a CO molecule; thus, design considerations for catalytic transformation directly to hydrocarbons, rather than *via* Fischer Tropsch, should look to weaken this binding mode, or investigate alternative oxide materials with weaker CO_2 binding. In this

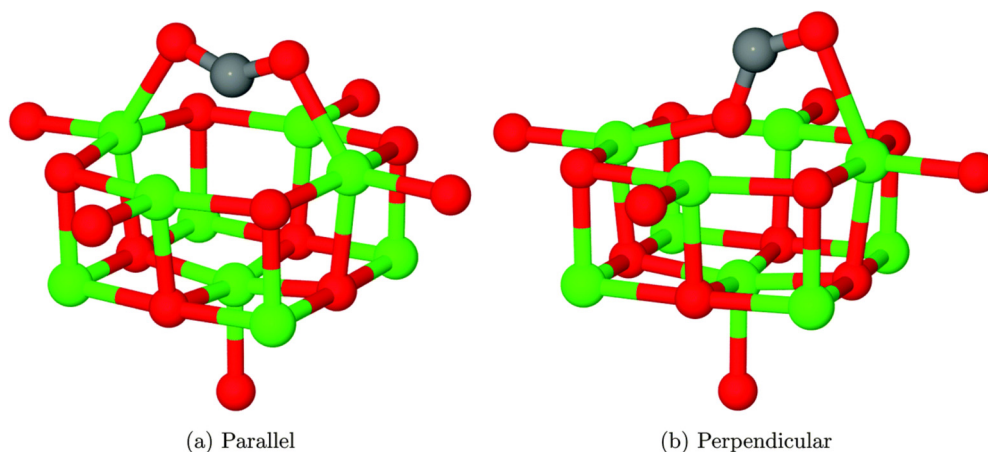


Fig. 13 Parallel and perpendicular adsorption modes of CO_2 adsorption on an MgO surface with an $F^{\cdot}$ defect containing two localised electrons. Figure reprinted from ref. 194 under a Creative Commons license (CC BY 3.0).



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