



Task-farming parallelism in ChemShell

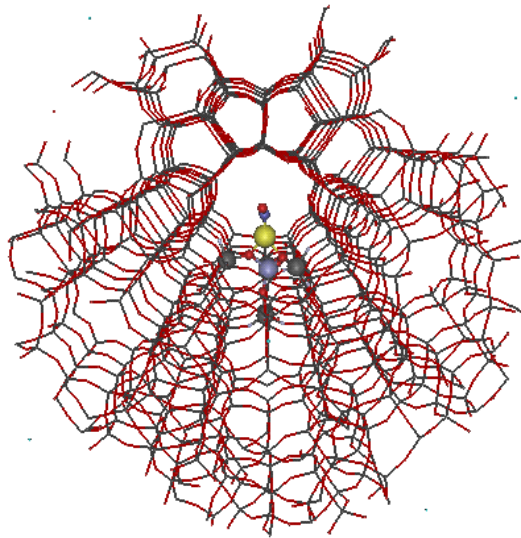
Tom Keal
STFC Daresbury Laboratory

8 May 2013

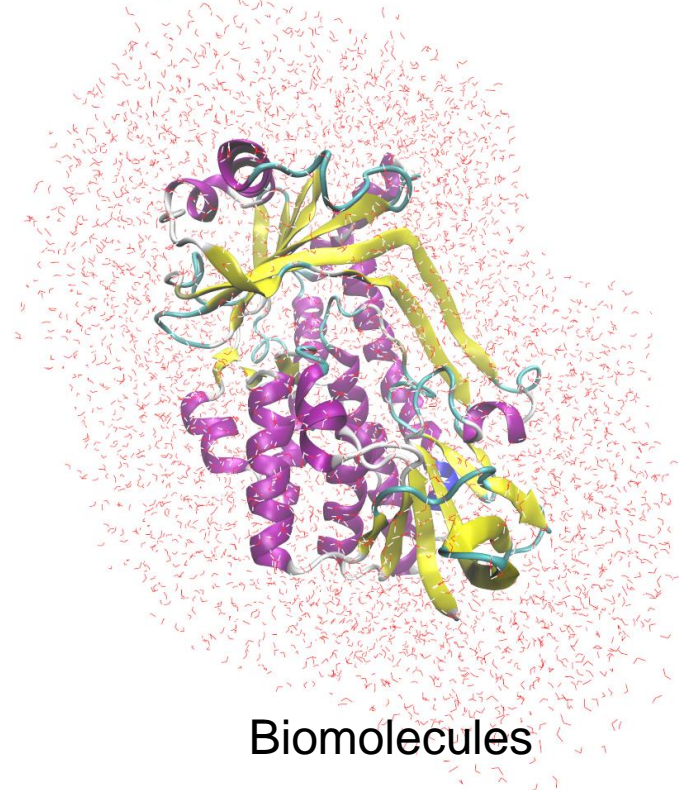


QM/MM modelling

- **ChemShell combines high-level quantum mechanical calculations for reactive region with classical force fields for environment**



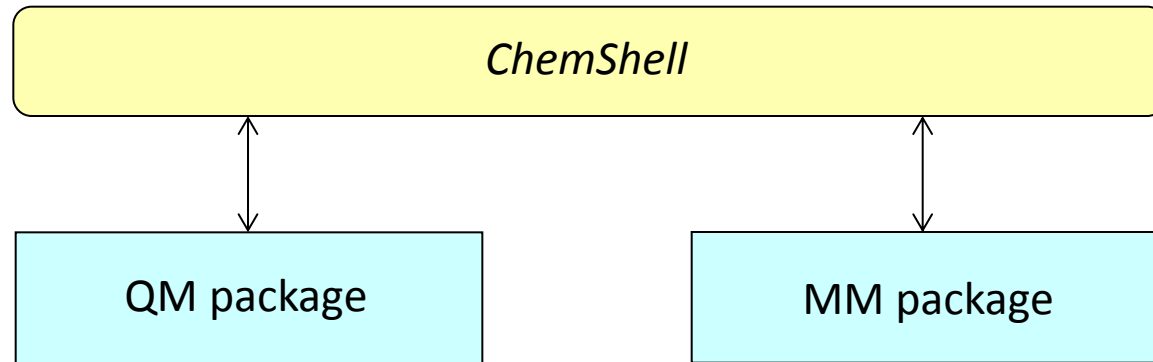
Materials



Biomolecules



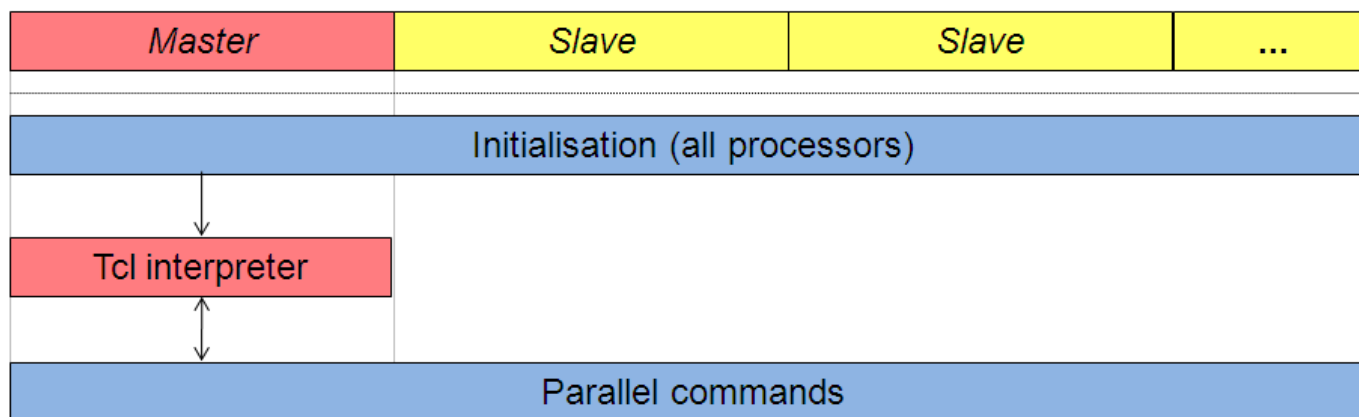
QM/MM calculations



- **User controls ChemShell using Tcl commands**
- **ChemShell modules written in C/Fortran**
- **External programs called to obtain energy & gradient**
- **ChemShell forms combined QM/MM gradient**

ChemShell in parallel

- ChemShell can run in parallel using MPI

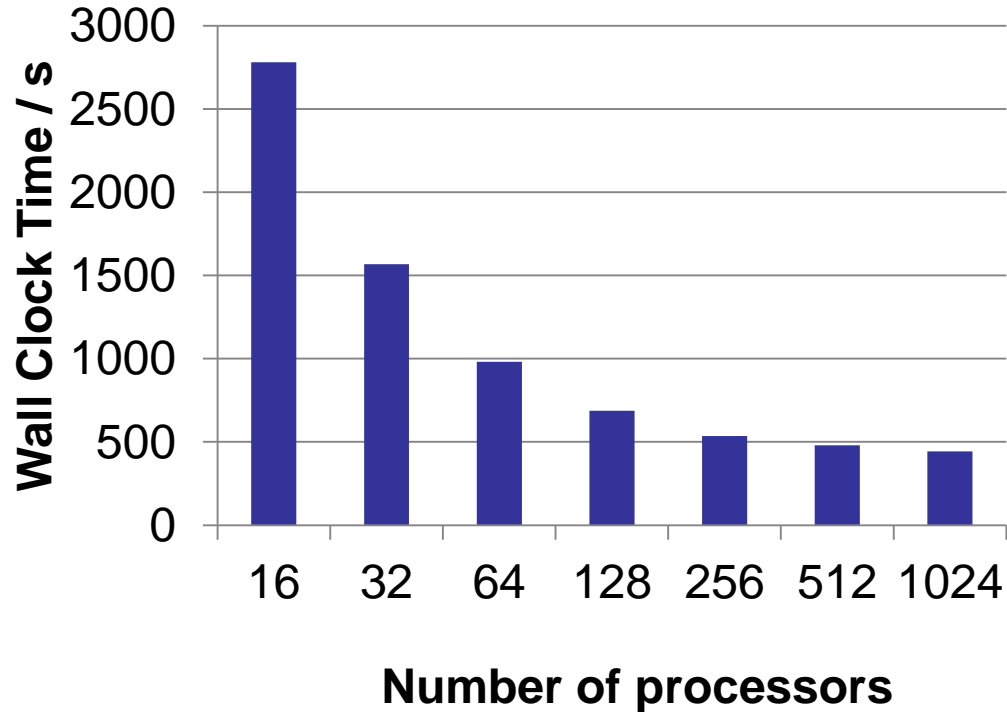


- Takes advantage of parallel external programs
 - e.g. parallel energy/gradient evaluation in GAMESS-UK
- However, this approach does not scale well to large numbers of processors



Typical scaling behaviour

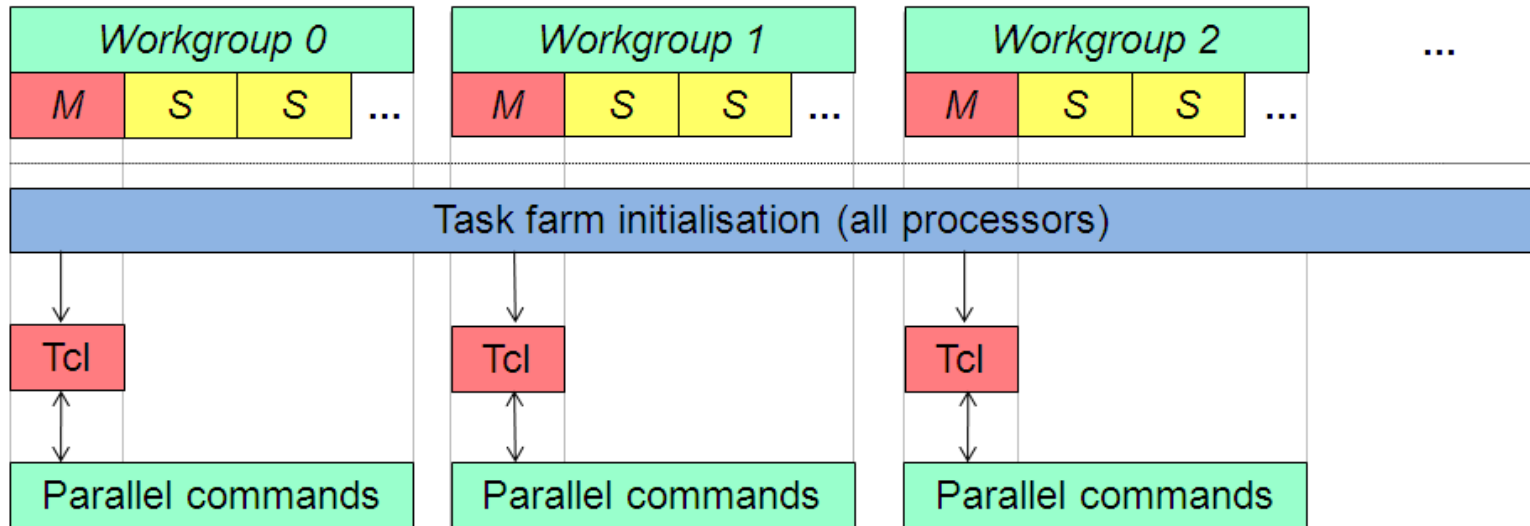
- Single-point GAMESS-UK timings (57 atom silicate cluster, TZVP)





Task-farming parallelism

- Aim is to parallelise ChemShell algorithms as well
 - e.g. parallel Hessian evaluation, optimisations, etc.
- Task-farming approach: divide up processors into workgroups working independently on tasks





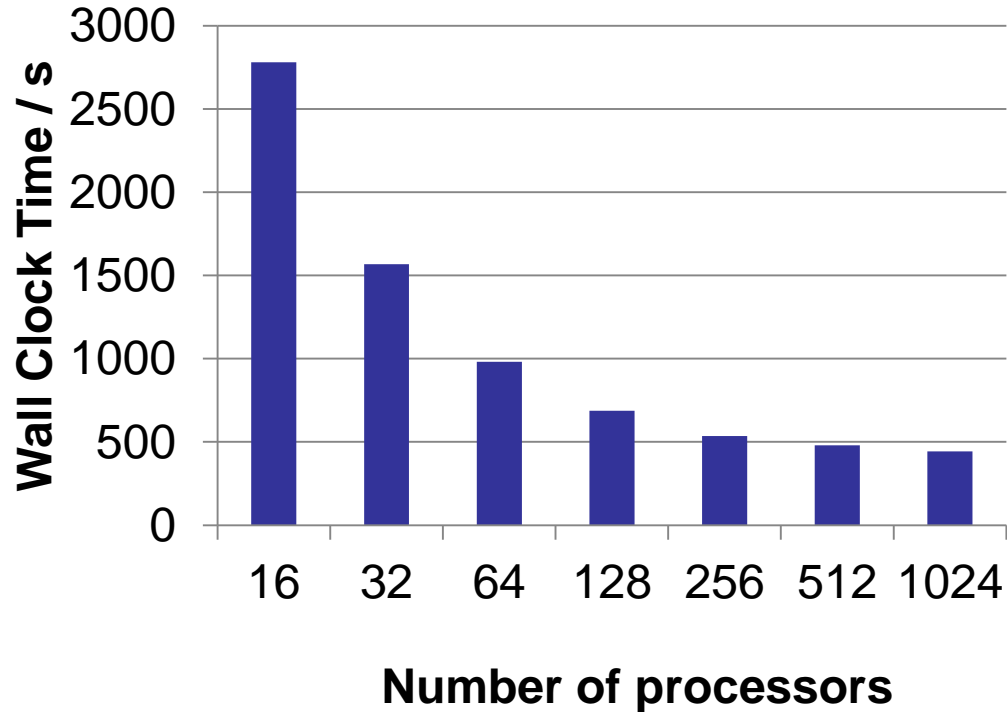
Task-farming parallelism

- **Workgroups are essentially independent**
 - Separate outputs, working directories
- **All lower-level parallelism (e.g. external calculations) occurs within a single workgroup.**
 - GAMESS-UK, GULP, etc. modified to work within workgroups
 - MPI communicator split by ChemShell and passed to external program
- **All workgroups are controlled via a single Tcl input script**
 - Tcl commands to allow workgroup-specific tasks, synchronise workgroups, share data, etc.



Example: finite-difference Hessian

- Single-point GAMESS-UK timings (57 atom silicate cluster, TZVP)





Example: finite difference Hessian

- Performance can be improved by dividing up the set of 1024 processors into workgroups:

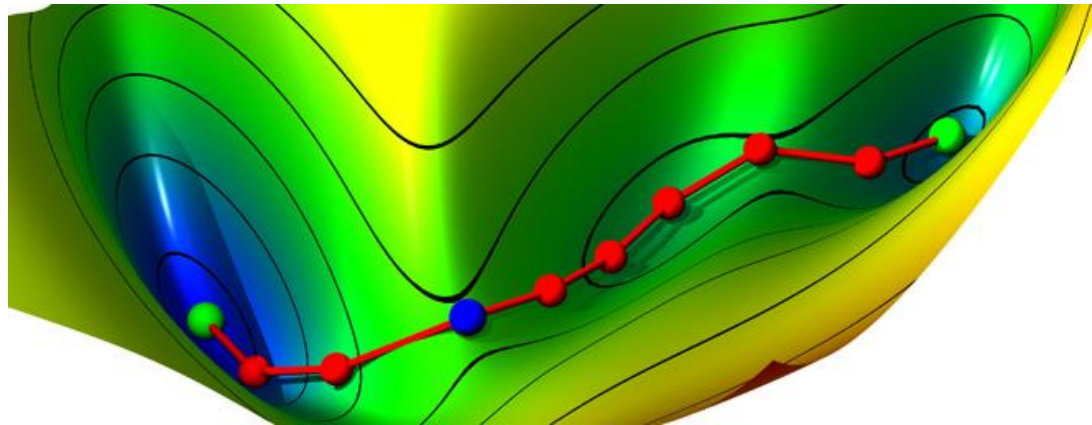
Workgroups	Procs / workgroup	Time / s	Speed-up
1	1024	52762	
64	16	7812	6.8

- Optimal number of workgroups depends on system size / total processor count / algorithm



Example 2: nudged elastic band method

- Optimising reaction paths: finds the minimum energy path

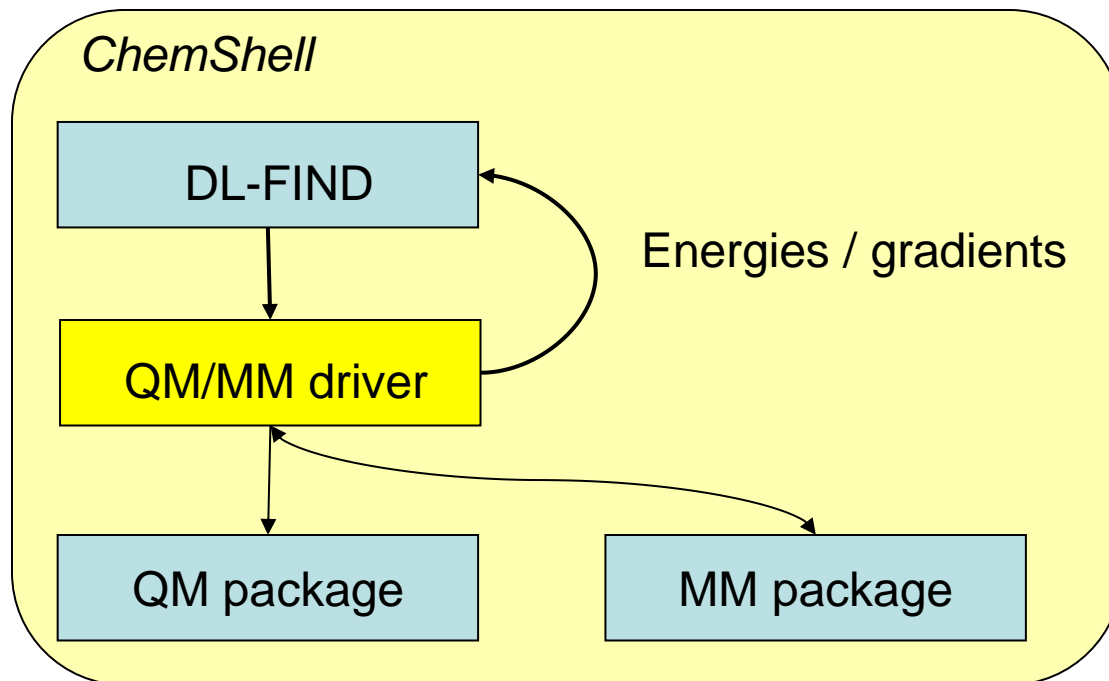


- Multiple images, connected by spring forces.
- Climbing image to find transition state
- Image E/g evaluations are independent, so can be parallelised



DL-FIND

- An open-source geometry optimisation library
- Interface to ChemShell for QM/MM optimisations
- NEB routines implemented in DL-FIND





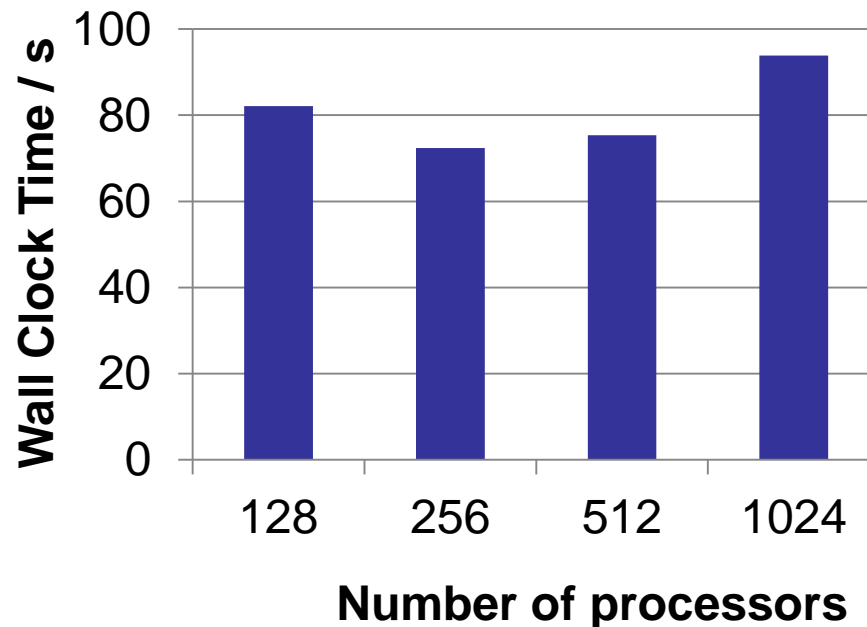
Parallel NEB in DL-FIND

- **ChemShell/DL-FIND parallel interface implemented**
 - Pass the relevant workgroup information
- **Each workgroup runs DL-FIND**
 - Images allocated according to workgroup ID
 - Images remain with same workgroup throughout calculation (benefit from restart information)
 - Energies/gradients shared between workgroups at the end of each cycle



Parallel NEB timings

- **Test system: hydrogen interchange on Al-doped ZnO (see later)**
 - 3207-atom QM/MM cluster (32 QM atoms, PVDZ)
- **Number of workgroups is limited by number of NEB images**
 - Test system: 10 images (2 fixed)
- **Single point calculations:**





Parallel NEB timings

- Test over 50 cycles of NEB
- Again divide up set of 1024 processors into workgroups:

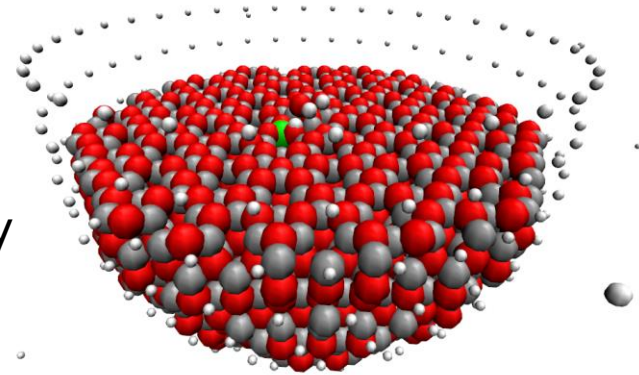
Workgroups	Procs / workgroup	Time / s	Speed-up vs 1024	Speed-up vs 256
1	1024	26404		
1	256	23536		
2	512	14673	1.8	1.6
4	256	7089	3.7	3.3
8	128	3110	8.5	7.6

- Task-farming is essential to make use of > 256 cores



Hydrogen dissociation study

- **Al-doped ZnO important in industrial catalysis (methanol synthesis)**
 - Important to understand interaction of hydrogen with ZnO
 - Catalytically important polar oxygen-terminated surface studied
 - Active site postulated to be vacant oxygen interstitial surface site
- **Embedded cluster model: 3200-atom system**
 - 61 QM atoms around a VOISS centre with nearby Al impurity
 - DFT ionic pseudopotentials (e.g. Zn^{2+}) associated with 63 atoms in the boundary region (cations only)
 - Polarised MM forcefield (shell model)



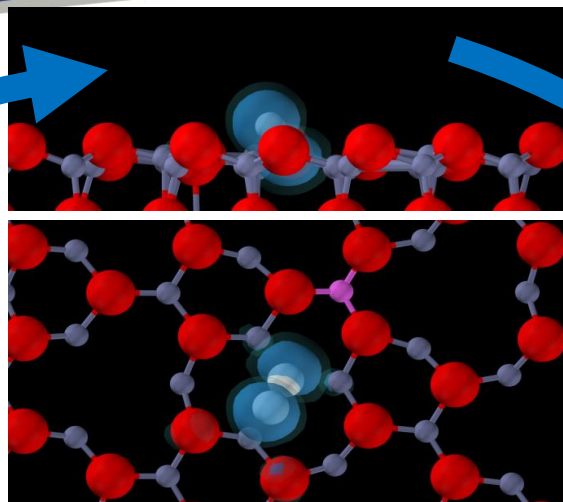
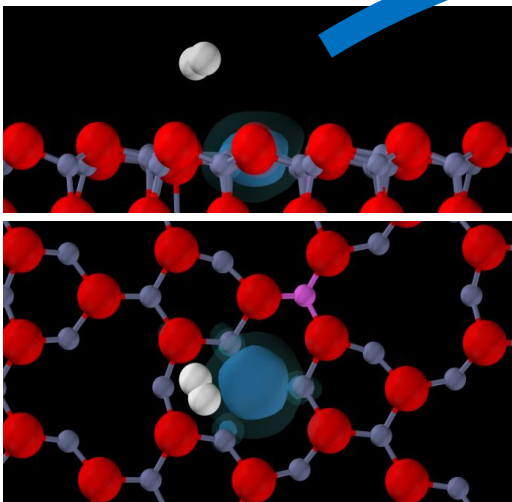


Exploring the potential energy surface

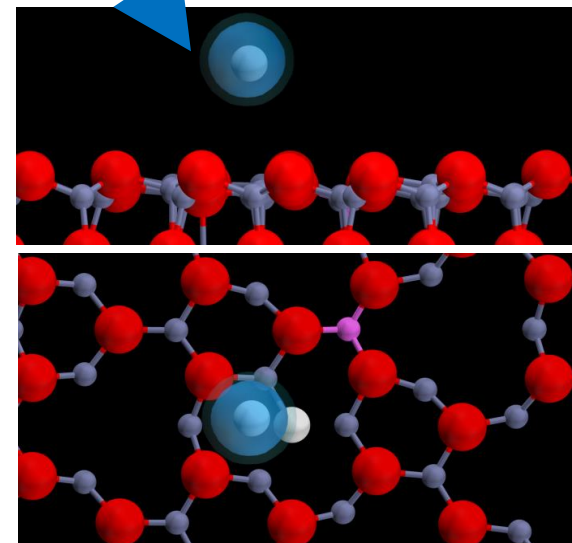
- **DFT: B97-2 functional, TZVP basis set**
- **Starting point: physisorption of hydrogen molecule on surface**
- **Endpoint: previous studies showed dissociation is heterolytic**
- **NEB method used to calculate barrier for dissociation**
 - First run indicated a stable intermediate, so two stage reaction
 - Two NEB runs then required: R->I and I->P



30.2



-9.1

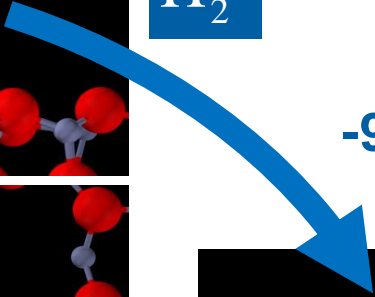
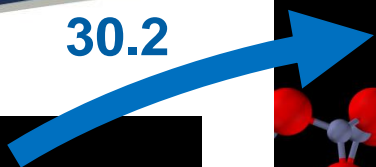
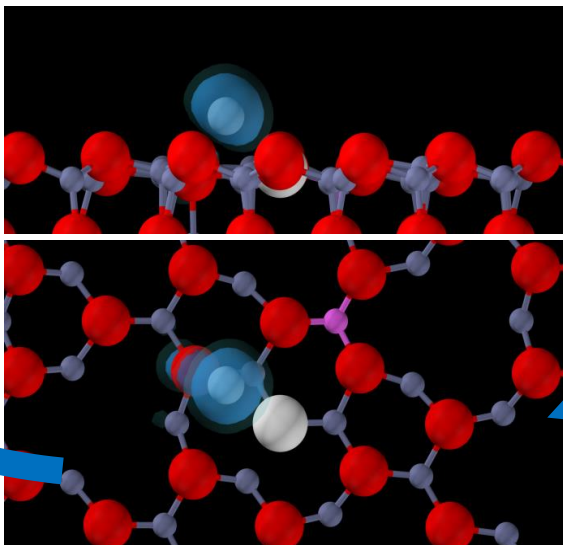


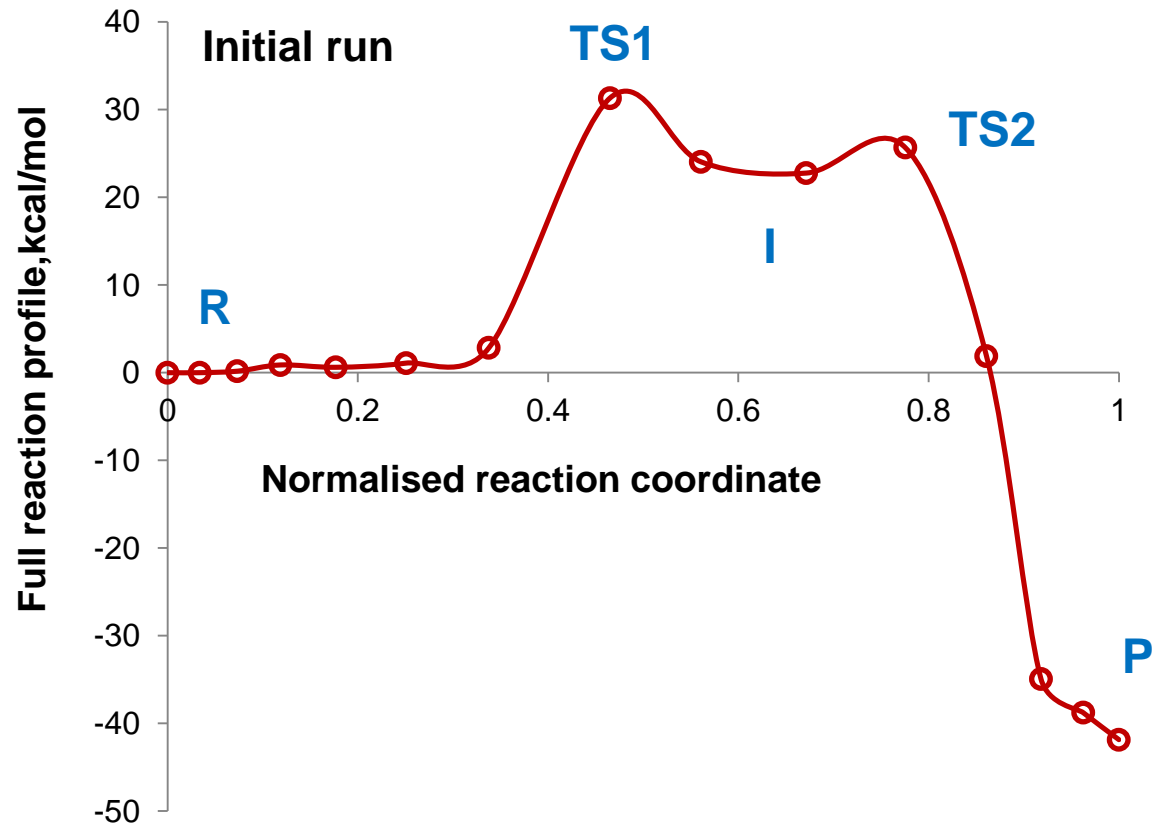
4.5



Energies
in
kcal/mol

-67.5







Comparison with experiment

- **Total activation energy of H₂ desorption = E(TS1) - E(P)**
 - 72.1 kcal/mol
- **Bowker et al: 71.7 kcal/mol**
- **Kunat et al: 77.9 kcal/mol**
- **Consistent with irreversible reaction**
- **Dissociated surface H's useful for further catalytic reactions over Al/ZnO (methanol synthesis)**



Applicability of task-farmed ChemShell

- **Only relevant to large scale calculations**
- **External programs generally need to be modified and linked in**
 - QM codes: GAMESS-UK, DALTON, LSDALTON, FHI-AIMS
 - MM codes: GULP, DL_POLY
- **Works with parallel diagonalisers: PEIGS, ScaLAPACK (MPIBLACS)**
- **Not yet compatible with Global Arrays**
 - Required to support NWChem
 - Possible in principle?



ChemShell on BlueGene/Q

- **ChemShell ported successfully with GAMESS-UK and GULP**
 - Some modifications required (released in v3.6)
 - GAMESS-UK needs optimising
- **FHI-AIMS also working**
- **NWChem awaits a Global Arrays implementation on BGQ**



Outlook

- **Microiterative optimisation in DL-FIND**
- **New embedding methods (NWChem)**
- **New QM/MM techniques, e.g. adaptive QM regions**
- **Replacing Tcl user interface with Python**



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www.chemshell.org