

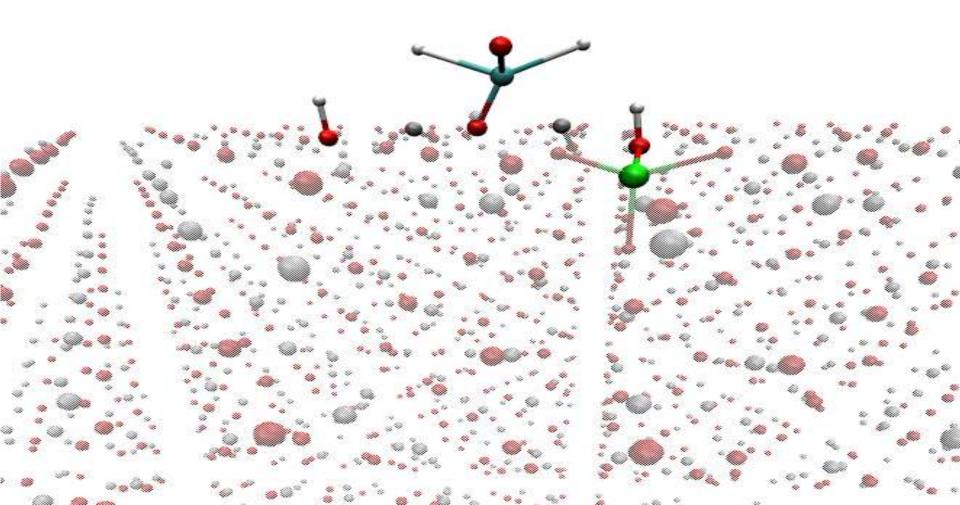
# Efficient massively-parallel tools for the study of catalytic chemistry

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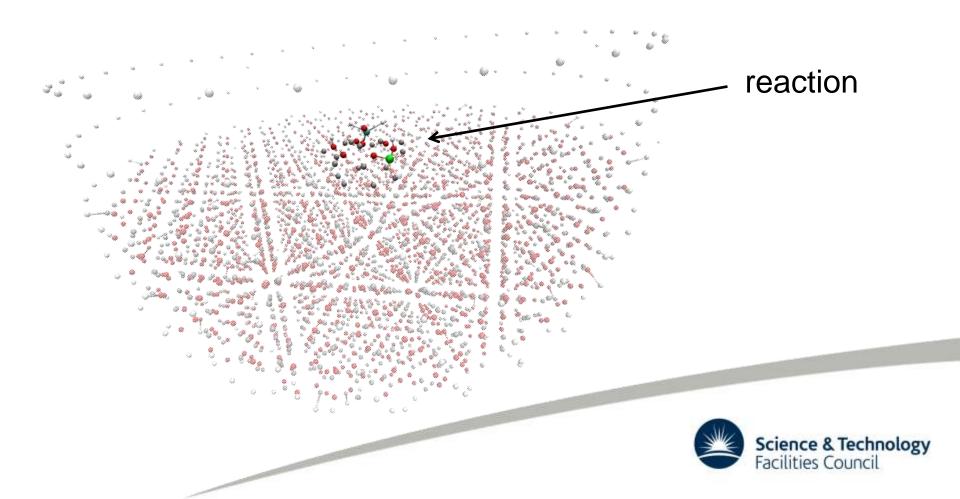
#### Catalytic chemistry example

Methanol synthesis on Al-doped zinc oxide (UCL)



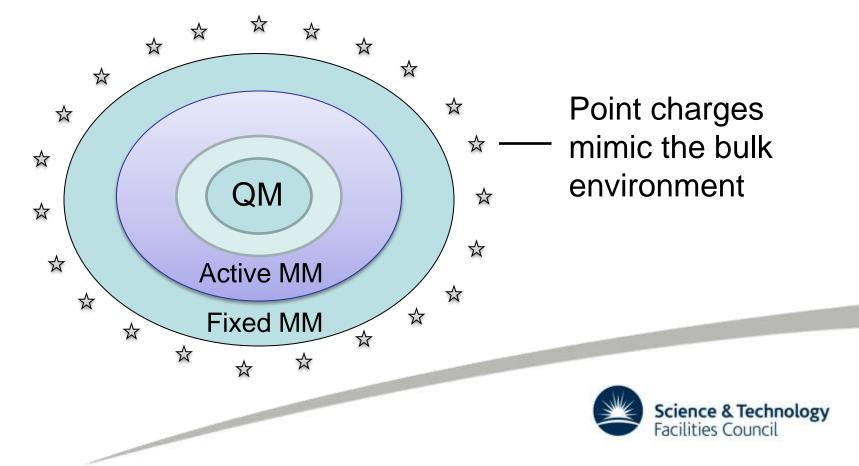
#### **Cluster calculations**

• Cut out a representative part of the surface



## QM/MM cluster calculations

- Quantum mechanical description required for reaction
- Molecular mechanical description for environment



## ChemShell

- Computational chemistry environment
  - www.chemshell.org
  - Tcl front end, C/Fortran behind the scenes
- Particularly useful for QM/MM calculations
  - Interfaces with external QM and MM programs to obtain E/g
  - ChemShell forms combined QM/MM energy/gradient
- Utilities for cutting clusters
- For our cluster calculations...
  - GAMESS-UK for the QM region
  - GULP for the MM region



# **ChemShell** in parallel

• ChemShell can run in parallel using MPI

Node 0	Node 1 N			
Initialisation				
Tcl				
Parallel code				

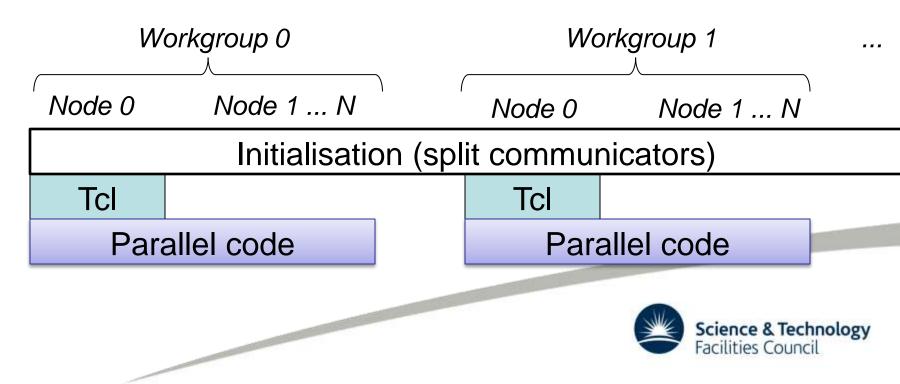
Takes advantage of parallel external programs

- E.g. parallel energy/gradient evaluation in GAMESS-UK
- However, this approach does not scale well to 000's of processors



# Task-farming parallelism

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



# Task-farming parallelism

- Workgroups are essentially independent
  - Separate stdout/stderr

- Separate working directories to prevent file conflicts
- All lower-level parallelism (e.g. GAMESS-UK calculations) occurs within a single workgroup. Therefore the workgroup communicator must be passed to GAMESS.
- All workgroups are controlled via a single Tcl input script
  - Tcl commands to allow workgroup-specific tasks
  - Workgroups can be explicitly synchronised
  - Local ChemShell objects can be made available globally



### Finite difference Hessians

- Second derivative matrix of the energy, calculated numerically using first derivatives (gradients)
- ChemShell's 'force' command split up into three stages to allow task-farmed execution:

Science & Technology

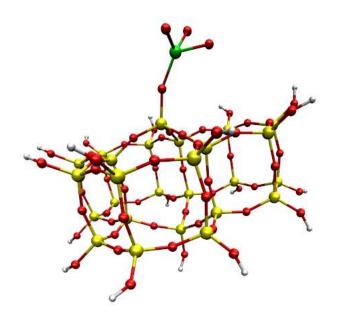
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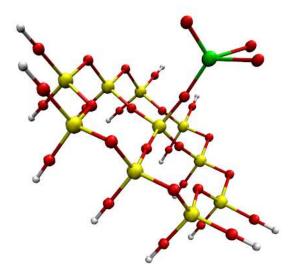
1. Precalculate the required gradients

- Divided up by atom number (static load-balancing)
- 2. Make gradient objects available globally
- 3. Build the Hessian from the precalculated gradients

#### Hessian benchmarks

- GAMESS-UK, DFT (B3LYP), lanl2 basis (pure QM)
- Two VO<sub>3</sub>/silicate clusters:





Cluster 2: 57 atoms

Cluster 1: 111 atoms

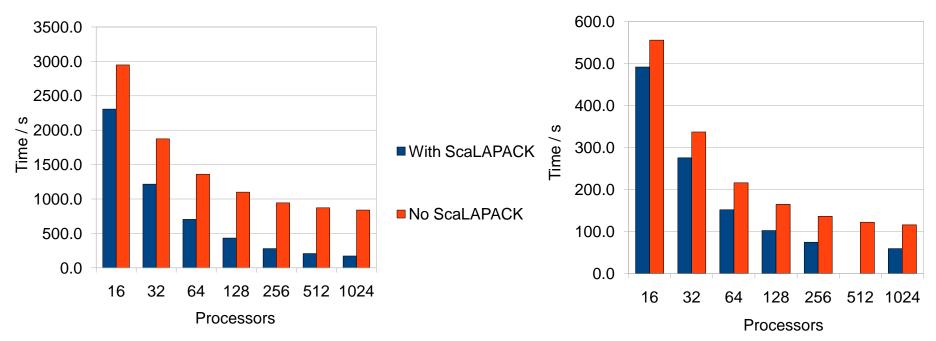


### Single point calculations

 Problem: ScaLAPACK routines in GAMESS-UK do not support split communicators

**Cluster 2** 

compare with and without ScaLAPACK



Cluster 1

## Hessian benchmark results

 Single gradient timings suggest that 32 workgroups is optimal

CLUSTER 1	Finite difference type		CLUSTER 2	Finite difference type	
Time / s	Single point	Two point	Time / s	Single point	Two point
Origina	28441.2	57842.1	Original	5769.8	11430.5
32 workgroups	12335.4	23690.8	Single workgroup	9150.9	18219.6
			32 workgroups	1256.6	2454.6
Task farming speed up	factor				
vs Original	2.3	2.4	Task farming speed up	factor	
			vs Single workgroup	7.3	7.4

vs Original

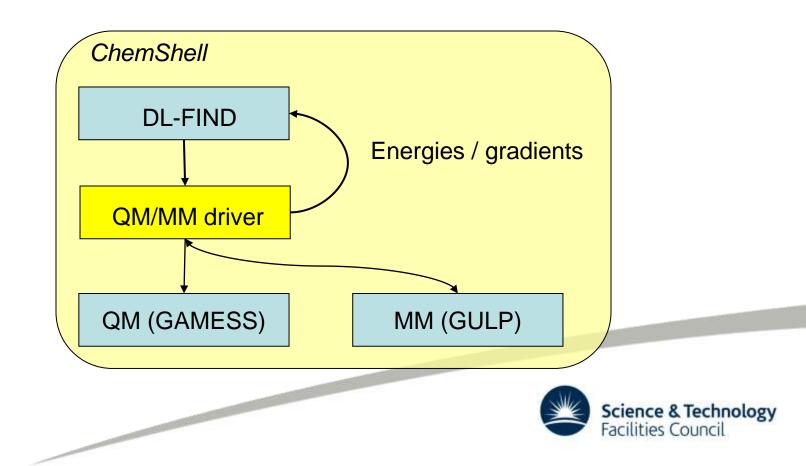
- Original = single workgroup with ScaLAPACK
- Single workgroup = no ScaLAPACK (like with like)



4.6

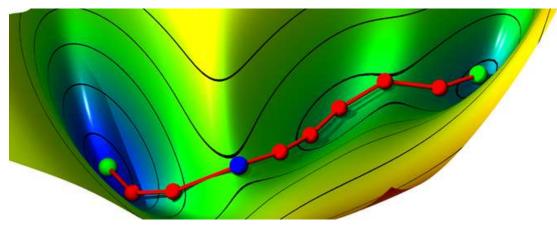
#### **DL-FIND**

- An open-source geometry optimisation library
- Interface to ChemShell for QM/MM optimisations



## 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



#### Parallel NEB

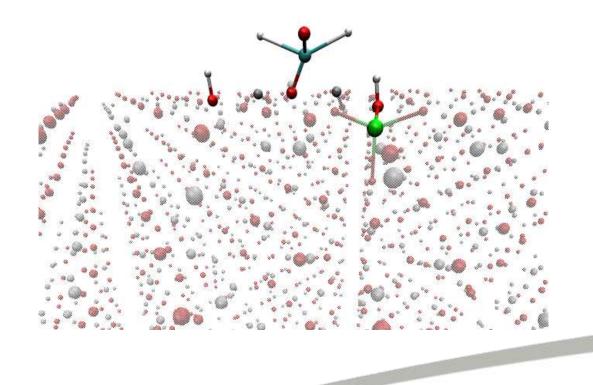
- ChemShell/DL-FIND parallel interface required
  - Pass the relevant MPI communicators, workgroup information
- Each workgroup runs DL-FIND

- Tasks allocated according to workgroup ID
- Energies/gradients shared between workgroups at the end of each cycle (allreduce)
- Some complications compared to serial case
  - Status, restarts
  - First cycle in serial for benefit of QM program (guess vectors)
  - Frozen images speed up serial calculations vs. parallel



#### Parallel NEB test

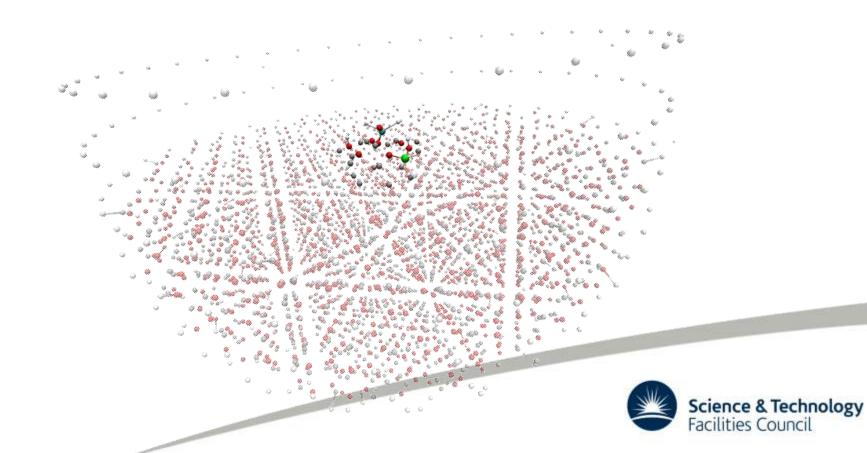
 NEB to find barrier of hydrogen exchange for CO<sub>2</sub> on Aldoped ZnO surface:





#### Parallel NEB test

- QM/MM cluster calculation
  - MM: GULP must use MPI communicator from ChemShell



#### Parallel NEB test details

- 3207 atoms
  - QM: 32 atoms, DFT(B97-1), double zeta basis (ECP for Zn)
  - MM includes shell model for polarisation

- 10 NEB images (including end points and climbing images)
- Speed up > 2 expected over standard 1024-processor run
  - Preliminary results (Time for 50 cycles of NEB):

1 workgroup of 1024 procs	10 hours
4 workgroups of 256 procs	3 hours



## Summary and outlook

- Task-farm parallel framework implemented in ChemShell
  GAMESS-UK and GULP made 'task farm aware'
- Finite difference Hessian code parallelised
- Parallel interface between ChemShell & DL-FIND
- Parallel NEB implemented

- Next milestone: interface ChemShell with DL-FIND's parallel optimisation algorithms
  - Genetic algorithm, stochastic search



#### Acknowledgements

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