



European Union Funding for Research & Innovation

CP2K – QM/MM Practical

Results – Single node

- Using 1 thread results in out of memory error
 - There is not enough memory per process (128 here)
 - 2 threads allow double the memory per process, remember threads share memory so they do not need their own copy of data
- 4 threads has better performance.
- 32 threads very slow. The threads span 2 memory regions on the node which means data is slow to access

Threads	Time CP2K (s) T_total	Time mp_alltoall_z2 2v (s)	Time mp_sum_dm 3 (s)	Time mp_waitan y (s)	Total time mp_ routines (s) T_mp
1	n/a	n/a	n/a	n/a	n/a
2	224.49	18.377	14.758	4.767	37.902
4	212.471	15.599	8.821		24.42
8	236.41	20.451	5.373	7.207	33.031
16	279.018	24.936		24.819	49.755
32	379.674	35.208		26.258	61.466

Results – Threading on multiple nodes

- Using 4 threads has the best performance on 1 and 2 nodes
 - 8 threads better on 4 nodes
 - More threads means less processes
 - Potential for less communication overhead
 - Also may exploit threading in areas of code that are not suitable for MPI
- Always a good idea to run tests to see what value is best



Results – Communications

- Run time of mp_sum decreases with the number of threads
 - Possibly due to less messages being sent
- Run time of mp_alltoall less clear
 - Some improvement with using threads
 - 2-4 threads look to optimal
- Hard to predict how routines and the code as a whole will be affected by changing the number of threads
 - Really have to test this





Results – Communications

- Fraction of time spent in MP routines increases with nodes
 - Other areas of the program are improving performance at a greater rate, i.e.. they have greater potential for parallel speed up.
- Changing the number of threads has the potential to reduce this fraction



Results – QM system size

- The large QM system scales better, though perhaps not as well as you would think.
- The system with the large QM region spends a greater fraction of its time doing communications.





BioExcel Partners





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