



NWChem: Enabling cutting-edge open-source science from the teraflop to the exaflop

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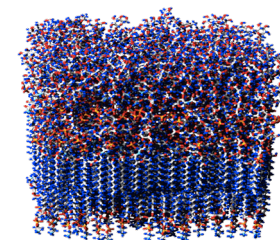
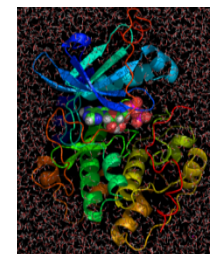
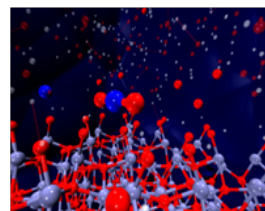
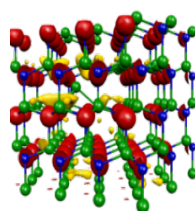
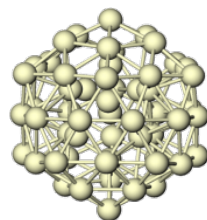
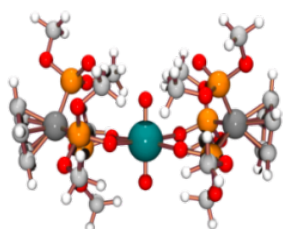


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Science driven software development



QM-CC

QM-DFT

AIMD

QM/MM

MM

- DOE's Premier computational chemistry software
- One-of-a-kind solution scalable with respect to scientific challenge and compute platforms
- From molecules and nanoparticles to solid state and biomolecular systems
- Portable – runs on a wide range of computers
 - ◆ Supercomputer to Mac or PC with Windows

<http://www.nwchem-sw.org>

NWChem is Open-Source



- NWChem consortium delivers code and infrastructure for computational chemistry community to build upon
- Establish more collaborative development environment
- Close to 55K downloads in under 2 years
- License is Educational Community License (ECL 2.0)
 - ◆ Apache style license



NWChem is expanding its developers community



Charge Transfer and Charge Transport in Photoactivated Systems

Developing Advanced Methods for Excited State Chemistry in the NWChem Software Suite

CAS/RASPT2 approaches

Approaches for ultrafast and excited state dynamics in solution, conformational sampling

Strong links with SciDAC SUPER and FASTMATH Institutes

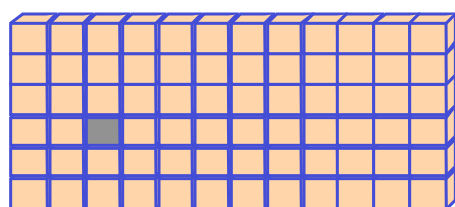
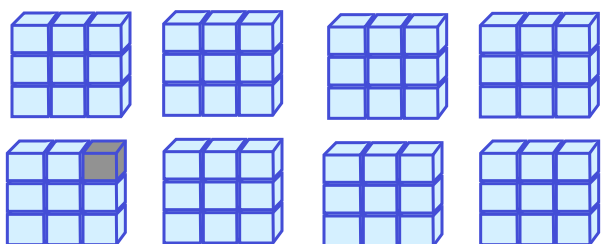


Parallel scalability in NWChem through Global Array Toolkit

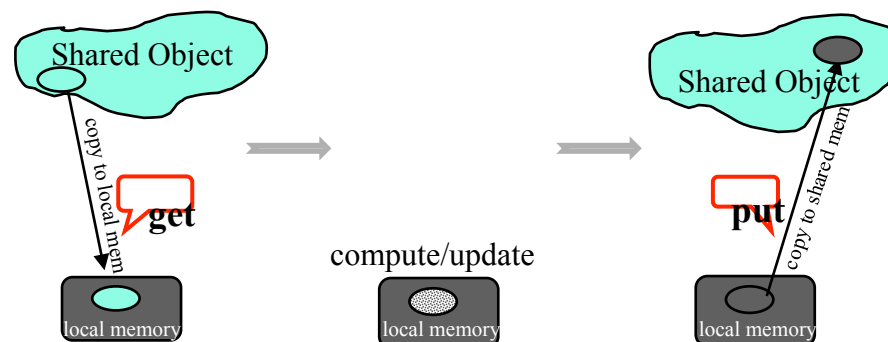


- NWChem uses Global Array Toolkit to scale and perform on parallel architectures
- Programming model: Distributed dense arrays that can be addressed through a shared memory-like style and one-sided access

Physically distributed data

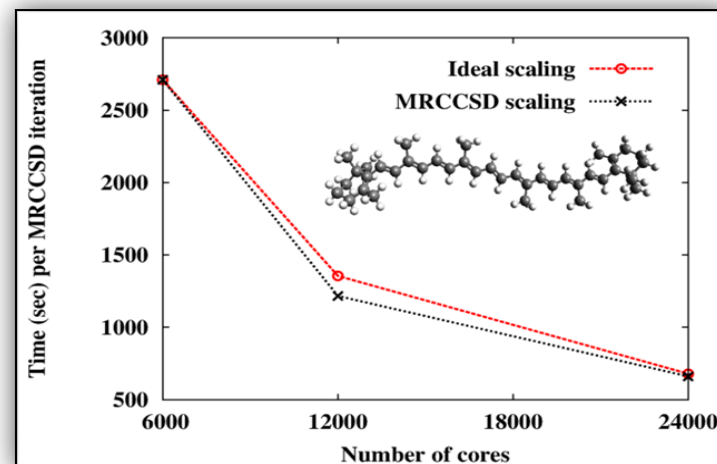
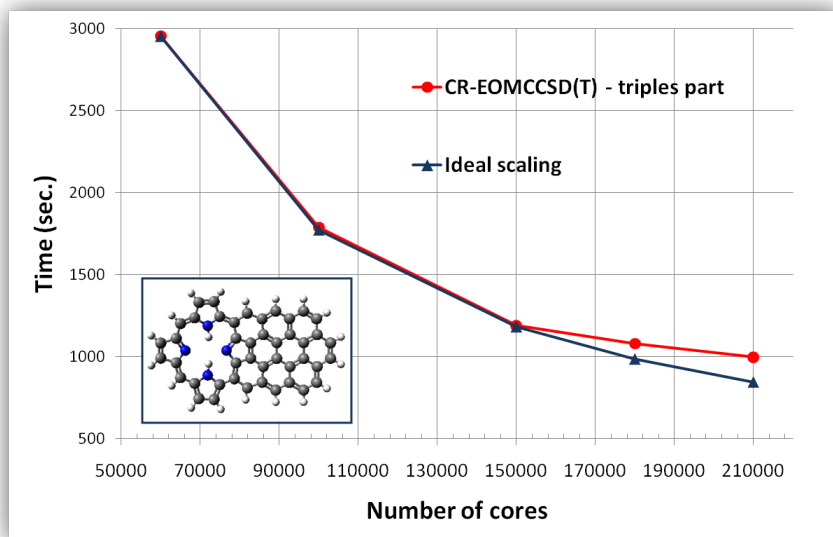


Global Address Space



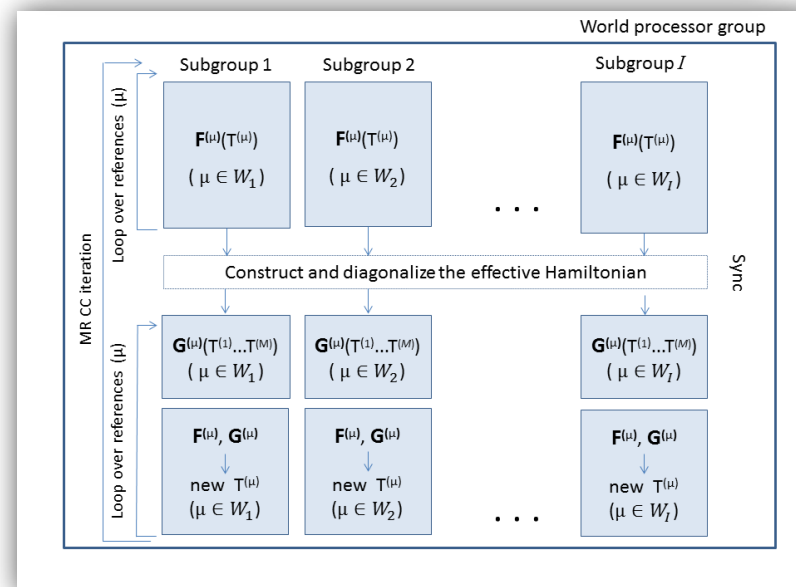
- Hides much of the necessary parallel infrastructure but awareness of locality of data for scalability
- Core-based model has to change change on exascale machines!!

Scalable single and multi-reference coupled cluster methods

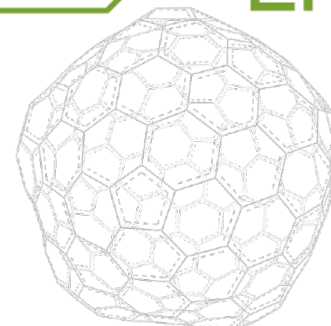


MRCC in NWChem

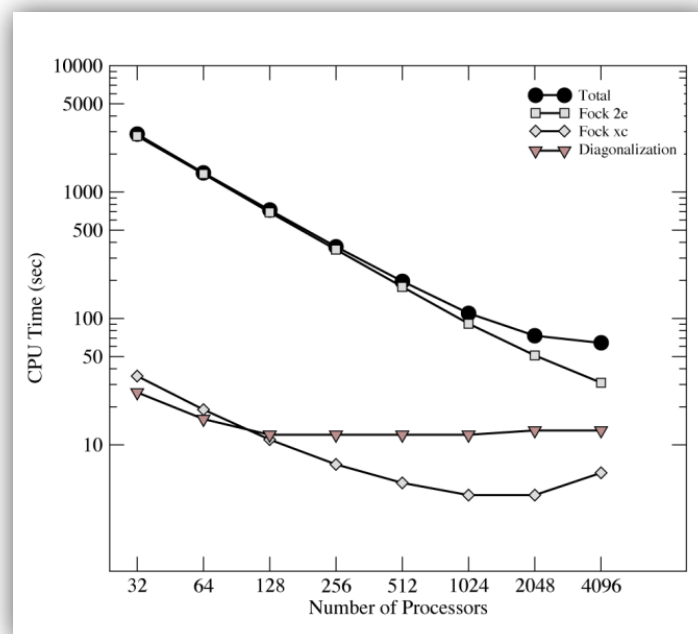
- ◆ Brillouin-Wigner MRCCSD
- ◆ Mukherjee Mk-MRCCSD
- ◆ State-Universal MRCCSD
- ◆ Perturbative triples corrections MRCCSD(T)
- ◆ Demonstrated scalability of MR-CCSD across 24,000 cores



Parallel Gaussian DFT performance



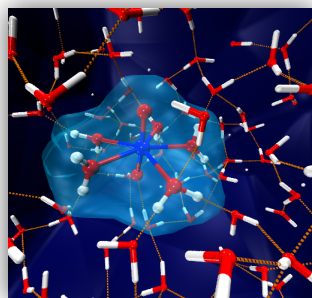
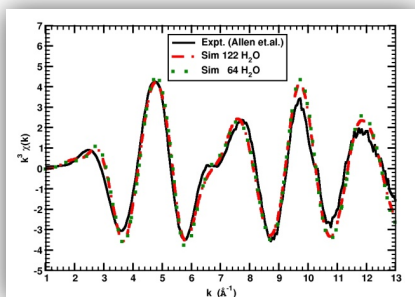
- Calculation on C_{240}
 - ◆ PBE0 functional, 6-31G*
 - ◆ Direct integral evaluation
 - ◆ Size 3600 basis functions
- Timings for different components of the Kohn-Sham matrix construction (scalar, vector, BLAS-2)
 - ◆ Fock 2e – two electron integrals
 - ◆ Fock xc – the DFT contribution
 - ◆ Diagonalization – eigenvector solver
- Scalability limited by diagonalization
 - ◆ Avoiding direct diagonalization, switched to quadratic convergence solver
- Other efforts are focused on
 - ◆ Reduced data movement
 - ◆ Reduced (linear scaling) approaches
 - ◆ Compute at lower precision



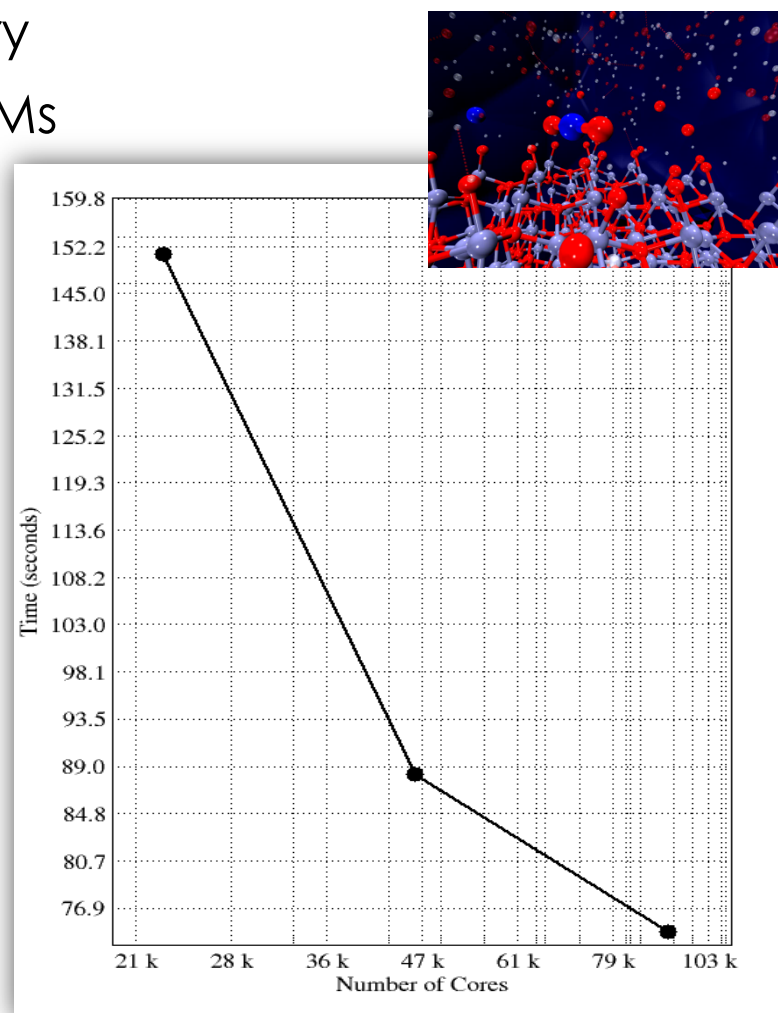
Plane wave DFT and dynamics for solution, surfaces, and materials



- Plane wave density functional theory
 - ◆ Driven by FFTs and global DGEMMs
- Extensive dynamics functionality with Car-Parrinello
 - ◆ Parallel-in-time development
- Various exchange-correlation functionals
 - ◆ Exact exchange is very efficient
 - ◆ Lots of FFTs here



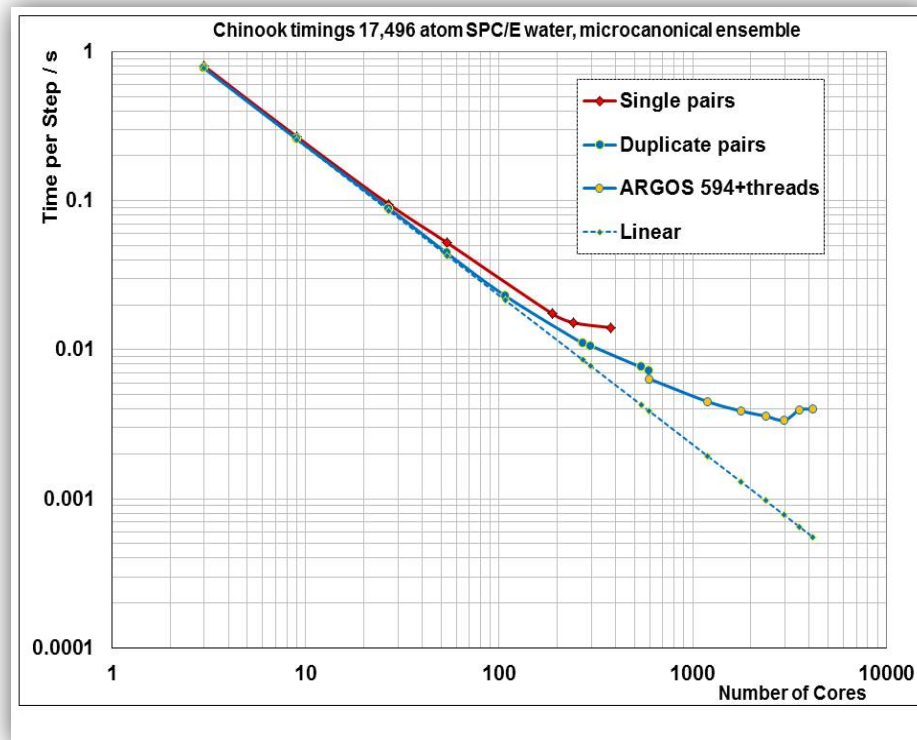
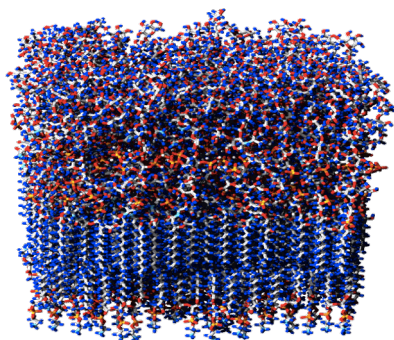
AIMD accurately models EXAFS of uranyl in water



Next-generation Molecular Dynamics



- Removal of all explicit global synchronizations from the basic MD time steps
- More effective hiding of communication latency and bandwidth through computation scheduling
- Improved scalability through cell-cell pair instead of cell distribution
 - ◆ Minimizing communication
 - ◆ Improved load balancing

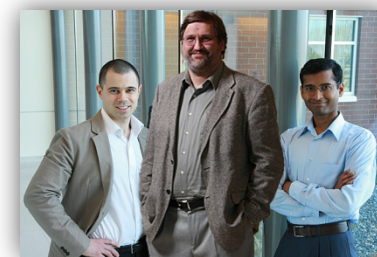
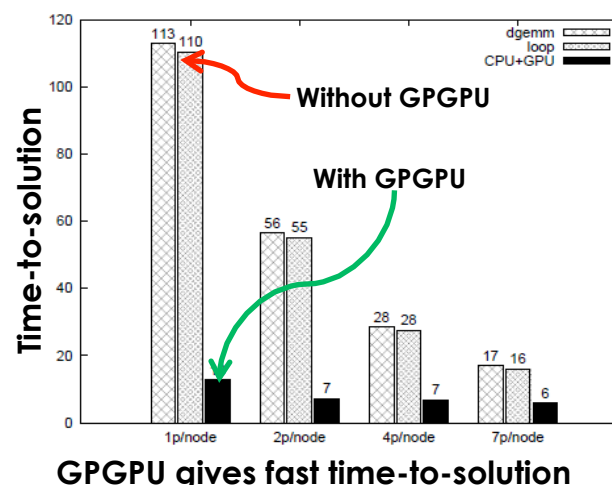
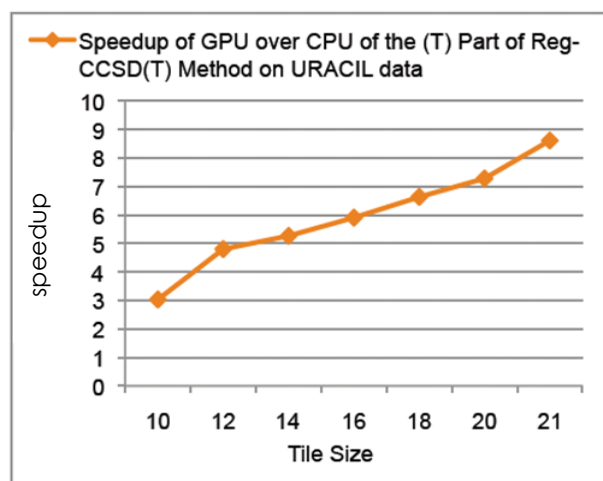


Scalability for 18,000 atoms
on 3,000 cores

That is only 6 atoms per core!

Coupled-cluster on GPGPUs

- The CCSD(T)/Reg-CCSD(T) codes have been rewritten in order to take advantage of GPGPU accelerators
- 8X speedup obtained by moving coupled cluster DGEMMs to GPGPU
 - ◆ Efficient overlap of data movement and computation is crucial for performance



Oreste Villa
Sriram Krishnamoorthy
Karol Kowalski

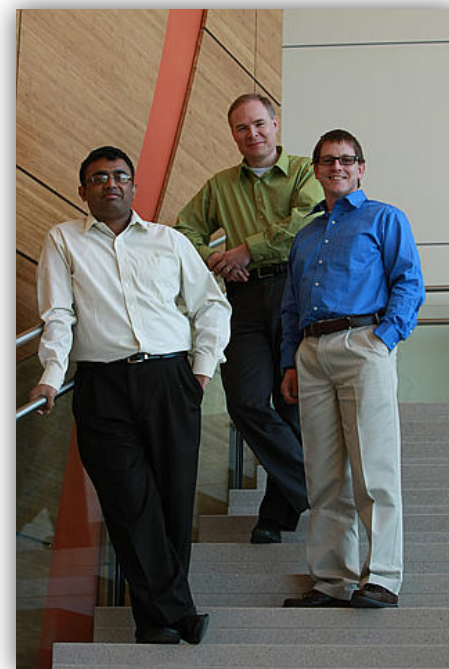
- Ongoing work on GPGPUs: Cholesky decomposition methods

Ma, Krishnamoorthy, Villa, Kowalski, JCTC 7,1316 (2011)

Handling faults in chemistry codes



- NWChem's compute model is task based and data centric:
 - ◆ Continue with (N-1) after failure
 - ◆ Ensure access to critical data through redundancy
- Reliability Benefits
 - ◆ Significantly increased MTBF; handle decent number of failures
 - ◆ Recovery proportional to degree of failure
 - ◆ Continued execution with existing nodes
- Overheads
 - ◆ Replication with synced writes - Critical data structures are significantly smaller

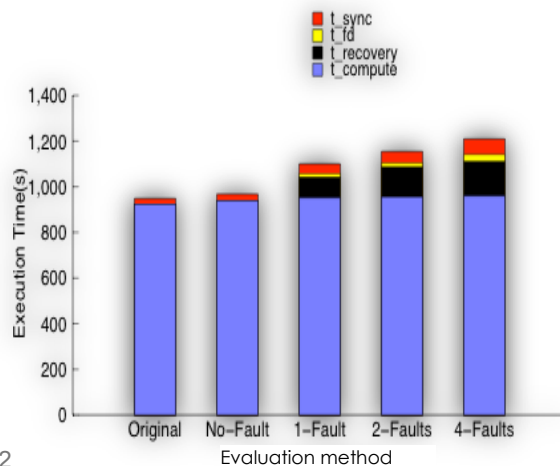
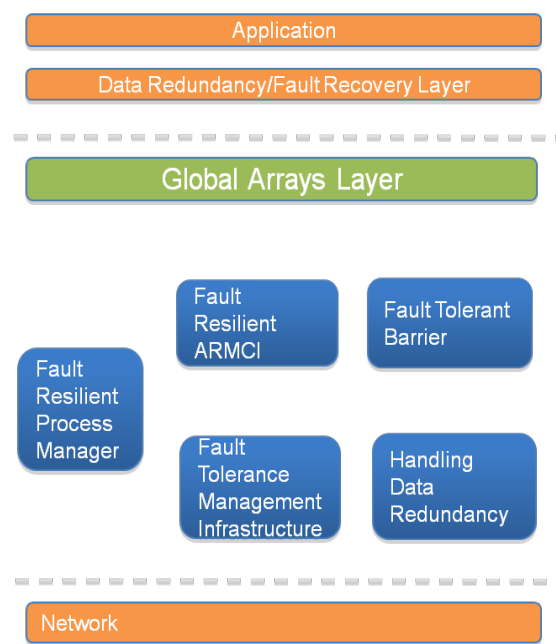


*Huub van Dam
Abhinav Vishnu
Bert de Jong*

NWChem CCSD(T) fault tolerant




Design of Fault Tolerant Communication Runtime System



- Handling faults increases scientific productivity and discovery
- Unique Fault Tolerant Management Infrastructure Developed
- Fault tolerance to hardware errors in NWChem demonstrated on various platforms
- We're now addressing soft errors within the Hartree-Fock algorithm
 - ◆ Using algorithmic numerical bounds
 - ◆ Work also drove us to look at reduced numerical precision is needed

van Dam, Vishnu, de Jong,
JCTC **7**, 66 (2011)


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Where we want to go with computational chemistry programming models



- Crucial to treat data and compute at the same footing
 - ◆ Moving from core to node as a basic data and task unit improves memory management
 - ◆ Cores within node performing different tasks

- Work load management
 - ◆ Next generation computational chemistry models will inherently be hierarchical
 - Dynamic task scheduling including futures
 - Tasks can spawn tasks (could include Python)
 - ◆ Avoid sync through data flow and task graphing
 - ◆ Move task to data and localization
 - ◆ Compute at minimally needed accuracy

NWChem development is funded by:

- US Department of Energy: BER, ASCR, BES**
- PNNL LDRD**

EMSL: A national scientific user facility integrating experimental and computational resources for discovery and technological innovation

