Implementation of First-Principles Molecular Dynamics on Large Scale Computers

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- Our goal: Simulate high-Z metals (Mo, Ta, Pu, ...) from first principles, without input from experiments
- The approach: Molecular dynamics: an atomic-scale simulation method
  - Compute the trajectories of all atoms
  - extract statistical information from the trajectories



Atoms move according to Newton's law:

$$m_i \ddot{\mathbf{R}}_i = \mathbf{F}_i$$



- Why "First-Principles"?
  - Avoid empirical models and adjustable parameters
    - Goal: applications to extreme conditions (high pressure, etc.) where no experimental data is available
  - Use fundamental principles: Quantum Mechanics
  - Must describe ions and electrons consistently and simultaneously







- The approach is applicable to very diverse
  - Chemistry
  - Nanotechnology
  - Semiconductors
  - Biochemistry
  - High-pressure physics





Biotin on silicon carbide



Silicon quantum dot



Growth of a carbon nanotube on an iron catalyst



• The computation of the electronic structure is the most expensive part of the simulation







# Computing the electronic structure

- Density Functional Theory: the Kohn-Sham equations
  - solutions represent molecular orbitals (one per electron)
  - molecular orbitals are complex scalar functions in R<sup>3</sup>
  - coupled, non-linear PDEs

$$\begin{cases} -\Delta \varphi_{i} + V(\rho, \mathbf{r})\varphi_{i} = \varepsilon_{i}\varphi_{i} & i = 1... N_{el} \\ V(\rho, \mathbf{r}) = V_{ion} (\mathbf{r}) + \int \frac{\rho(\mathbf{r}')}{|\mathbf{r} - \mathbf{r}'|} d\mathbf{r}' + V_{XC} (\rho(\mathbf{r}), \nabla \rho(\mathbf{r})) \\ \rho(\mathbf{r}) = \sum_{i=1}^{N_{el}} |\varphi_{i}(\mathbf{r})|^{2} \\ \int \varphi_{i}^{*}(\mathbf{r}) \varphi_{j}(\mathbf{r}) d\mathbf{r} = \delta_{ij} \end{cases}$$



# Computing the electronic structure

Periodic boundary conditions: all solutions of the form

$$\psi_{\mathbf{k},n}(\mathbf{r}) = e^{i\mathbf{k}\cdot\mathbf{r}}\varphi_{\mathbf{k},n}(\mathbf{r})$$

must be included (Bloch theorem)

• Solutions are represented as Fourier series

$$\varphi_{\mathbf{k},n}(\mathbf{r}) = \sum_{|\mathbf{k}+\mathbf{q}|^2 < E_{\text{cut}}} c_{\mathbf{k}+\mathbf{q},n} e^{i\mathbf{q}\cdot\mathbf{r}}$$

• Electronic charge density

$$\rho(\mathbf{r}) = \sum_{n} \int_{BZ} \left| \psi_{\mathbf{k},n}(\mathbf{r}) \right|^2 d^3 \mathbf{k}$$



# Computing the electronic structure

 A periodic solution is represented by the matrix of complex Fourier coefficients c<sub>qn</sub>

$$\varphi_n(\mathbf{r}) = \sum_{|\mathbf{q}|^2 < E_{\text{cut}}} c_{\mathbf{q},n} e^{i\mathbf{q}\cdot\mathbf{r}}$$

- The matrix of coefficients c<sub>qn</sub> must have orthogonal columns
- Dimensions of C: 10<sup>6</sup>x10<sup>4</sup>



# **Controlling Numerical Errors**

- The goal is high accuracy
- All numerical errors must be controlled
  - Convergence of Fourier series  $\varphi_n(\mathbf{r}) = \sum_{|\mathbf{q}|^2 < E_{\text{cut}}} c_{\mathbf{q},n} e^{i\mathbf{q}\cdot\mathbf{r}}$
  - Convergence of system size (number of atoms)
  - Convergence of k-space integration

$$\rho(\mathbf{r}) = \int_{BZ} \left| \psi_{\mathbf{k},n}(\mathbf{r}) \right|^2 d^3 \mathbf{k}$$

- We need to systematically increase
  - Plane-wave energy cutoff
  - Number of atoms

# BlueGene/L allows us to ensure convergence of all three approximations



# Algorithms used in FPMD



- Solving the KS equations: a constrained optimization problem in the space of coefficients c<sub>qn</sub>
- Poisson equation: 3-D FFTs
- Computation of the electronic charge: 3-D FFTs
- Orthogonality constraints require dense, complex linear algebra (e.g. A = C<sup>H</sup>C)



### The Platform: BlueGene/L

- 65,536 nodes, 128k CPUs
- 3D torus netwc
- 512 MB/node
- 367 TFlop peak





### **Obox code: main features**

- C++/MPI implementation of First-Principles Molecular Dynamics
- DFT/GGA exchange-correlation
- Plane-wave, norm-conserving pseudopotentials
- Designed for large-scale parallel platforms
- Main design constraint: small memory footprint (< 512MB per task)</li>
- Built on optimized parallel libs: PBLAS, ScaLAPACK
- XML interface
- Used on various parallel platforms (BG/Ls, Linux clusters)



#### **Qbox code structure**





### Data layout

• Distributed plane-wave coefficients

$$\varphi_{n\mathbf{k}}(\mathbf{r}) = \sum_{|\mathbf{k}+\mathbf{G}|^2 < E_{\text{cut}}} c_{n,\mathbf{k}+\mathbf{G}} e^{i(\mathbf{k}+\mathbf{G})\cdot\mathbf{r}} \mathbf{G}$$



# Data layout

- Single k-point wavefunction:
  - ScaLAPACK matrix block distribution

$$\varphi_{n\mathbf{k}}(\mathbf{r}) = \sum_{|\mathbf{k}+\mathbf{G}|^2 < E_{\text{cut}}} c_{n,\mathbf{k}+\mathbf{G}} e^{i(\mathbf{k}+\mathbf{G})\cdot\mathbf{r}}$$

- Dimensions of C: 10<sup>6</sup>x10<sup>4</sup>
- Typical process grid: 512x16



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#### **Communication** patterns

• 3-D Fourier transforms  $\varphi_{n\mathbf{k}}(\mathbf{r}) \leftrightarrow c_{n,\mathbf{k}+\mathbf{G}}$ 





#### **Communication** patterns

• Accumulation of electronic charge density





### **Communication** patterns

- Other operations: (orthogonalization, non-local potential energy, Ritz diagonalization)
  - use the ScaLAPACK linear algebra library





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# Mapping tasks to physical nodes

• Mapping a 2-D process grid to a 3-D torus





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### Single-node kernels



- Dual core
- Dual FPU
- Three-level cache memory hierarchy
- L1 caches not coherent
- L2, L3 coherent

# Single-node kernels

- Exploiting the BG/L hardware
  - Use double FPU instructions ("double hummer")
  - Use both CPUs on the node
    - use virtual node mode, or
    - program for two cores
  - We use BG/L in co-processor mode
    - 1 MPI task per node
    - Use second core using dual-core kernels
- DGEMM/ZGEMM kernel (John Gunnels, IBM)
  - Hand optimized, uses double FPU very efficiently
  - Algorithm tailored to make best use of L1/L2/L3
  - Dual-core version available: uses all 4 FPUs on the node
- FFTW kernel (Technical University of Vienna)
  - Uses hand-coded intrinsics for DFPU instructions



#### ZGEMM Performance

- ZGEMM on one processor per node
  - In excess of 98% of peak (2.76/2.80 GF/node)
- ZGEMM in co-processor mode
  - 97% of peak (5.43/5.60 GF/node)
  - Uses fork/join construct
  - Flushes caches to maintain coherence
- Kernel performance
  - Over 99.5% of peak



# The Test Problem

- Electronic structure of a 1000-atom Molybdenum sample
- p semi-core electrons included
- 12,000 electrons
- 32 non-local projectors for pseudopotentials
- 112 Ry plane-wave energy cutoff
- High-accuracy parameters





#### **Obox performance results**

- Single k-point calculation (k=0)
- single-core dgemm library
- co-processor mode
- bipartite y-z mapping at 64k nodes



#### 46.70 TFlops on 64k nodes





#### **Obox performance results**

- Single k-point calculation (k=0)
- dual-core dgemm library
- co-processor mode
- bipartite y-z mapping



#### 64.0 TFlops on 64k nodes





#### **Obox performance results**

- Multiple k-point calculations 1 k-point: 108.8 TFlop/s (30% of peak)
- complex arithmetic
- dual-core zgemm library
- co-processor mode

4 k-points: 187.7 TFlop/s (51% of peak)









Node mapping significantly affects performance



# Node Mapping Optimization

- Analysis of the MPI tree broadcast algorithm in sub-communicators
  - The performance of MPI\_Bcast is unsatisfactory for checkerboard mappings: the tree algorithm results in communication bottlenecks
  - Using a space-filling curve to assign tasks to nodes improves performance
  - performance gain: ~7 TFlops



- unnecessary Type\_commit operations were removed in the BLACS
- performance gain: ~3 TFlops





# History of First-Principles MD performance



## History of FPMD performance





# Computational cost of firstprinciples simulations

• The computational cost of solving the Kohn-Sham equations is  $O(N^3)$ 

$$\begin{cases} -\Delta \varphi_i + V(\rho, \mathbf{r}) \varphi_i = \varepsilon_i \varphi_i & i = 1...N_{el} \\ V(\rho, \mathbf{r}) = V_{ion}(\mathbf{r}) + \int \frac{\rho(\mathbf{r}')}{|\mathbf{r} - \mathbf{r}'|} d\mathbf{r}' + V_{XC}(\rho(\mathbf{r}), \nabla \rho(\mathbf{r})) \\ \rho(\mathbf{r}) = \sum_{i=1}^{N_{el}} |\varphi_i(\mathbf{r})|^2 \\ \int \varphi_i^*(\mathbf{r}) \varphi_j(\mathbf{r}) d\mathbf{r} = \delta_{ij} \end{cases}$$



# O(N<sup>3</sup>) algorithms onlarge parallel platforms

- Storage of electronic wavefunctions: O(N<sup>2</sup>) (N wavefunctions described in a volume O(N))
- Number of operations: O(N<sup>3</sup>) (orthogonalization)

Problem size	# ops	# CPUs	ops/CPU	storage	storage/CPU
Ν	N <sup>3</sup>	Ν	N <sup>2</sup>	N <sup>2</sup>	Ν
Ν	N <sup>3</sup>	N <sup>2</sup>	N	N <sup>2</sup>	1
Ν	N <sup>3</sup>	N <sup>3</sup>	1	N <sup>2</sup>	1/N



# Linear-scaling methods

- Achieve *linear scaling: O(N)* operations
- Introduce approximations to reduce the computational cost from *O(N<sup>3</sup>)* to *O(N)*.
- Several approaches proposed in the past 10 years
- Most successful approach: represent the solutions of the Kohn-Sham equations in terms of nonorthogonal, localized functions.
- An important goal is *controlled accuracy*, i.e.
  - Simple parameters (e.g. grid spacing) to control numerical accuracy
  - As robust as *O(N<sup>3</sup>)* methods



### Linear-scaling methods

 Domain decomposition approach: wavefunctions are localized in spherical, overlapping domains.

$$S_{ij} = \left\langle \phi_i \left| \phi_j \right\rangle = \int_{\Omega} \phi_i^*(\mathbf{r}) \phi_j(\mathbf{r}) d^3 \mathbf{r} \right\rangle$$
$$H_{ij} = \left\langle \phi_i \left| H \phi_j \right\rangle = \int_{\Omega} \phi_i^*(\mathbf{r}) H \phi_j(\mathbf{r}) d^3 \mathbf{r} \right\rangle$$



 $E(Y) = \operatorname{tr}(S^{-1}Y^T HY) \quad Y \in R^{M \times N} \quad S = Y^T Y$ 



#### Localization of orbitals

 Spherical domains are attached to atoms or bonds



 "Linear-Scaling First-Principles Molecular Dynamics with Controlled Accuracy", J.L.Fattebert and F.Gygi, Comp. Phys. Comm. 162, 24 (2004)

# O(N) with controlled accuracy

- Errors in computed ionic forces are decaying exponentially for large localization radii
- Errors are computed by comparison with O(N<sup>3</sup>) method with same numerical approximations





# O(N) with controlled accuracy

• The accuracy of ionic forces is critical for molecular dynamics simulations



J.L. Fattebert and F. Gygi, Phys. Rev. B 73, 115124 (2006)



# MD: using adaptive localization centers (ALC)

- Spherical domains move during the MD simulation
- The positions of the centers are recalculated at each time step using the center of charge of each orbital



J.L. Fattebert and F. Gygi, Phys. Rev. B 73, 115124 (2006)



# O(N) Molecular Dynamics

H<sub>2</sub>O 32 molecules MD simulation with adaptive localization centers



FIG. 2. Total energy and ALC displacements during molecular dynamics simulation of water (512 molecules) at 300 K with localization radius of 9 Bohr.



# Controlling energy drift in MD

• H<sub>2</sub>O 32 molecules MD simulation

TABLE I. Measure of energy drift and number of SC steps required for convergence.

Localization radius (Bohr)	Energy drift (mHa/at/ps)	No. SC iteration/ MD step
8	–0.222 (–47 K/ps)	29
9	–0.103 (–22 K/ps)	23
10	+0.001 (0 K/ps)	14



# Comparison with O(N<sup>3</sup>): simulations of liquid water

Timings for 1 electronic step (processors Itanium 2, 1.4 GHz, Quadrics switch)





# Linear-scaling methods

- Review articles:
  - G.Galli, "Linear Scaling Methods For Electronic Structure Calculations and Quantum Molecular Dynamics Calculations", *Current Opinion in Solid State and Materials Science*, 1, 864 (1996).
  - S.Goedecker, "Linear Scaling Electronic Structure Methods", *Rev. Mod. Phys.* 71, 1085 (1999).
- Our recent work on *O(N)* with controlled accuracy
  - J.L.Fattebert and F.Gygi, Comp. Phys. Comm. 162, 24 (2004).
  - J.L. Fattebert and F. Gygi, Phys. Rev. B 73, 115124 (2006)



# Linear-scaling methods

- Outstanding remaining issues
  - The choice of localization radii is difficult a priori
    - Some orbitals may require larger radii than others
    - The description of unoccupied orbitals is more difficult than occupied orbitals
  - The description of metallic systems is not satisfactory
  - The initial choice of localization centers is not obvious
    - Use atoms in some systems, bonds in others
  - Moving localization centers in a smooth, continuous way

More research is needed to develop linear-scaling electronic structure methods that are suitable for MD simulations



# Summary

- High-performance implementations of First-Principles Molecular Dynamics can be developed on large scale computers (up to 128k CPUs)
- Highly tuned single-node kernels are used
- Node mapping is critical for this application
- Linear scaling (O(N)) FPMD is the subject of more investigations to achieve *controlled accuracy*

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