

Max Webinar

BigDFT

*The operators of BigDFT code. From convolutions to Poisson Solver in High Performance Computing*

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Virtual Room

BigDFT Real-Space  
approach

Luigi Genovese

Wavelets and ISF

GPU

Wavelet Convolutions

Optimization Approach

BOAST

Challenges

Poisson Solver

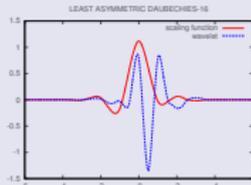
Implicit Solvents

Summary

Daubechies  $f(x) = \sum_{\mu} c_{\mu} \phi_{\mu}(x)$

Orthogonal set

$$c_{\mu} = \int dx \phi_{\mu}(x) f(x) \equiv \langle \phi_{\mu} | f \rangle$$

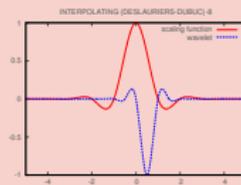


No need to calculate overlap matrix of basis functions  
Used for **wavefunctions**,  
**scalar products**

Interpolating  $f(x) = \sum_j f_j \phi_j(x)$

Dual to dirac deltas (... is it?)

$$f_j = f(j)$$



The expansion coefficients are the point values on a grid  
Used for **charge density**,  
**function products**

Magic Filter method (A. Neelov, S. Goedecker)

The passage between the two basis sets can be performed without losing accuracy  $c_{\mu} = \sum_k w_{\mu-k} f_k$ ,  $w_k = \langle \phi | L_k \rangle$ .

## The SCF cycle

Orbital scheme:

- Hamiltonian
- Preconditioner

Coefficient Scheme:

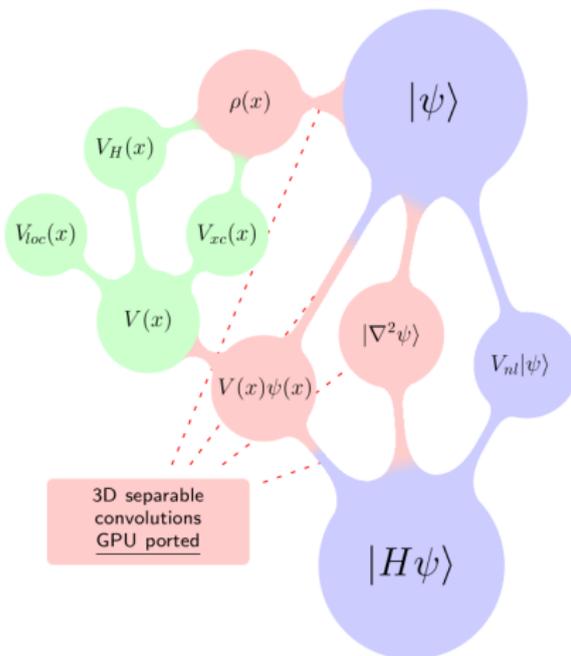
- Overlap matrices
- Orthogonalisation

## Comput. operations

- **Convolutions**
- **BLAS routines**
- FFT (Poisson Solver)

Real Space

Daub. Wavelets



Why not GPUs?

Wavelets and ISF

GPU

- Wavelet Convolutions
- Optimization Approach
- BOAST
- Challenges

Poisson Solver

- Implicit Solvents
- Summary

## A DFT code conceived with a HPC mindset

- DFT calculations up to many thousands atoms
- An award-winning HPC code
- BigDFT has been conceived for massively parallel etherogeneous architectures since more than 10 years (MPI + OpenMP + GPU)



## Code able to run routinely on different architectures

- GPGPU since the advent of double-precision (2009)
- Itanium, BG-P and BG-Q (Incite award 2015)
- ARM architectures (Mont Blanc I)
- KNL Accelerators (Marconi)
- K computer (RIKEN) Fugaku – preparatory
- European Supercomputers (Archer, IRENE-Rome, ...)

## Developer and user dilemmas

- Does my code fit well? For which systems?
- How much does porting cost?
- Should I always use GPUs?
- How can I interpret results?

## Evaluating GPU convenience

### Three levels of evaluation

1. Bare speedups: GPU kernels vs. CPU routines  
Are the operations suitable for GPU?
2. Full code speedup on one process  
Amdahl's law: are there hot-spot operations?
3. Speedup in a (massively?) parallel environment  
The MPI layer adds an extra level of complexity

## Initially, naive routines (FORTRAN?)

$$y(j, l) = \sum_{\ell=L}^U h_{\ell} x(l + \ell, j)$$

- Easy to write and debug
- Define reference results

```

do j=1, ndat
  do i=0, n1
    tt=0.d0
    do l=lowfil, lupfil
      tt=tt+x(i+l, j)*h(l)
    enddo
    y(j, i)=tt
  enddo
enddo

```

## Optimisation can then start (2010: X5550, 2.67 GHz)

Method	GFlop/s	% of peak	SpeedUp
Naive (FORTRAN)	0.54	5.1	1/(6.25)
<b>Current (FORTRAN)</b>	<b>3.3</b>	<b>31</b>	<b>1</b>
Best (C, SSE)	7.8	73	2.3
<b>OpenCL (Fermi)</b>	<b>97</b>	<b>20</b>	<b>29 (12.4)</b>

A trade-off between benefit and effort

## FORTRAN based

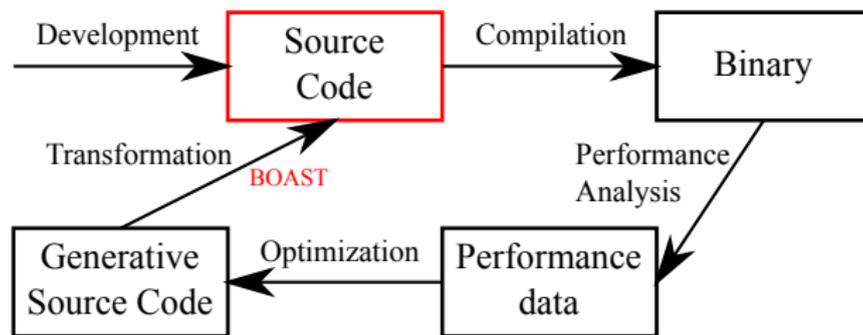
- ✓ Relatively accessible (loop unrolling)
- ✓ Moderate optimisation can be achieved relatively fast
- ✗ Compilers fail to use vector engine efficiently

## Push optimisation at the best

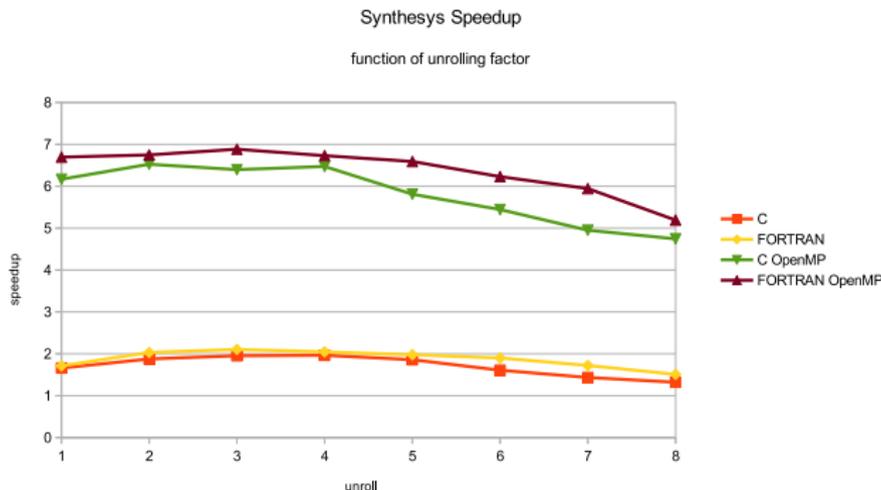
- Only one out of 3 convolution type has been dissected
  - About 20 different patterns have been studied for one 1D convolution
  - Tedious work, huge code → Maintainability?
- 👉 Automatic code generation with BOAST engine

## Metaprogramming the kernels

Optimization hand-in-hand with *design* of the operations



- Generate combination of optimizations
- C, OpenCL, FORTRAN and CUDA are supported
- Compilation and analysis are automated
- Selection of best version can also be automated



## Take-home messages of the BOAST strategy

- Meta-programming of Hot-spot operations
- Optimization can be adapted to the characteristics
- Portability, best effort and maintainance may go together
- Towards a BLAS-like library for wavelets convolutions

## Not all codes can benefit from BLAS/LAPACK/FFT

- Domain specific libraries need to be optimized for the architecture
- Autotuning is possible through the usage of tools like BOAST
- Generality can be achieved through meta-programming

## Example: libconv

- BOAST can generate all wavelet families  $\Rightarrow$  the library has more functionalities than the original code
- BOAST can adapt the routines to the architecture using OpenMP and vector instructions
- BOAST can target FORTRAN and C with OpenMP, CUDA, OpenCL

- Less data → **lower** number of flops per atom
- Communication scheduling **completely** different

## Different problems 🖱️ new opportunities

- No hot spot operations anymore!  
Convolutions are a less important percentage of the run
- Linear algebra becomes sparse  
Extra complexity in handling parallelisation

## The time-to-solution is considerably improved

18 k Atoms → less than 20 minutes on 13k cores

- The computational physicist is happier
- What about the computer scientist?

## (Screened) Poisson Equation for **any BC in vacuum**

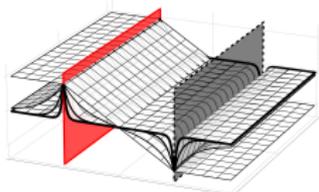
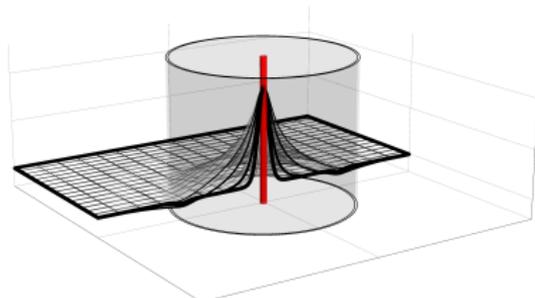
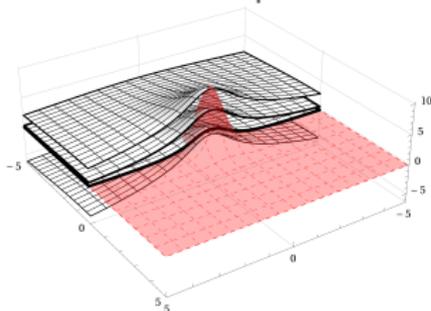
Non-orthorhombic cells (periodic, surface BC):

$$(\nabla^2 - \mu_0^2)V(x, y, z) = -4\pi\rho(x, y, z)$$

Machine-precision accuracy

J. Chem. Phys. **137**, 13 (2012)

Extended to implicit solvents (JCP 144, 014103 (2016))



## Future developments

Range-separated Coulomb operator

$$\frac{1}{r} \left[ \operatorname{erf} \frac{r}{r_0} + \operatorname{erfc} \frac{r}{r_0} \right]$$

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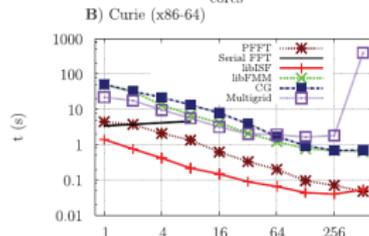
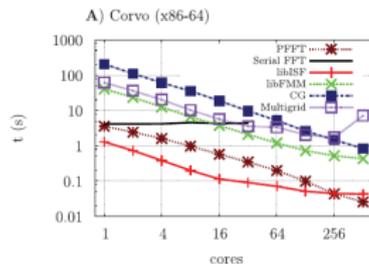
## A Survey of the Parallel Performance and Accuracy of Poisson Solvers for Electronic Structure Calculations

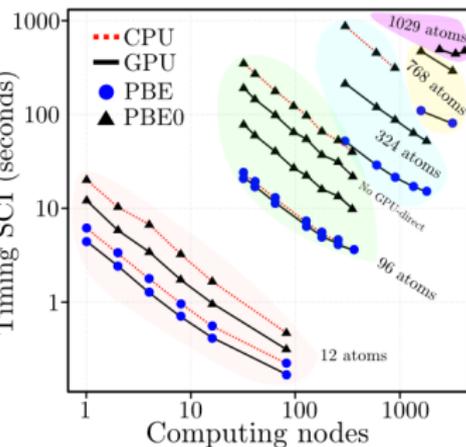
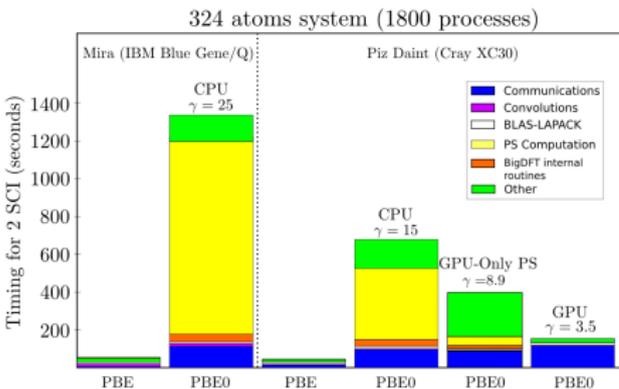
Pablo García-Risueño,<sup>\*[a,b,c]</sup> Joseba Alberdi-Rodriguez,<sup>[d,e]</sup> Micael J. T. Oliveira,<sup>[f]</sup> Xavier Andrade,<sup>[g]</sup> Michael Pippig,<sup>[h]</sup> Javier Muguerza,<sup>[d]</sup> Agustin Arruabarrena,<sup>[d]</sup> and Angel Rubio<sup>[e,i]</sup>

We present an analysis of different methods to calculate the classical electrostatic Hartree potential created by charge distributions. Our goal is to provide the reader with an estimation on the performance—in terms of both numerical complexity and accuracy—of popular Poisson solvers, and to give an intuitive idea on the way these solvers operate. Highly parallelizable routines have

been implemented in a first-principle simulation code (OCTOPUS) to be used in our tests, so that reliable conclusions about the capability of methods to tackle large systems in cluster computing can be obtained from our work. © 2013 Wiley Periodicals, Inc.

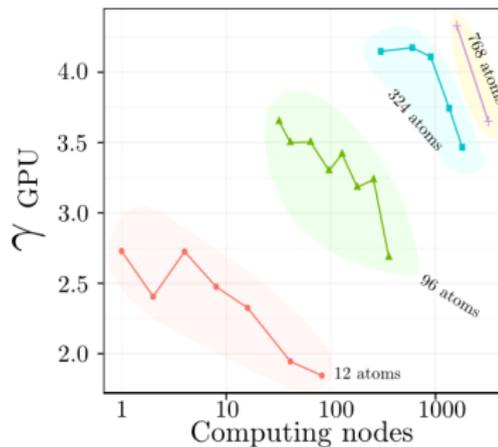
DOI: 10.1002/jcc.23487





**UO<sub>2</sub> systems:**

Atoms	Orbitals
12	200
96	1432
324	5400
768	12800
1029	17150



## Poisson solver for implicit solvents

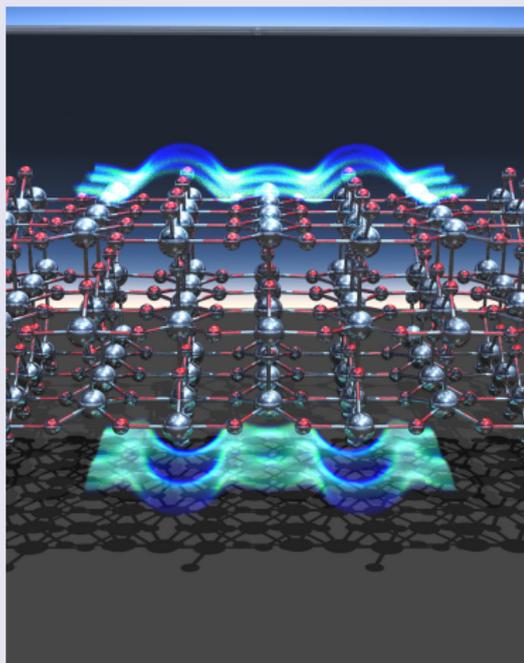
JCP 144, 014103 (2016)

Allows an efficient and accurate treatment of implicit solvents

The dielectric function determine the **cavity** where the solute is defined.

The cavity can be

- rigid (PCM-like)
  - determined from the Electronic Density (SCCS approach)
- ☞ Can treat various BC  
(here  $\text{TiO}_2$  surface)



Can be used in conjunction with  $O(N)$  BigDFT

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Neutral or ionic wet environments

Generalized Poisson eq.

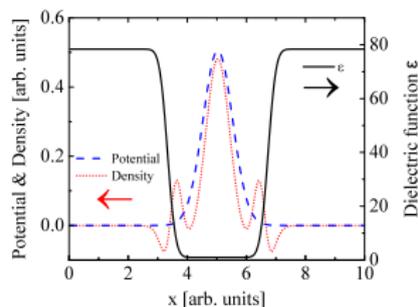
$$\nabla \cdot \varepsilon(\mathbf{r}) \nabla \phi(\mathbf{r}) = -4\pi \rho(\mathbf{r})$$

Poisson-Boltzmann eq.

$$\nabla \cdot \varepsilon(\mathbf{r}) \nabla \phi(\mathbf{r}) = -4\pi [\rho(\mathbf{r}) + \rho^{ions}[\phi](\mathbf{r})]$$

Various algorithms implemented

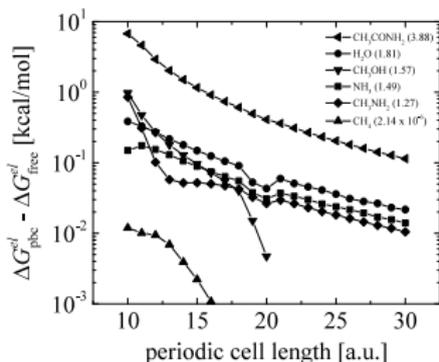
- Polarization Iteration (use gradient of polarization potential)
- Preconditioned Conjugate Gradient (need only function multiplications)



Solution found by iterative application of PS in vacuum BC

## Advantages of explicit BC

Difference between the electrostatic solvation energy computed with periodic and free boundary conditions as function of the periodic cell length.



## Molecule dipole norm (D)

	vacuum	H <sub>2</sub> O
CH <sub>3</sub> CONH <sub>2</sub>	3.88	5.76
H <sub>2</sub> O	1.81	2.41
CH <sub>3</sub> OH	1.57	2.14
NH <sub>3</sub>	1.49	1.98
CH <sub>3</sub> NH <sub>2</sub>	1.27	1.78

Explicit BC avoid error due to supercell aliasing

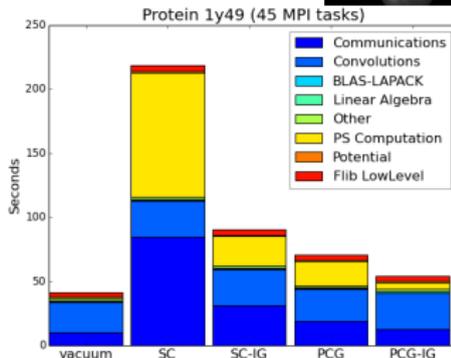
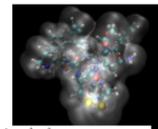
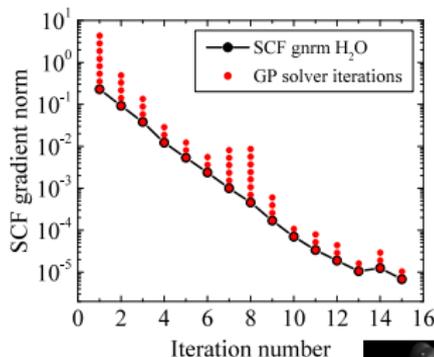
## Blackbox-like usage

The Generalized PS only needs few iterations of the vacuum poisson solver

## Time-to-solution

Timings for the protein PDB ID: 1y49 (122 atoms) in water

- Full SCF convergence 49 s
- Solvent/vacuum runtime ratio  $\alpha = 1.16$



## Features of Systematicity

- ISF provide a rigorous framework to interpret real-space coefficients
- High interpolating power (precise results)
- Scaling relation  $\rightarrow$  computational efficiency, arbitrary precision
- (Quasi) variational
- **In conjunction with (Daubechies) wavelets offer a good formalism to get rid of uncertainties**

## Opportunities from new quadratures

- Generalize the concept of **compensating charge** (multipoles)
- Combine together cartesian and polar meshes