

**Solvers: diagonalization, OMM,
PEXSI, CheSS. Parallelization issues.**



**Barcelona
Supercomputing
Center**

Centro Nacional de Supercomputación

**Solvers: diagonalization, OMM,
PEXSI, CheSS.**

Calculating the density kernel

A typical SCF cycle in DFT codes looks as follows:

1 calculate

- Hamiltonian matrix $H_{\alpha\beta} = \langle \phi_\alpha | \mathcal{H} | \phi_\beta \rangle$
- overlap matrix $S_{\alpha\beta} = \langle \phi_\alpha | \phi_\beta \rangle$

2 calculate the density kernel **K** out of **H** and **S**

3 calculate

- energy $E = Tr(\mathbf{KH})$
- electronic density $\rho(\mathbf{r}) = \sum_{\alpha,\beta} \phi_\alpha^*(\mathbf{r}) K_{\alpha\beta} \phi_\beta(\mathbf{r})$

4 update the Hamiltonian operator \mathcal{H} according to the new electronic density ρ

5 start over again

Limiting factor in most calculations (in particular big ones) with SIESTA:
Calculation of the density matrix



SIESTA offers various solvers to calculate the density kernel:

- Diagonalization (various flavors)
- Orbital minimization method (OMM)
- PEXSI
- CheSS

The choice of the most suited method depends on the specific calculation:

- system size
- sparsity of the matrices
- HOMO-LUMO gap
- used basis set
- etc.



Most straightforward approach to calculate the density kernel:

- 1 solve the generalized eigenvalue problem $\mathbf{H}\mathbf{c}_i = \epsilon_i \mathbf{S}\mathbf{c}_i$
- 2 calculate the density kernel as $K = \sum_i \mathbf{c}_i \mathbf{c}_i^T$

Advantages:

- exact calculation without any approximations
- universally applicable
- there exist highly optimized libraries

Shortcomings:

- possible sparsity of the matrices cannot be exploited
- cubic scaling with system size
- hard to parallelize

Various flavors for the diagonalization

There are various libraries to diagonalize a matrix:

- ScaLAPACK: parallel version of LAPACK
 - most popular library for dense general purpose linear algebra
 - various diagonalization algorithms:
 - PDSYEV: based on tridiagonal QR iteration
 - PDSYEV D: based on Divide and Conquer algorithm
 - PDSYEV X: based on Bisection and Inverse Iteration
 - PDSYEV R: based on the parallel MRRR algorithm
 - often limited in parallel performance
- ELPA
 - better performance than ScaLAPACK using the same API
- MAGMA (Matrix Algebra on GPU and Multicore Architectures)
 - linear algebra library for heterogeneous architectures (CPU, Xeon Phi, GPU)
 - Has interfaces to LAPACK and ScaLAPACK routines, so easy to port

In SIESTA:

- ScaLAPACK (various flavors)
- ELPA

Orbital minimization method

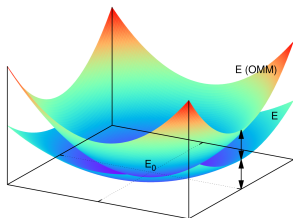
Find $n = N/2$ Wannier functions describing the occupied subspace by direct unconstrained minimization

- The original OMM functional [F. Mauri et al., Phys. Rev. B 47, 9973 (1993); P. Ordejón et al., Phys. Rev. B 48, 14646 (1993)]:

$$\tilde{E} = 2 \text{Tr}\{[\mathbf{I} + (\mathbf{I} - \mathbf{S})]\mathbf{H}\}, \quad \text{with } S_{ij} = \langle \psi_i | \psi_j \rangle, H_{ij} = \langle \psi_i | \mathcal{H} | \psi_j \rangle$$

- For orthonormal set ($\mathbf{S} = \mathbf{I}$): $\tilde{E} = E = \text{Tr}(\mathbf{H})$
- For all other cases: $\tilde{E} \geq E$ if \mathbf{H} is negative definite (easily fulfilled by shift)

Unconstrained (i.e. no explicit orthogonalization) global minimum of the function \tilde{E} coincides with E

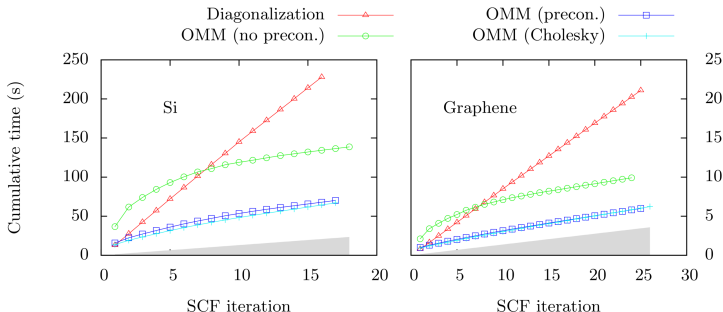


Orbital minimization method – results

OMM has the potential to be a $\mathcal{O}(N)$ method, even if the version implemented in SIESTA is not (no localization constraints).

Still an interesting alternative to diagonalization

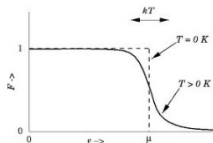
- At each SCF iteration, the results from the previous one can be reused as starting guess
- Especially towards the end of the SCF cycle potentially faster than diagonalization





The density matrix can be calculated directly from the Hamiltonian:

$$\mathbf{K} = f(\mathbf{H}), \quad \text{with } f(\epsilon) = \frac{1}{1 + e^{\beta(\epsilon - \mu)}}$$



All we need is a computationally convenient representation of the Fermi function f .

Two possibilities:

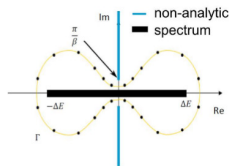
- rational expansion: PEXSI
- Chebyshev expansion: CheSS



Approximate the Fermi function using a pole expansion in the complex plane:

$$f(\epsilon) \approx \text{Im} \sum_{l=1}^{n_p} \frac{w_l}{\epsilon - (z_l + \mu)}$$

$$K \approx \text{Im} \sum_{l=1}^{n_p} \frac{w_l}{\mathbf{H} - (z_l + \mu)\mathbf{S}}$$



Advantages:

- Only a small number of poles is required (typically about 40)
- Each pole independent

Most important task: Inversion of the matrices $\mathbf{H} - (z_l + \mu)\mathbf{S}$
 \implies done with the Selected Inversion algorithm



- PEXSI only calculates those elements of the density matrix which are required to calculate physical quantities (charge density, energy, forces, ...)
- Thus PEXSI can exploit the sparsity of the matrices (consequence of the localized character of the basis set $\{\phi_\alpha\}$)

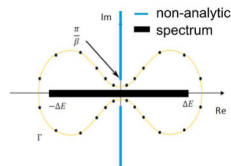
Exact method (no approximations!), but reduced scaling.

For sufficiently big problems:

- 1-dimensional: $\mathcal{O}(N)$
- 2-dimensional: $\mathcal{O}(N^{3/2})$
- 3-dimensional: $\mathcal{O}(N^2)$

Number of poles depends on the inverse electronic temperature β and the spectral width ΔE .

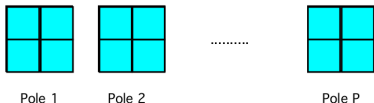
However fast convergence with the number of poles
 \Rightarrow still applicable to metals!



PEXSI – Scaling



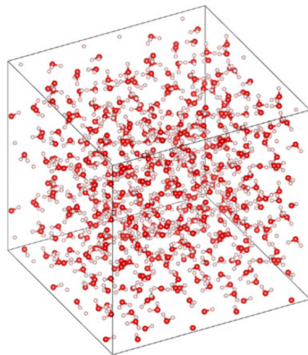
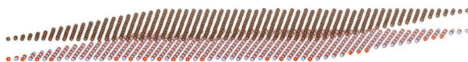
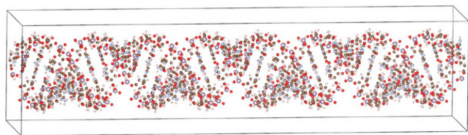
PEXSI exhibits ideal parallel scaling with respect to the poles:



40 processors per pole \times 40 poles:
160 processors

Test systems:

- 1D: DNA (up to 17875 atoms)
- 2D: C–BN (up to 12770 atoms)
- 3D: H₂O (up to 24000 atoms)

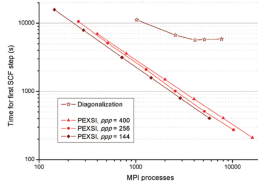
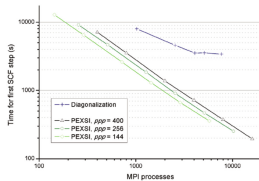
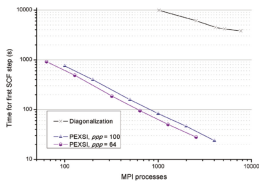
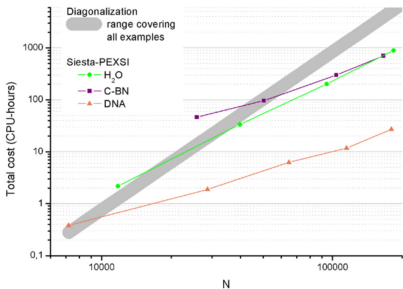


PEXSI – Scaling



Fitted slopes:

- DNA: 1.3 ; C–BN: 1.7 ; H₂O: 2.2
- Deviations from perfect scaling:
Due to parallelization issues
- Prefactor: “sparsity” of the system
- For large systems always faster than diagonalization



$\mathcal{O}(N)$ schemes

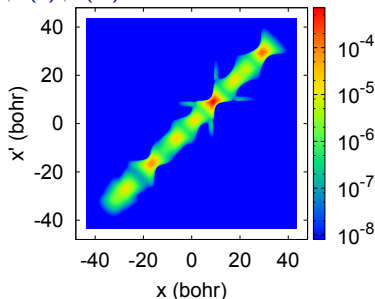


The key for $\mathcal{O}(N)$ schemes is locality: “nearsightedness” principle
(W. Kohn, Phys. Rev. Lett. **76**, 3168 (1996))

Example: Density matrix $F(\mathbf{r}, \mathbf{r}') = \sum_i f_i \psi_i(\mathbf{r}) \psi_i(\mathbf{r}')$

The matrix elements $F(\mathbf{r}, \mathbf{r}')$ decay rapidly with $|\mathbf{r} - \mathbf{r}'|$:

- insulators and metals at finite temperature: exponentially
- metals at zero temperature: algebraically



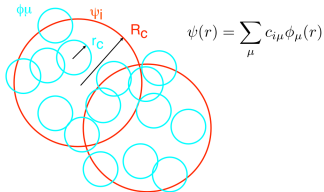
The decay length depends on the HOMO-LUMO gap of the system.
 $\mathcal{O}(N)$ schemes thus work best for insulators.

$\mathcal{O}(N)$ schemes



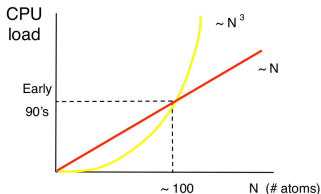
Basic idea: Localization

- Confine the orbitals within a sphere with cutoff R_c .
- Equivalent: Enforce the density matrix F to be sparse.



Justified by the nearsightedness.

$\mathcal{O}(N)$ usually have a larger prefactor than classical $\mathcal{O}(N^3)$ calculations
 \Rightarrow crossover point





Basic idea: Approximate the density matrix by a polynomial expansion:

- Use Chebyshev polynomials to avoid instabilities
- Shift and scale \mathbf{H} such that eigenvalues lie in the range $[-1, 1]$ ($\bar{\mathbf{H}}$)
- Calculate the density matrix as

$$\mathbf{K} = f(\mathbf{H}) \approx \frac{c_0}{2} \mathbf{I} + \sum_{i=1}^{n_{pl}} c_i \mathbf{T}^i(\bar{\mathbf{H}})$$

Efficient and flexible approach:

- The coefficients c_i can be cheaply calculated using textbook formulas (also for other expansions than the density matrix, e.g. the inverse)
- The Chebyshev polynomials fulfill the following recursion relation:

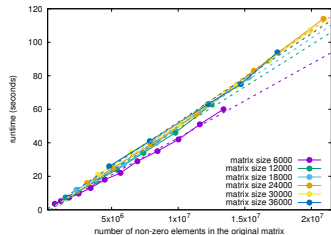
$$\mathbf{T}^0(\bar{\mathbf{H}}) = \mathbf{I} \quad ; \quad \mathbf{T}^1(\bar{\mathbf{H}}) = \bar{\mathbf{H}} \quad ; \quad \mathbf{T}^{j+1}(\bar{\mathbf{H}}) = 2\bar{\mathbf{H}}\mathbf{T}^j(\bar{\mathbf{H}}) - \mathbf{T}^{j-1}(\bar{\mathbf{H}}).$$

From this we see:

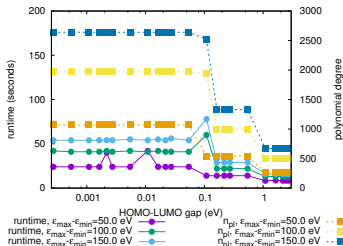
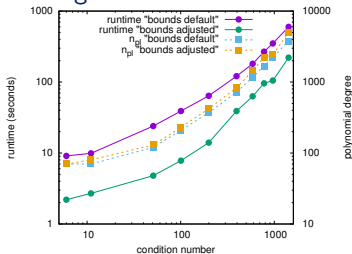
- Only matrix vector multiplications required, easily parallelizable
- $\mathcal{O}(N)$ method by restricting the multiplications to a sparsity pattern



The performance of CheSS only depends on the number of non-zero entries of the matrices

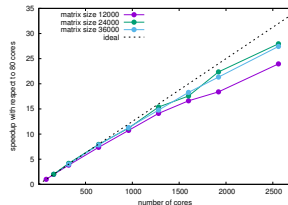


The algorithm works best for matrices with a small eigenvalue spectrum:

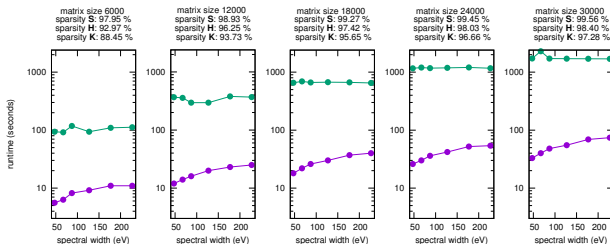




CheSS shows a very good parallel scaling (for both MPI and OpenMP)



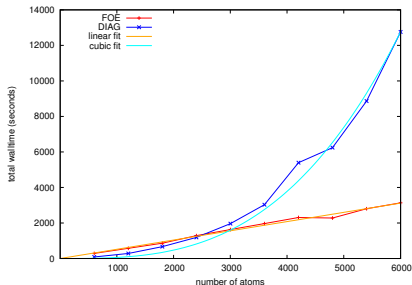
For hybrid MPI/OpenMP setup, CheSS can outperform PEXSI for appropriate systems (small eigenvalue spectrum, non-zero HOMO-LUMO gap)





SIESTA has been interfaced with CheSS (not yet in the official version).

Very simple test system:
alkane chain, DZP basis



Most important remaining bottleneck: Reduce the spectral width of the SIESTA matrices (sort of contracted basis set?)

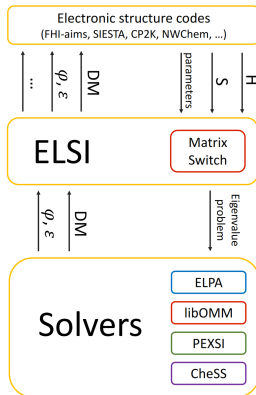


The ELSI project (ELectronic Structure Infrastructure) is an effort to unite various of the aforementioned solvers:

- ELPA
- OMM
- PEXSI
- CheSS

Ultimate goal:

- Provide one single interface to all these libraries
- A DFT code interfacing ELSI gets access to all these methods
- Easy case-by-case choice of the solver



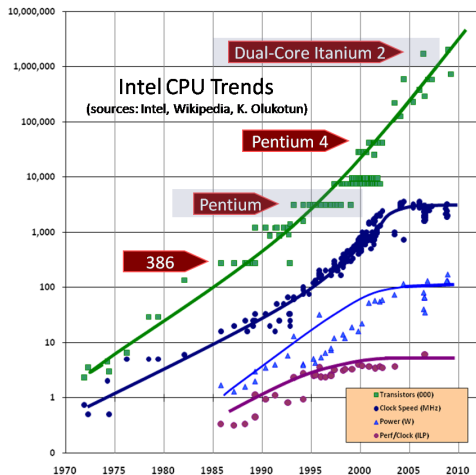
Parallelization issues

Need for parallelization

Clock speed of a single core saturates since about 2005

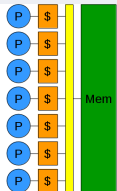
Performance gain only possible by using more cores at the same time

Applications must be highly parallelized

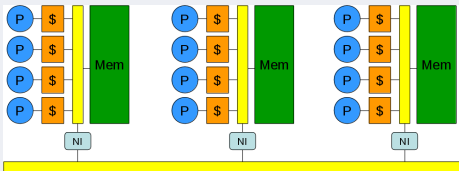


Types of parallelism

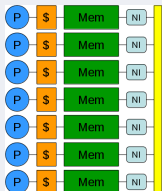
shared memory



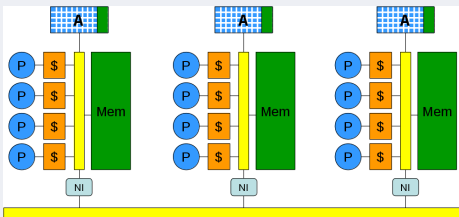
hybrid distributed/shared memory



distributed memory



accelerators



Characteristics of the parallelization schemes

Shared memory:

- done using OpenMP
- can often be added relatively easily on top of existing code (but getting good performance is not that easy!)
- limiting factor often memory bandwidth
- maximal speedup limited by number of cores per node

Distributed memory:

- distributing data is essential for large applications
- done using MPI (Message Passing Interface)
- introducing MPI often requires a major refactoring of the code
- limiting factor is the communication
- maximal speedup depends on application and architecture, in principle no upper bound

Characteristics of the parallelization schemes

Hybrid distributed/shared memory:

- combination of MPI and OpenMP
- most complex form of parallelism, careful implementation required
- allows to increase the maximal speedup (multiplicative)
- allows to overcome the aforementioned constraints and limitation (e.g. memory, bandwidth, etc.)

Accelerators:

- often used to accelerate specific intensive parts of the code
- most popular ones: GPUs and MICs
- can be combined with other parallelization schemes
- maximal speedup limited by accelerator
- usually hard work to get good performance



Parallel resources can be exploited by SIESTA in various ways:

- various levels of distributed memory parallelization using MPI:
 - k-point parallelism
 - Distributing orbitals and gridpoints
- recently shared memory using OpenMP was added
- parallelization of the various external solver:
 - BLAS: OpenMP
 - ScaLAPACK: MPI and OpenMP
 - PEXSI: heavy two-level MPI parallelization, limited OpenMP parallelization
 - CheSS: efficient MPI and OpenMP parallelization



Compile options:

```
## MPI wrappers to compiler
FC=mpif90

## Compile with OpenMP
FFLAGS= -fopenmp

## Flag activating MPI
FPPFLAGS= -DMPI

## Parallel linear algebra libraries
LIBS = <your_scalapack_lib> <your_blacs_lib>

## additional external libraries
LIBS += <your_chess_lib>

## additional preprocessor flags
FPPFLAGS += -DSIESTA__CHESS

MPI_INTERFACE=libmpi_f90.a
MPI_INCLUDE=.
```

Parallel execution



Execution on local workstation:

```
mpirun -n 8 siesta < example.fdf
```

Execution on cluster

Submit script (e.g. submit.sh)

```
#!/bin/bash
#BSUB -J Siesta
#BSUB -n 16
#BSUB -oo output_%J.out
#BSUB -eo output_%J.err
#BSUB -R "span[ptile=16]"
#BSUB -W 00:20
<load required modules>
mpirun siesta < example.fdf
```

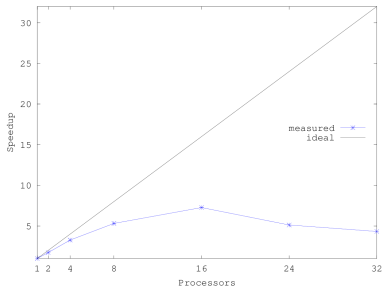
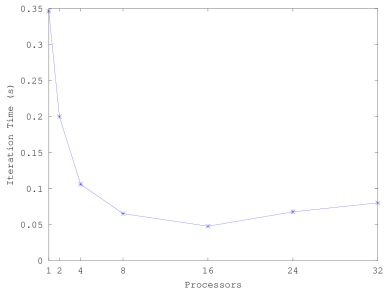
Submission on cluster

```
bsub < submit.sh
```

Parallel scaling: Time and speedup

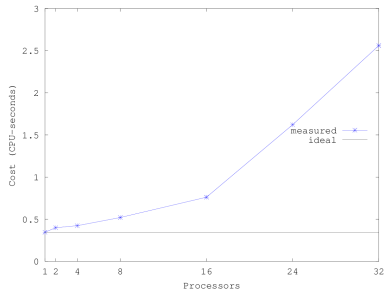
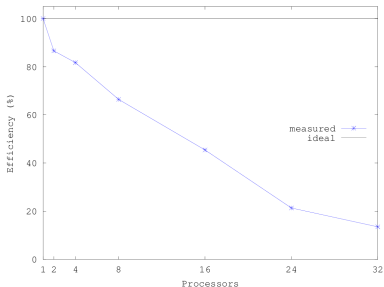
Using more cores does NOT always make a calculation faster (e.g. communication overhead)!

Careful estimation of the parallel resources is requested.



Parallel scaling: Efficiency and cost

Choosing a bad parallelization scheme strongly affects the efficiency and cost:



To get a rough estimate of the parallel performance, see also


<http://departments.icmab.es/leem/siesta/siestimator/siestimator.php>

Access to parallel computers

- 1 Clusters at universities, research institutes, companies, ...
- 2 National networks
e.g. in Spain: RES <http://www.res.es/>
- 3 European networks
e.g. PRACE: <http://www.prace-ri.eu>
 - “Preparatory access” for testing, benchmarks, etc.
 - “Project access for” production runs

RES and PRACE:

- Regular calls
- Significance of project and scalability of software
- Grants certain amount of CPU-hours (free of charge)



Thank you for your attention!