

# Sakurai-Sugiura algorithm based eigenvalue solver for Siesta

Georg Huhs

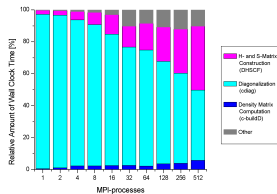
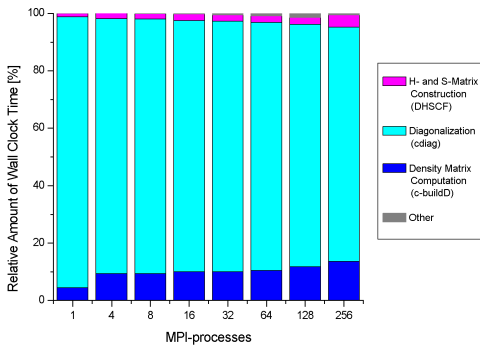


**Barcelona  
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Centro Nacional de Supercomputación

# Motivation



Timing analysis for one SCF-loop iteration:



left: CNT/Graphene, right: DNA

# Siesta

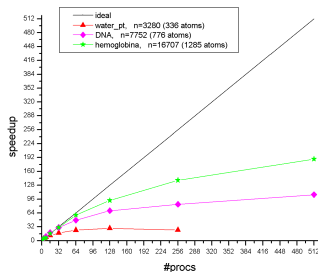


## Specifics

- High fraction of EVs needed (e.g. 30%)
- Sparse matrices

## Actual solver: ScaLAPACK

- Dense solver
- Limited scaling



# Siesta

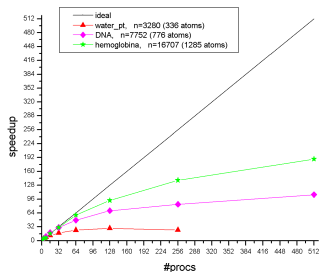


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# Characteristics



SS-method, Tetsuya Sakurai and Hiroshi Sugiura, 2002

- Solver for GEV  $A\mathbf{x} = \lambda B\mathbf{x}$   
 $A, B \in \mathbb{C}^{n \times n}$
- Projection method, based on complex contour integration
- Meant for finding EVs in a certain domain
- Suitable for
  - large, sparse matrices
  - parallelisation

# Basic steps



Divide domain of interest into subdomains, in each:

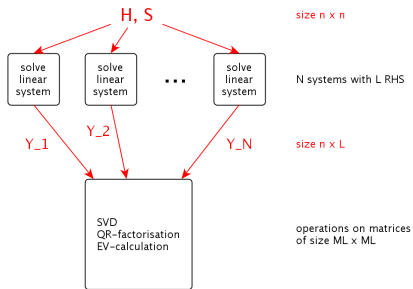
- 1 Solve a set of linear systems
- 2 Use results for numerical integrations
- 3 Construct subspace
- 4 Find eigenvalues/-vectors in subspace and backtransform
- 5 Select correct eigenpairs

# Parallelisation



## Three levels of parallelisation

- 1 Handling subdomains in parallel
- 2 Solving  $N$  linear systems in parallel
- 3 Use parallel solver for each linear system

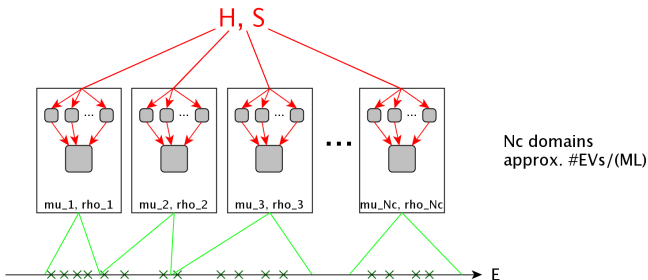


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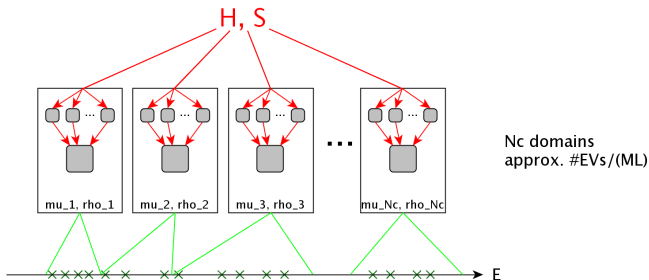


# Parallelisation



## Three levels of parallelisation

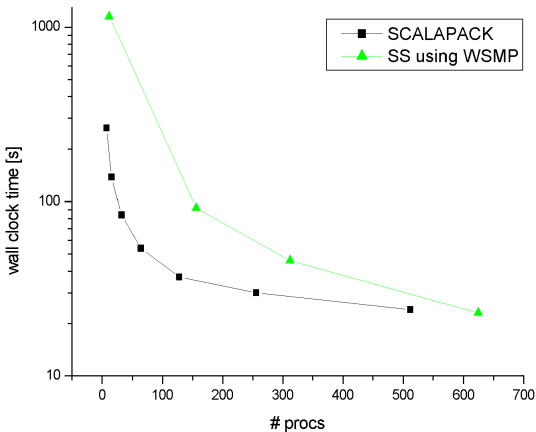
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# Timing expectations



Example: DNA, 776 atoms,  $n = 7752$



# Using GPUs



**Implemented:** Solving the linear systems with CUSP.

**Algorithm:** gmres without preconditioner!

**Gain:** Speedup of 3-4 compared to WSMP (for the smaller examples)

Timing solving all lin. systems in parallel (per subdomain)

Problem	Setup	Solve	Comm	total	ScaLAPACK
water_pt ( $n = 3280$ )	4.0	3.7	0.1	11	7
DNA ( $n = 7752$ )	3.3	7.6	0.5	13	$\sim 20$
hemoglobina ( $n = 16707$ )	8.0	10	1.0	<b>22</b>	<b><math>\sim 70</math></b>

All times in seconds

# Conclusions



## Issues

- Inefficient in serial mode
- Memory parallelisation only in linear solver

## Benefits

- Robust algorithm, parameterisation can be hidden from user
- **Actual problems:** SS can be faster than ScaLAPACK, depending on scaling of lin. solver
- **Future problems:** high potential for larger numbers of processors
- Possibility to use many GPUs efficiently

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# Thanks to

Alberto García, Pablo Ordejón (Siesta)

Tetsuya Sakurai (Algorithm)

Anshul Gupta (WSMP)

BSC support team

José María Cela (BSC)

# More details (1)



- 1 Define EV-range  $\Rightarrow \gamma, \rho$  and a set of vectors  $V \in \mathbb{C}^{n \times L}$ .
- 2 Set quadrature points:

$$z_j = \gamma + \rho e^{(2\pi i(j + \frac{1}{2}) \frac{1}{N})}, \quad j = 0, \dots, N - 1$$

- 3 Solve the linear systems

$$(z_j B - A) Y_j = B V$$

- $N$  systems,  $L$  right hand sides each
- independent!

# More details (2)



## 4 Numerical integration

$$S_k = \frac{1}{N} \sum_{j=0}^{N/2-1} \left( \frac{z_j - \gamma}{\rho} \right)^{k+1} Y_j, \quad k = 0, \dots, M-1$$

Matrix  $S = [S_0, S_1, \dots, S_{M-1}] \in \mathbb{C}^{n \times ML}$

## 5 Find the rank $K$ of $S$ with an SVD and shrink it to $n \times K$

## 6 QR-decomposition of $S \Rightarrow$ orthogonal basis $Q$

## 7 $\tilde{A} = Q^T(A - \gamma B)Q$

$$\tilde{B} = Q^T B Q$$

$$\tilde{A}, \tilde{B} \in \mathbb{C}^{K \times K}$$

## 8 Eigenpairs of $\tilde{A}\tilde{x} = \tilde{\lambda}\tilde{B}\tilde{x}$



# More details (3)



- 8 Eigenpairs of  $\tilde{A}\tilde{\mathbf{x}} = \tilde{\lambda}\tilde{B}\tilde{\mathbf{x}}$
- 9 Eigenpairs of the original problem

$$\lambda_j = \tilde{\lambda}_j + \gamma$$
$$\mathbf{x}_j = Q \cdot \tilde{\mathbf{x}}_j$$

- 10 Remove exterior eigenvalues and ghosts.