

# Hierarchic Data Structures for Sparse Matrix Representation in Large-scale DFT/HF Calculations

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# Talk Outline

Hierarchic  
Matrices

P. Salek

Sparse QM

Error Control

HMLib

Parallelization

- Running large-scale ab-initio calculations – scaling.
- Solving problems with sparse matrices.
- OpenMP parallelization – problems.

# Performance of the SCF Cycle

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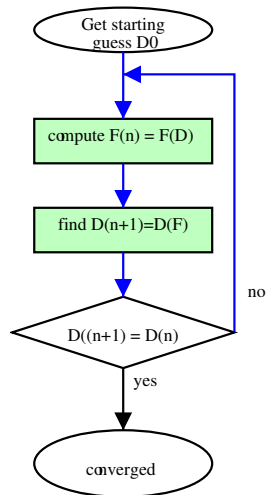
Sparse QM

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Parallelization

- Computation of Kohn-Sham matrix  $F$  is time-consuming.
- For really large systems, density evaluation ( $F \rightarrow D$ ) is time-consuming as well.
- Matrix memory usage grows quadratically.
- Local basis set – basis functions localized on atoms.



# Research Group

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Parallelization

- Elias Rudberg: Interaction evaluation (PhD in December).
- Emanuel Rubensson: Sparse matrices.

# $F \rightarrow D$ Step

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Density traditionally obtained via diagonalization and *aufbau* principle:

$$FC = \epsilon SC \quad D = C_{\text{occ}} C_{\text{occ}}^T$$

- Diagonalization does not scale linearly.
- Density optimization and purification algorithms scale linearly when sparsity is used.

# Sparse Matrices in Quantum Chemistry

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Parallelization

- Matrices are used to represent operators  $D$  and  $F$ .
- Overlap matrix, Density matrix, Fock Matrix, Kohn-Sham matrix.
- Matrices must be represented in such a way that common operations are fast.
- Sparsity appears only for larger molecules ( $> 50$  atoms).

# Sparsity Patterns

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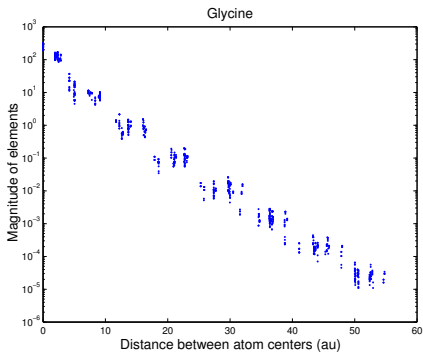
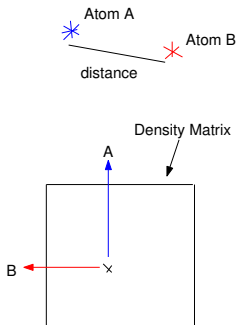
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Matrix sparsity depends on the basis set and geometry and to some extent on the band gap.

# Taking Advantage of Sparsity Patterns

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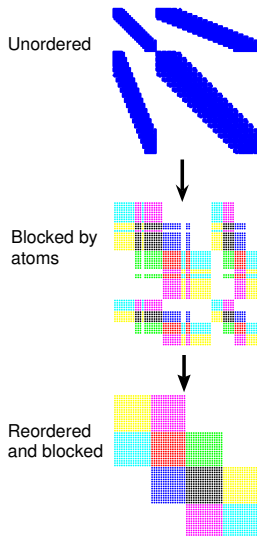
Error Control

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Parallelization

- Sparsity appears in blocks.
- Reorder atoms to merge the atom blocks in larger ones.
- Use BLAS for operations on blocks and Compressed-Sparse Row (CSR) format for block storage.
- Enforce sparsity by small element truncation.

## Alkane chain Fock matrix





# $F \rightarrow D$ Step With Sparse Matrices

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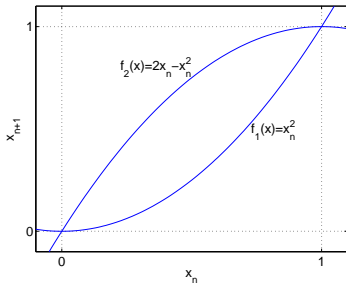
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Parallelization

- Use Trace-Correcting Purification – a series of spectral transformations.
- Performance limited by the sparse matrix multiplication speed.



```
compute P = (lmax I-F)/(lmax-lmin)
while abs(trace(P)-N)>threshold
  if(trace(P)>N) then
    P := P*P
  else
    P := 2*P-P*P
end while
```

# Example TC2 Application

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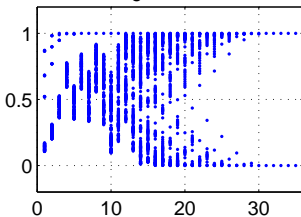
Error Control

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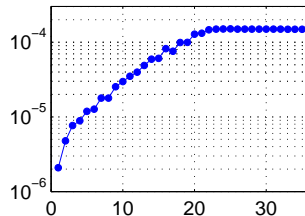
Parallelization

## Glycine Molecule

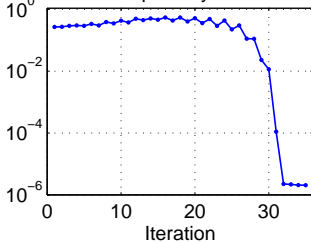
Eigenvalues



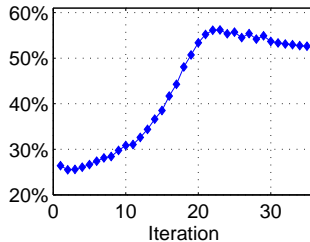
Accumulated error



Idempotency error



Share of nonzero elements



# Problems with TC2

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- Number of TC2 iterations depends the bandgap.
- Control the error: TC2 error grows exponentially with the number of iterations.
- More flexible representation than Compressed Sparse Row is needed for easy implementation of other algorithms.

# Systematic Small-Submatrix Selection Algorithm

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- Error introduced by truncation of small elements.
- First approaches considered only distance between atoms and empirical threshold factors – unreliable!
- More advanced approaches look at the norms of neglected blocks – more reliable but strict error control still impossible.
- **SSSA looks at the error of the entire matrix. Provides strict error control.**

# The Effect of SSSA on Total Energy Error

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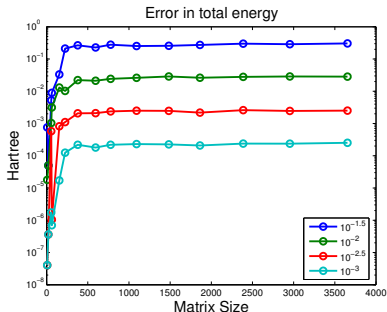
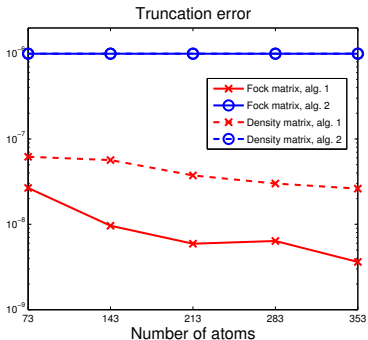
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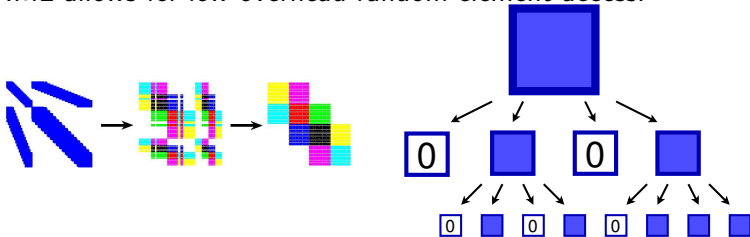
Benchmark on water clusters with Hartree-Fock and STO-3G.  
alg. 1: threshold based filtering, alg. 2: SSSA.



SSSA provides rigorous error control. → Saves time and gives trustworthy results. Energy extrapolation possible.

# Hierarchic Matrix Library (C++)

HML allows for low-overhead random element access.



```
typedef Matrix<Matrix<Matrix<double> >  
> MyMatrixType;  
typedef Matrix<Matrix<Matrix<long double> >  
> MyAccurateMatrixType;
```

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# HML Features

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Parallelization

- Easy to code and maintain.
- Block size determined by the architecture performance, not chemistry.
- Low overhead random element access.
- Blocked algorithms easy to express:
  - ① Matrix multiplication, also by transposed matrices.
  - ② Use of matrix symmetry.
  - ③ INverse CHolesky factorisation (INCH).

# Block Size Tradeoff

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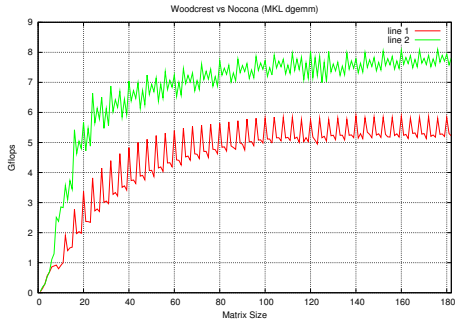
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- Smaller block size → more opportunity for screening.
- Larger block size → better block-block multiplication performance.





# Example Implementation of $C := \text{beta} * C + A * B$

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```
static void multiply(const Matrix<Telement>& A,
                   const Matrix<Telement>& B,
                   Matrix<Telement>& C, double beta) {
    for (int colC = 0; colC < C.ncols; colC++)
        for (int rowC = 0; rowC < C.nrows; rowC++) {
            Telement::multiply(A(rowC, 0), B(0, colC),
                               C(rowC, colC), beta);
            for (int colA = 1; colA < A.ncols; colA++)
                Telement::multiply(A(rowC, colA), B(colA, colC),
                                    C(rowC, colC), 1);
        }
}
```

- Lowest level (block) multiplication expressed in terms of BLAS calls.
- Template expansion will generate (*instantiate*) code for all the remaining hierarchy levels.

# Intel MKL vs HML Benchmark

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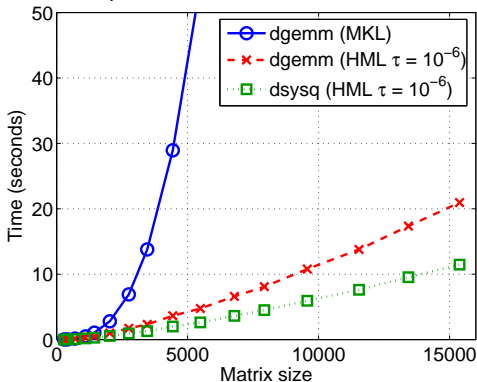
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Parallelization

- HML design allows for easy implementation of symmetric matrix multiplication (sysq:  $S = \alpha T^2 + \beta S$ ) as needed by TC2:
- sysq twice faster than general sparse multiplications.

Matrix multiplication benchmark: water clusters/3-21G



# OpenMP Parallelization

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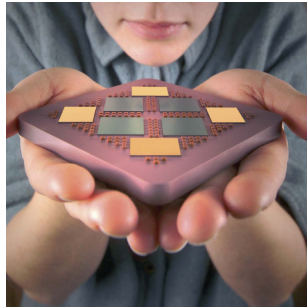
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Parallelization

- Parallel programs necessary to efficiently use modern multi-core hardware.
- OpenMP less *invasive* and easier to load-balance.
- Problems with scaling and... compiler support.
- Poor compiler support! GNU gcc is the only reliable, OpenMP-enabled compiler known to us so far.



# Details of OpenMP Parallelization

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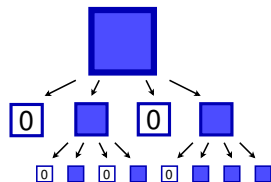
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Parallelization

- Pick a level in the hierarchy, run a parallel loop with dynamic scheduling over it.
- Approach trivial to implement.
- Higher levels: coarse load distribution.
- Lower levels: thread startup overhead.



# Exceptions and OpenMP

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Parallelization

- OpenMP and C++ exceptions *do* interact.
- Threads must catch any exceptions that are generated. The behavior is undefined otherwise.
- We do the *right* thing (in case you ask).

```
#pragma omp parallel for
for (int i = 0; i < MAX; i++) {
    try {
        // Heavy lifting here
    } catch (...) { /* Handle it nicely. */ }
}
```

# Compiler Problems

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**GNU C++** OpenMP support since 4.1(?). No problems found. Sequential performance lower than its competitors.

**Portland C++** fairly warns that it cannot handle exceptions and OpenMP at the same time. A honest warning but. . .

**Intel C++** 3 versions tried. *All* of them had bugs either in sequential code or in OpenMP parallelization.

**8.1** fails to generate correct sequential code; miscompiles OpenMP code as well.

**9.1** works sequentially; compiler crashes with executed with `-openmp` flag.

**10.0** fails to generate correct sequential code. Support tickets with Intel are open.

# OpenMP Speedup

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Parallelization

- Timings taken on 1.5GHz Itanium2, 4 CPU (luc2, PDC), 4 threads.
- Glycine-Alanine chain with 1600+ atoms, HF method. GNU C++.

Operation	CPU time [s]	Wall time [s]	Speedup
FDS-SDF	133.54	53	2.66
Purification	947.59	454	2.13

- Acceptable multiplication load balancing (3.5/4.0) but serial data management has negative impact on scalability.
- Additionally, purification involves serial error estimation routines.

# Summary & Outlook

Hierarchic  
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Parallelization

- SSSA – for strict error control.
- HML – flexible sparse matrix representation.
- A number of algorithms (arbitrary  $M \times M$  multiplications, inverse Cholesky factorisation) already implemented.
- OpenMP parallelization.

## OUTLOOK

- Analyse the sparsity in the QM methods beyond the algorithms relevant for SCF: Linear response for calculation of molecular properties.



# Small Submatrix Selection Algorithm (SSSA)

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Given a matrix norm  $\|\cdot\|$  and an error limit  $\varepsilon$  we want to find a sparse approximation  $\tilde{A}$  of  $A$  so that  $\|A - \tilde{A}\| < \varepsilon$ .

SSSA:

- 1 Compute the Frobenius norm of each submatrix.
- 2 Sort the values in descending order.
- 3 Remove submatrices from the end as long as the error is within desired accuracy.

⇒ Error very close to the requested value in the Frobenius norm.