

Locality-aware parallel block-sparse matrix-matrix multiplication using the Chunks and Tasks programming model



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ABSTRACT

We present a method for parallel block-sparse matrix-matrix multiplication on distributed memory clusters. By using a quadtree matrix representation, data locality is exploited without prior information about the matrix sparsity pattern. A distributed quadtree matrix representation is straightforward to implement due to our recent development of the Chunks and Tasks programming model [Parallel Comput. 40, 328 (2014)]. The quadtree representation combined with the Chunks and Tasks model leads to favorable weak and strong scaling of the communication cost with the number of processes, as shown both theoretically and in numerical experiments.

Matrices are represented by sparse quadrees of chunk objects. The leaves in the hierarchy are block-sparse submatrices. Sparsity is dynamically detected by the matrix library and may occur at any level in the hierarchy and/or within the submatrix leaves. In case graphics processing units (GPUs) are available, both CPUs and GPUs are used for leaf-level multiplication work, thus making use of the full computing capacity of each node.

The performance is evaluated for matrices with different sparsity structures, including examples from electronic structure calculations. Compared to methods that do not exploit data locality, our locality-aware approach reduces communication significantly, achieving essentially constant communication per node in weak scaling tests.

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1. Introduction

Sparse matrix-matrix multiplication, sometimes referred to as SpGEMM, is a key operation in large-scale electronic structure calculations based on for example Hartree–Fock or Kohn–Sham density functional theory [1]. Sparse matrix-matrix multiplication is used in particular in polynomial expansion [2] and minimization methods [3] to compute the density matrix. Such methods are used in a number of electronic structure codes such as CONQUEST [4], CP2K [5], ERGO [6], FREEON [7], HONPAS [8], ONETEP [9], and LATTE [10] to achieve a computational cost that increases only linearly with system size. The matrix sparsity varies from tens to thousands of nonzeros per row depending on the underlying model and the basis set used. It is often beneficial to use a block-sparse data structure. The optimal block size depends on the model and on the order of the matrix rows and columns. The present work is mainly motivated by Hartree–Fock and Kohn–Sham density

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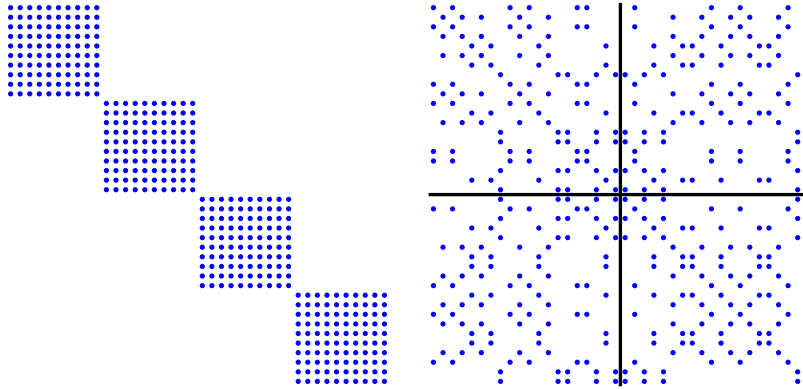


Fig. 1. A trivial example of possible effects of random permutation to load balance static work and data distribution. Left: Example matrix for which we want to compute the square on 4 identical compute nodes. The best distribution of data and work is obvious and leads to perfect load balance and no communication of matrix elements. Right: Random permutation of the matrix columns and rows and a two-dimensional data decomposition indicated by solid black lines. Although the workload may still be roughly load balanced, communication of matrix elements is now needed.

functional theory calculations using Gaussian basis sets in which the matrices have up to thousands of nonzero elements per row and a priori unknown sparsity patterns [6,11]. This work is also relevant for the general parallel SpGEMM problem as no application specific information such as atomic positions is built into the presented method.

Algorithms based on dense matrix-matrix multiplication are generally considered attractive because of the existence of efficient linear algebra libraries, e.g. [12,13], and parallelization through e.g. Cannon's algorithm or SUMMA [14]. Parallel sparse and block-sparse matrix-matrix multiplication has received less attention and represents a greater challenge, particularly when the nonzero pattern is not known in advance. Nevertheless, several parallel sparse matrix-matrix multiplication methods have been presented. Several methods assume some a priori knowledge about the input matrix sparsity structure and use that knowledge to improve performance [15–18]. Other methods require beforehand knowledge of the computational pattern or the sparsity structure of the output matrix, requiring a preparatory symbolic multiplication step before the actual parallel computation can start [19,20]. Here, we will focus on the general case where no a priori knowledge about the structure is assumed, and no symbolic multiplication step is needed. Recent methods for this general case are first employing a random permutation of the rows and columns of the matrix to destroy any structure in the sparsity pattern and decrease data locality [21,22]. The goal of this maneuver is to obtain about the same density of nonzero elements everywhere in the matrix. Then, a static distribution of work and data is used in the same way as for dense matrices, but with the local block-block multiplies replaced by sparse products. This random permutation approach prevents load imbalance, but the obvious drawback is that the possibility to exploit the nonzero structure to reduce communication or make efficient use of the memory hierarchy is spoiled, see Fig. 1 for a trivial yet illustrative example. On the other hand, such exploitation is difficult to achieve since it requires that the mapping of data and work to physical resources is performed dynamically during the calculation [23]. We believe that the difficulties are mainly associated with the programming model used to tackle the problem. While conventional programming models like message passing protocols work well for static distribution of work and data, they are inconvenient if you want to distribute data and work dynamically. The programmer has to make decisions about where data should be located, where every piece of work should be executed, and see to it that data is communicated as needed.

Recently, we proposed a new programming model named Chunks and Tasks, designed to work well for algorithms with dynamic work and data [24]. We describe and analyze in this article an approach to parallel block-sparse matrix-matrix multiplication based on a hierarchical quaternary tree (quadtrees) representation implemented using the Chunks and Tasks programming model. The method is locality-aware in the sense that it is able to exploit a priori unknown structure in the sparsity pattern to reduce communication and thereby improve performance. The library used for performance evaluation here is a further developed version of the code that was briefly described and used for test calculations in [24].

This article is organized as follows: in Section 2, we briefly discuss the Chunks and Tasks programming model. Our new matrix library based on Chunks and Tasks is presented in Section 3. In the present work, the Chunks and Tasks matrix library is used together with a block-sparse leaf matrix type, described in Section 4. An analysis of the computational costs due to the quadtree representation is given in Section 5, followed by results of test calculations in Section 6 and concluding remarks in Section 7.

2. Programming model

Our block-sparse matrix library has been implemented using the Chunks and Tasks programming model [24]. In Chunks and Tasks the programmer writes her program in terms of small pieces of data and work, chunks and tasks, respectively.

The programmer is responsible for dividing work and data into smaller pieces but not for the mapping of work and data onto physical resources. The programmer need not worry about message passing, all communication is handled by the Chunks and Tasks library. The programmer does neither have to worry about race conditions nor non-deterministic behavior. The computation is driven by the registration of tasks, similarly to other task-based models. Recursive nesting of tasks is allowed, i.e. during task execution new tasks can be registered as in for example Cilk [25], Scioto [26], SuperGlue [27], and XKaapi [28]. This is important for scalability of dynamic algorithms, since otherwise only a single process can generate new tasks, or multiple processes generate predetermined (static) task graphs.

A key feature of the Chunks and Tasks model is that abstractions are not only provided for work but also for data. The Chunks and Tasks library takes care of the distribution of both work and data. The user creates data objects called chunks. The transfer of responsibility of such a chunk object to the runtime library is referred to as registration of a chunk; in return, the user gets an identifier that can be used to specify dependencies later on. After the point of registration the chunk object is read-only. This is in a way similar to e.g. Linda [29] and Concurrent Collections [30] that also have a “space” to which you can add a piece of data and later retrieve it, possibly on another process. A key difference is that in Linda and Concurrent Collections the identifier is chosen by the application programmer whereas in Chunks and Tasks, the identifier is chosen by the runtime library. On one hand, being able to choose identifiers makes it possible for a process or task to ask for data without any prior communication or interaction whatsoever with the process or task that registered the data. On the other hand, the fact that the runtime library does not control the identifiers means that inconsistencies can be introduced (supposedly unintentionally) where for example several different pieces of data with the same identifier exist, e.g. on distant nodes in a cluster. Perhaps of even greater importance is that such a model makes it difficult for the runtime library to make data available efficiently. Any process may ask for any piece of data at any time possibly without any information being available locally about the location of the piece being asked for. This stands in contrast to Chunks and Tasks where the library for example can store information about the location in the chunk identifier. In this way, Chunks and Tasks, by imposing appropriate restrictions, makes life easier both for the application programmer and the runtime library developer.

2.1. Library implementations

A Chunks and Tasks program can be compiled, linked and executed with any Chunks and Tasks runtime library implementation. We will in our performance evaluation use the publicly available Chunks and Tasks library CHT-MPI [24,31], which is written in C++ and uses the Message Passing Interface (MPI) for communication between computational nodes, enabling Chunks and Tasks programs to run on distributed-memory clusters.

The CHT-MPI implementation uses work stealing to distribute tasks, see e.g. [32], meaning that there is no central “master” node responsible for all scheduling. Each worker process is responsible for its own set of tasks, and steals work from some other (randomly selected) worker when it has no work left. For recursive algorithms operating on hierarchical data structures, some tasks recursively generate new tasks leading to a tree of tasks, and each worker process effectively executes its own local part of that tree. Work stealing always occurs as high up as possible in the local task tree of the victim process.

Besides distributing tasks, the runtime library must also make sure that the necessary input data is available for each task. Thanks to the possibility of storing the MPI rank of the owner process in the chunk identifier, this becomes straightforward: when a particular chunk is needed as input to a task, the library implementation simply inspects the chunk identifier to find out from which worker the data should be fetched. In this way data can be made available efficiently, without need for any central authority storing information about the location of all chunks.

Each chunk object is by default owned by the worker process that created that chunk. This has the advantage that no communication is needed to create a chunk, and temporary chunks used within a local part of the task tree can often be reused directly without need for any communication, since each worker processes its own local part of the task tree. CHT-MPI also implements a chunk cache for each worker process, meaning that if the same chunk is needed multiple times it only needs to be fetched the first time. Thus, for Chunks and Tasks programs corresponding to recursive algorithms operating on hierarchies of chunks, data re-use happens automatically. Note that the data distribution is determined dynamically and follows from the work stealing distribution of tasks. The distribution of chunks among worker processes will therefore in general be different for different runs of the same program.

For the reasons outlined above, CHT-MPI can be used to achieve scalable parallelization for Chunks and Tasks programs; both work and data is distributed dynamically without need for any “master” node that all workers must communicate with. From an application programmer's point of view, what is needed to make use of these features is to express the program using hierarchical representations and recursive algorithms. As will be seen in the following section, a quadtree-based representation is a natural way to achieve this for matrix operations. See [24,31] for more information about CHT-MPI.

3. Quadtree representation of matrices in the Chunks and Tasks model

Hierarchical data structures based on a two-dimensional block decomposition of the matrix at each level in a hierarchy have both been used to block for the memory hierarchy in dense matrix computations [33–35] and to avoid operations on zero elements (or entire submatrices that are zero) in sparse matrix computations [36]. Quadtree representations have also been advocated for simplicity and expressiveness in particular leading to ease of programming for multiprocessing (shared

```

1: input:  $A, B$ 
2: if  $A$  not NIL and  $B$  not NIL then
3:   if lowest level then
4:      $X = \text{leafMatrixMultiply}(A, B)$ 
5:      $C = \text{registerChunk}(X)$ 
6:   else
7:     for  $m = 1, 2$  do
8:       for  $n = 1, 2$  do
9:          $Y_1 = \text{registerTask}(\text{multiply}, A_{m1}, B_{1n})$ 
10:         $Y_2 = \text{registerTask}(\text{multiply}, A_{m2}, B_{2n})$ 
11:         $C_{mn} = \text{registerTask}(\text{add}, Y_1, Y_2)$ 
12:      end for
13:    end for
14:     $C = \text{registerTask}(\text{createFromIds}, C_{11}, C_{12}, C_{21}, C_{22})$ 
15:  end if
16: else
17:    $C = \text{NIL}$ 
18: end if
19: output:  $C$ 

```

} execute

→ fallback

Algorithm 1. Pseudo-code for quadtree based matrix-matrix multiplication using the Chunks and Tasks programming model.

memory) environments [33,34,36–39] and straightforward exploitation of symmetry [39]. As will be shown in Section 5, the quadtree representation is in principle also appropriate for distributed representation of sparse matrices on computer clusters. The caveat is that a distributed sparse quadtree representation is difficult to implement in conventional programming models, especially if a priori unknown sparsity patterns are to be handled efficiently. In this section, we describe how such a sparse matrix quadtree representation can be straightforwardly implemented in the Chunks and Tasks programming model [24].

In our Chunks and Tasks matrix library, matrices are represented by sparse quadtrees of chunks. At the lowest level in the hierarchy, different leaf matrix representations, for example dense or sparse, may be used. In this work we will focus on regular matrix-matrix multiplication on the form $C = AB$ and the symmetric matrix square operation $C = A^2$, where A and therefore also C is symmetric and only the upper triangles of A and C are stored. The sparse symmetric matrix square is a key operation and a major computational challenge in linear scaling electronic structure calculations. In all task type implementations, sparsity is dynamically exploited at all levels in the hierarchy by skipping operations on zero submatrices, which are represented by NIL chunk identifiers. Regular matrix-matrix multiplication without transpose $C = AB$ and matrix addition $C = A + B$ are illustrated as pseudo-code in Algorithms 1 and 2, respectively. For all task types, at the lowest level in the hierarchy, the corresponding functionality of the leaf matrix library is used, while at higher levels a straightforward implementation for the two by two case is used, see e.g. lines 7–14 in Algorithm 1. Checking for NIL chunk identifiers corresponds to the if statement on line 2 in each of Algorithms 1 and 2. In practice, the Chunks and Tasks C++ interface requires a regular execute function used when all input chunks are available, and a fallback execute used when some of the input chunk objects cannot be constructed due to NIL chunk identifiers. This is convenient from a programmer's point of view since the if statement on line 2 is checked by the runtime library and the appropriate function, regular or fallback execute, is called automatically. This also allows for compile-time type checking and programming errors such as attempts to access nonexistent chunks are not possible. We list and describe below all chunk and task types that are needed for the $C = AB$ and $C = A^2$ operations.

3.1. Chunk types for quadtree representation

- **Matrix:** A basic matrix chunk type is used to represent nonzero submatrices in the quadtree representation. At each but the lowest level in the hierarchy, the matrix is divided into four submatrices represented by their chunk identifiers. At the lowest level, a leaf matrix type is used for matrix representation. Storage and addressing of zero submatrices is avoided at all levels in the hierarchy. Zero submatrices are represented by NIL chunk identifiers. Note that a NIL chunk identifier can appear at any level in the hierarchy. The matrix dimension is also stored along with the maximum allowed dimension for leaf matrices. This basic chunk type is the natural Chunks and Tasks implementation of a quadtree matrix representation as defined by Wise and Franco [40]. When setting up the quadtree structure the matrix is split so that a predetermined uniform blocksize at each level in the hierarchy is achieved as far as possible. This ensures that when two matrices A and B are combined in e.g. a multiplication or addition operation, the submatrix dimensions of A and

```

1: input:  $A, B$ 
2: if  $A$  not NIL and  $B$  not NIL then
3:   if lowest level then
4:      $X = \text{leafMatrixAdd}(A, B)$ 
5:      $C = \text{registerChunk}(X)$ 
6:   else
7:     for  $m = 1, 2$  do
8:       for  $n = 1, 2$  do
9:          $C_{mn} = \text{registerTask}(\text{add}, A_{mn}, B_{mn})$ 
10:      end for
11:    end for
12:     $C = \text{registerTask}(\text{createFromIds}, C_{11}, C_{12}, C_{21}, C_{22})$ 
13:  end if
14: else
15:   if  $A$  not NIL then
16:      $C = A$ 
17:   else if  $B$  not NIL then
18:      $C = B$ 
19:   else
20:      $C = \text{NIL}$ 
21:   end if
22: end if
23: output:  $C$ 

```

Algorithm 2. Pseudo-code for quadtree based matrix addition using the Chunks and Tasks programming model.

B will match. The blocksize used at leaf level should be chosen to achieve an appropriate granularity of chunks and tasks. This is a trade-off: the leaf level chunks/tasks should be small enough to allow sufficient parallelism for the given computational resource. At the same time, each leaf level chunk/task should contain enough data/work to make the administration overhead of the runtime library negligible. The matrix dimensions at higher levels in the hierarchy are directly determined by the lowest level blocksize, by multiplying by a factor of 2 for each level. A submatrix in the quadtree represented by a matrix chunk does not contain any global information such as the global matrix dimension or its location in the entire matrix, i.e. row and column offsets.

- *Matrix parameters:* A chunk type for matrix parameters is used to convey information needed in the construction of matrix chunks. The chunk includes information about the matrix dimension and the leaf matrix dimension, and whenever needed information about the location of the matrix (its rowwise and columnwise offsets from the upper left corner) in the global matrix. It is also used to store information needed by the leaf matrix type.

3.2. Task types for regular matrix-matrix multiplication

- $C = AB$, $C = A^T B$, $C = AB^T$, and $C = A^T B^T$: The task types for regular and transposed matrix-matrix multiplication takes two matrix chunks, described above, and returns the product. If the input matrices are both at the lowest level in the hierarchy, the corresponding leaf matrix multiplication is invoked. Otherwise matrix multiplication and matrix addition tasks for child submatrix multiplication and addition are registered for execution. The results are collected into the result matrix with a task for creation of a matrix chunk from four submatrix chunk identifiers. See the pseudo-code in [Algorithm 1](#) for the $C = AB$ case.
- $C = A + B$: The task type for matrix addition takes two matrix chunks and returns their sum. If the input matrices are at the lowest level, the addition in the leaf matrix library is performed. Otherwise, tasks for child submatrix addition are registered for execution, and the results are collected with a task for creation of a matrix chunk from child submatrix identifiers. See the pseudo-code in [Algorithm 2](#).
- *Creation from submatrix identifiers:* A task type for creation of a matrix chunk from four submatrix chunk identifiers is needed since chunks are read-only after the point of registration. Since the submatrix chunk identifiers are included in the matrix chunk, it is not possible to construct the matrix chunk before the construction of the submatrices. Therefore, this task type is used whenever a matrix that depends on the results of other tasks needs to be constructed. This task type also takes a chunk of matrix parameters as input.

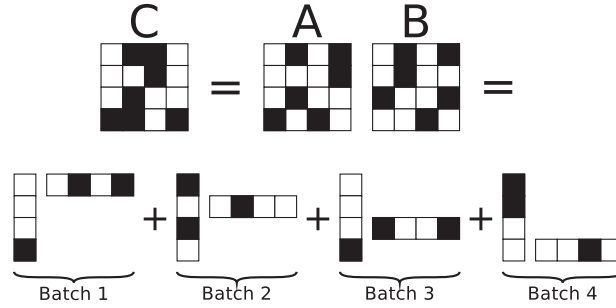


Fig. 2. Illustration of a sparse matrix product as a sum of sparse outer products. All multiplies within a batch are independent making it possible to use the batched gemm API in cuBLAS.

3.3. Additional task types for symmetric matrix square

- $C = A^2$ where A is symmetric: A symmetric matrix square task squares a symmetric matrix in upper triangular storage. At the lowest level the corresponding leaf matrix symmetric matrix square operation is executed. At higher levels, the symmetric matrix square task registers symmetric rank-k, symmetric square, symmetric multiply, and matrix addition tasks to compute the submatrices in the product matrix. The results are collected into the result matrix chunk with a task for creation of a matrix chunk from submatrix identifiers. In general, a symmetric matrix square task directly or indirectly makes use of all but the $C = A^T B^T$ task type in this section. Therefore, a benchmark of the symmetric matrix square operation covers nearly all task types presented here.
- $C = AB$, where A or B is symmetric: Symmetric matrix multiply is the task type for multiplication of two matrices where either the first or the second multiplicand is a symmetric matrix in upper triangular storage. At the lowest level the corresponding multiplication operation for symmetric matrix multiply is executed. At higher levels, tasks for regular matrix multiplication, symmetric matrix multiply, and matrix addition are registered and the results are collected into the result matrix chunk with a task for creation from submatrix identifiers.
- $C = AA^T$ and $C = A^T A$: With this task, the so-called symmetric rank-k operation is performed; a symmetric matrix in upper triangular storage is constructed from the product of a general matrix and its transpose. At the lowest level the symmetric rank-k operation of the leaf matrix library is used. At higher levels, symmetric rank-k, matrix multiplication, and matrix addition tasks are registered and a task for creation from submatrix identifiers is used to collect the results into the result matrix chunk, as for the other task types.

4. Leaf matrix types

As discussed above, different leaf matrix representations may be used at the lowest level in the quadtree. A leaf matrix type used together with our chunk quadtree representation has to implement some basic functionality such as serialization routines. The leaf matrix type also has to implement functionality needed by task types used together with the leaf matrix type. When implementing a leaf matrix type, one can assume that the matrix data fits in the memory of a single compute node. All functionality in the class has to be thread-safe, since the Chunks and Tasks library must be able to execute several leaf tasks and call the serialization routines simultaneously.

4.1. Block-sparse leaf matrix type

In the present work we are using a block-sparse leaf matrix type. The block-sparse matrix class uses a uniform block-size configurable via the matrix parameters chunk type as discussed in Section 3.1. Submatrices are kept in a simple two-dimensional array where only non-zero submatrices are allocated. The leaf matrix library makes use of the Basic Linear Algebra Subprograms (BLAS) [41] for submatrix-submatrix multiplications on CPUs and the NVIDIA CUDA Basic Linear Algebra Subroutines (cuBLAS) [42] for submatrix-submatrix multiplications on graphics processing units (GPUs). An advantage of using the BLAS and cuBLAS library interfaces is that we can take advantage of optimized BLAS and cuBLAS library implementations. We have for example observed substantial performance improvements of the cuBLAS library when going from Cuda 5.0 to Cuda 6.5 (see the caption of Table 2).

Our block-sparse leaf matrix type targets problems where a block size around 16–64 is appropriate. The regular gemm operation in cuBLAS is inefficient for such small matrix dimensions. Therefore, we are instead making use of the batched gemm API in cuBLAS. The routine executes a batch of small matrix-matrix multiplications. The operations in a batch should be independent in the sense that none of the multiplications are allowed to write to the same product matrix. The block-sparse multiplication can be expressed as a sum of outer products, see Fig. 2. Each outer product is a batch of small matrix-matrix multiplications and all multiplies within a batch are independent.

Algorithm 3 Algorithm for load balanced processing of batch lists.

```

1: Get list of batches (CPU)
2: while not done do
3:   if free GPU slot then
4:     Process remaining batches on GPU
5:   else
6:     Process one batch on CPU
7:   end if
8: end while

```

We would like to make use of both CPUs and GPUs, if any. If there is a GPU available, the multiply will be processed by the GPU. Otherwise the multiply will be processed by a CPU core. Since there will be other threads that execute similar tasks, a GPU may become available during the calculation. Then, it is generally good if the remaining work can be offloaded to the GPU. To achieve this we are using [Algorithm 3](#). Using this algorithm, load balancing between the CPUs and GPUs is achieved when several threads executing leaf matrix multiplies are running on the host. An alternative could be to use a Chunks and Tasks library that let idle host threads reexecute already running tasks whenever there is no more work, as in [\[43\]](#). The application programmer would then not have to worry about feeding smaller portions to the CPU, and such an approach could also help in case of various failures. On the other hand, some tasks would be executed more than once.

4.2. Efficient utilization of both GPUs and CPUs

To make efficient use of both GPUs and CPUs on each node, data transfers to/from GPUs should as far as possible be overlapped with computation and the load should be shared between the GPUs and the CPU cores. In addition, in case the amount of work in each task varies, it is preferable to run small tasks on the CPU cores and let the GPUs handle the computationally heavier tasks.

Overlap of data transfers and computation is achieved by keeping two slots for each device (GPU), allowing one thread to do computations while another thread is transferring data. In order to have a task ready for execution on each device, a bounded queue is used, with a queue length equal to the number of devices. To allow computationally heavy tasks to run primarily on the GPUs, the queue is prioritized according to a measure of the expected amount of work needed to execute each task. For tasks that do not fit in the queue, the work is performed on a CPU core until a GPU becomes available, see [Algorithm 3](#). This gives load balancing between GPUs and CPUs provided that the number of threads is large enough; for small numbers of threads only GPUs are used.

Since the preparatory work needed to get the list of batches for each task is performed on a CPU core before checking for a place in the queue, computation on the GPU is overlapped not only with data transfers to/from the device memory but also with the preparatory work.

Note that the Chunks and Tasks C++ interface presented in [\[24\]](#) does not include support for data transfers between the host and GPU memory nor does it assist in scheduling of tasks on GPUs, as does for example StarPU [\[44\]](#). The Chunks and Tasks library is unaware of any devices that may be installed on the compute nodes. The code for deciding when to offload work to the GPU, as well as data transfers to/from GPU memory, is part of the task execution. This means that data that is used by several tasks running on the same GPU will be transferred to the device memory once for each task. However, as will be seen in [Section 6.1](#), the design described above achieves both hiding of data transfer costs and sharing of the workload between GPUs and CPUs.

5. Computational costs associated with the quadtree representation

In this section, we first consider the number of tasks for different sparsity patterns, and then use those results to get theoretical estimates of computation and communication costs.

The total number of tasks includes both addition and multiplication tasks. However, for the purpose of studying the scaling behavior of the total number of tasks it is sufficient to consider only the number of multiplication tasks since this is always larger than the number of addition tasks. This can be understood in the following way. The input submatrices to any addition task are temporaries that come from previous addition or multiplication tasks. The input submatrices to any multiplication task come from the original input matrices (A and B). The output submatrix resulting from any multiplication or addition task can appear as input to at most one addition task. Therefore, the tasks generating the trace of temporaries that precedes any addition task form a binary tree in which every node has 0 or 2 children with multiplication tasks as leaves and addition tasks as non-leaves. Such a tree has more leaves than non-leaves. Thus, the total number of addition tasks is strictly bounded by the total number of multiplication tasks.

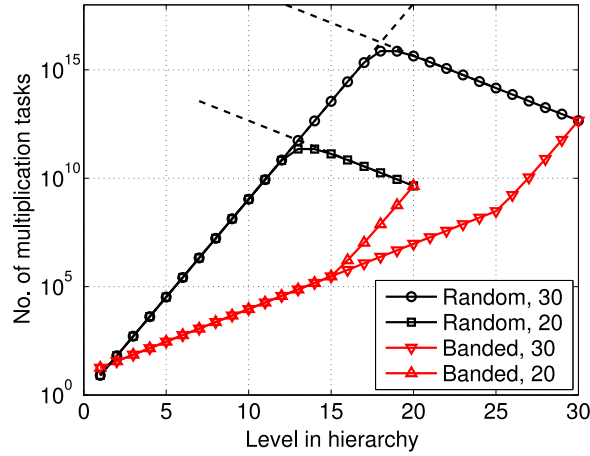


Fig. 3. Number of matrix-matrix multiplication tasks at different levels in the hierarchy for random and banded matrices with matrix dimension $N = 2^L$ with $L = 20$ and $L = 30$. The target number of nonzero elements per row is 65, corresponding to $\delta = 65/N$ in the random case and $k = 5$ in the banded case. The dashed lines indicate the upper bounds given by (2) and (3).

5.1. Total number of tasks

We will here consider the total number of matrix-matrix multiplication tasks for different sparsity patterns. Tasks executed at higher levels can be seen as administration work required to determine which low-level tasks are needed. A key issue is how much such extra administration work that is generated due to the quadtree representation.

In this section, we consider a quadtree representation with blocksize 1 at the lowest level. In practical calculations, a larger blocksize will typically be used for performance reasons; we use blocksize 1 here in order to more clearly see the effects of the quadtree structure. The use of a larger blocksize will correspond to merging several of the lowest levels, leaving it to the leaf matrix library to handle any sparsity there.

We first consider a case with little data locality, a random sparsity pattern where the nonzero matrix elements are uniformly randomly distributed. The probability δ to find a nonzero element at a given position is the same everywhere in the matrix and uncorrelated to the position of other nonzero matrix elements.

Let the levels in the hierarchy be labeled such that level $l = 0$ is the highest level (the root of the tree) and level $l = L$ is the lowest (leaf) level. Let N be the matrix dimension and $N_l = 2^l$ such that N_l^2 is the total number of submatrices at level l and $N_L = N$. If we denote the probability of a submatrix at level l being nonzero as δ_l , the expected number of matrix-matrix multiplication tasks at level l is

$$C_l^{\text{random}} = N_l^3 \delta_l^2 = 2^{3l} (1 - (1 - \delta)^{n_l})^2 \quad (1)$$

where $n_l = 2^{2(L-l)}$ is the total number of elements (including both zeros and nonzeros) in each submatrix at level l . The relationship (1) is illustrated in Fig. 3.

Since $\delta_l \leq 1$, it follows that

$$C_l^{\text{random}} \leq N_l^3 = 8^l \quad \text{for all } l. \quad (2)$$

Furthermore, the probability δ_l of a submatrix being nonzero satisfies the relation $\delta_l = 1 - (1 - \delta_{l+1})^4 = 4\delta_{l+1} - 6\delta_{l+1}^2 + 4\delta_{l+1}^3 - \delta_{l+1}^4 \leq 4\delta_{l+1}$. Therefore, we also have that

$$C_l^{\text{random}} \leq N_l^3 (4^{L-l} \delta)^2 = \frac{16^L \delta^2}{2^l} \quad \text{for all } l. \quad (3)$$

Although both inequalities (2) and (3) are valid for all l , (2) is tight for low levels while (3) is tight for high levels, as seen in Fig. 3.

Let $x = \log_2(N) + \frac{\log_2(\delta)}{2}$ be the point where the two bounds above intersect, given by solving $4^{L-x} \delta = 1$. Then, $8^x = (\delta N^2)^{\frac{3}{2}}$, $8^{\lfloor x \rfloor} \leq 8^x$, and $\frac{16^L \delta^2}{2^{\lfloor x \rfloor + 1}} < \frac{16^L \delta^2}{2^x} = 8^x$. Therefore, assuming $\delta \geq 1/N^2 \implies x \geq 0$ the expected total number of tasks

$$\sum_{l=0}^L C_l^{\text{random}} \leq \sum_{l=0}^{\lfloor x \rfloor} 8^l + \sum_{l=\lfloor x \rfloor + 1}^L \frac{16^L \delta^2}{2^l} \quad (4)$$

$$= 8^{\lfloor x \rfloor} \sum_{l=0}^{\lfloor x \rfloor} \frac{1}{8^l} + \frac{16^L \delta^2}{2^{\lfloor x \rfloor + 1}} \sum_{l=0}^{L-\lfloor x \rfloor - 1} \frac{1}{2^l} \quad (5)$$

$$< 8^x \left(\frac{1}{1-1/8} + \frac{1}{1-1/2} \right) \quad (6)$$

$$= (3^{1/7})(\delta N^2)^{3/2}. \quad (7)$$

We note that even though the number of tasks at leaf level is $\mathcal{O}(N^3\delta^2)$, the total number of tasks is $\mathcal{O}(N^3\delta^{3/2})$ due to excessive administration work at higher levels.

As a simple case with data locality, we consider banded matrices with bandwidth $b = 2d + 1$ where for simplicity we assume that $d = 2^k$ for some $k \geq 0$. In this case, the number of matrix-matrix multiplication tasks at level l is bounded by

$$C_l^{\text{banded}} < N_l b_l^2 = 2^l (2d_l + 1)^2 \quad (8)$$

where

$$d_l = \begin{cases} 1 & \text{for } l < L - k, \\ 2^{l-(L-k)} & \text{for } l \geq L - k. \end{cases} \quad (9)$$

As seen in Fig. 3, the case with data locality gives a very different behavior of the number of tasks on each level; most of the work is concentrated at the lowest levels in the hierarchy. The total number of tasks is bounded by

$$\sum_{l=0}^L C_l^{\text{banded}} < \sum_{l=0}^{L-k-1} 2^l 3^2 + \sum_{l=L-k}^L 2^l (2 \cdot 2^{l-(L-k)} + 1)^2 \quad (10)$$

$$< \left(4^{4/7} d^2 + 5^{1/3} d + 2 + \frac{9}{d} \right) N. \quad (11)$$

We note that the total number of tasks is proportional to the number of tasks at the lowest level, i.e. no excessive administration work is going on at higher levels.

As an example of sparsity structures appearing in physical applications, we consider overlap matrices for systems of evenly distributed particles in $D \geq 1$ spatial dimensions with one spherically symmetric basis function per particle, ordered using a recursive divide-space procedure. We consider finite systems (no periodicity). A matrix element A_{ij} is nonzero if the distance between particles i and j is smaller than some radius R . Note that this is a kind of sparsity structure found in many applications in physics and chemistry, where each matrix element is often related to a pair of particles or other objects in a physical system; sparsity arises from the fact that only matrix elements that correspond to objects that are sufficiently close in space are nonzero. In D spatial dimensions with the chosen ordering of basis functions, the N_l blocks at level l can be seen as corresponding to a set of N_l spatial boxes such that all basis functions (or particles) in a given block are contained in the corresponding box. The number of multiplication tasks at a given level can be estimated by

$$C_l^{\text{overlap}} < N_l M_l^2 = 2^l M_l^2 \quad (12)$$

where M_l is the number of spatial boxes that can be reached by a sphere of radius R . For high levels where the width of spatial boxes is larger than R , M_l is determined by the number of neighboring boxes, $M_l = 3^D$, independently of l . For lower levels M_l is proportional to the volume of a D -dimensional sphere of radius $\frac{R}{h_l}$ where the width of boxes $h_l \propto 2^{\frac{L-l}{D}}$, giving $M_l \propto R^D 2^{l-L}$. Therefore, analogously to the banded matrix case, for high levels $C_l^{\text{overlap}} \propto 3^{2D} 2^l$ and for lower levels $C_l^{\text{overlap}} \propto R^{2D} 2^{3l-2L}$. We note also that $C_l^{\text{overlap}} \geq 2C_{l-1}^{\text{overlap}}$ for all $l \geq 1$. Therefore, as for the banded matrix case, the total number of tasks is proportional to the number of tasks at the lowest level. Numerical experiments for $D = 1, 2, 3$ are shown in the left panel of Fig. 4. The test matrices were created using the Ergo program [6] to compute overlap matrices for artificially generated 1d, 2d, and 3d molecules with one basis function per atom from the standard Gaussian basis set STO-3G, applying the default recursive divide-space procedure to order the atoms. The molecules were generated by placing hydrogen atoms on a D -dimensional grid with separation 2 Å and a uniform random displacement of up to ± 1 Å in each coordinate direction. The matrix size was $2^{16} = 65536$ for $D = 1, 2$ and $40^3 = 64000$ for $D = 3$. Blocksize 1 was used for the 1d case while the 2d and 3d cases used blocksize 2 and 4, reducing the number of hierarchy levels for those cases by 1 and 2, respectively.

As an example where different degrees of data locality can be easily investigated we consider multiplication of graph matrices constructed using the R-MAT model [45]. We choose the R-MAT parameters such that $b = c = d = \frac{1-a}{3}$ and perform tests for different values of a in the range $0.25 \leq a < 1$. Setting $a = 0.25$ essentially corresponds to the random case described above, while increasing a corresponds to increasing the data locality. We use matrix dimension $2^{13} = 8192$ and a number of graph edges corresponding to 5 nonzero elements per row, although for large a values matrices become more sparse due to multiple graph edges between the same nodes. The right panel of Fig. 4 shows the number of matrix-matrix multiplication tasks for a set of different a values. We note that most of the work is pushed towards lower levels as the degree of data locality increases.

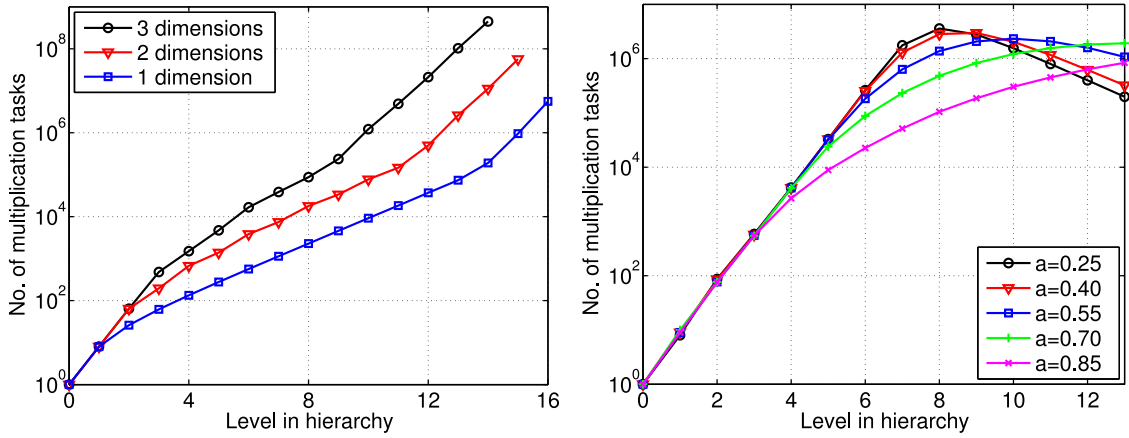


Fig. 4. Number of matrix-matrix multiplication tasks for matrices with different sparsity structures. Left: overlap matrices for artificially constructed 1d, 2d, and 3d molecules. Right: R-MAT graph matrices with parameters corresponding to different degrees of data locality. See the text for details.

5.2. Computation and communication costs

We will here assume that the tasks are as far as possible evenly distributed over the computational nodes, i.e. that the total execution time is given by $\mathcal{O}(T_1/p + T_\infty)$ where T_1 and T_∞ are the serial and critical path execution times, respectively and p is the number of processes. Such load balancing can for example be achieved by work stealing [32]. While the total number of addition tasks is always smaller than the total number of multiplication tasks, the number of tasks along the critical path is up to $\mathcal{O}((\log(N))^2)$ for the additions and $\mathcal{O}(\log(N))$ for the multiplications. Therefore, we take T_1 to be proportional to the total number of multiplication tasks and $T_\infty = \mathcal{O}((\log(N))^2)$ given by the worst case critical path length. Thus, for the random case we have that the execution time is

$$\mathcal{O}\left(\frac{(\delta N^2)^{3/2}}{p} + (\log(N))^2\right) \quad (13)$$

and for the banded case we have that the execution time is

$$\mathcal{O}\left(\frac{d^2 N}{p} + (\log(N))^2\right). \quad (14)$$

We are interested in the communication costs in the weak and strong scaling limits. We consider first a weak scaling test constructed by keeping the number of nonzero matrix elements per row fixed and increasing the matrix dimension together with the number of processes. In the random case, this means that $\delta \propto \frac{1}{N}$ and the number of leaf level tasks is $\mathcal{O}(N)$ but the total number of tasks is $\mathcal{O}(N\sqrt{N})$. Assuming that all data for each task needs to be communicated this means that each process needs to receive data scaling as $\mathcal{O}(\sqrt{p})$ with the number of processes. In the banded and overlap cases both the leaf level and total number of tasks is $\mathcal{O}(N)$ and the average amount of data received per process is $\mathcal{O}(1)$. Since these results are based on the number of tasks, the latency cost behaves in the same way; the number of messages exchanged per process is proportional to the number of tasks per process. For strong scaling the number of tasks is constant, so the average number of tasks per process and thereby also the average amount of data communicated per process scales as $\mathcal{O}(1/p)$ for all sparsity structures.

The above results for the quadtree representation can be compared to the approach where a random permutation is employed to destroy data locality followed by application of the Sparse SUMMA algorithm [46]. Assuming that the random permutation succeeds to evenly distribute the nonzero matrix elements, the number of matrix elements that each process needs to fetch from other processes becomes (see e.g. equation (3.1) in [21])

$$\frac{2mN}{\sqrt{p}} \quad (15)$$

where m is the number of nonzeros per row. Similarly to the above, a weak scaling test can be constructed by keeping m fixed and letting $N \propto p$, leading to each process receiving data scaling as $\mathcal{O}(\sqrt{p})$ with the number of processes. The weak and strong scaling results are summarized in Table 1.

6. Performance evaluation

In this section, we will examine the performance of our block-sparse matrix-matrix multiplication presented in Section 3 when linked to the publicly available Chunks and Tasks library implementation CHT-MPI described in Section 2.1,

Table 1

Communication costs. Scaling of the average amount of data received by each process with the number of processes p in the weak and strong scaling limits for matrices with different sparsity patterns.

	Weak	Strong
Quadtree - random	$\mathcal{O}(\sqrt{p})$	$\mathcal{O}(1/p)$
Quadtree - banded	$\mathcal{O}(1)$	$\mathcal{O}(1/p)$
Quadtree - overlap	$\mathcal{O}(1)$	$\mathcal{O}(1/p)$
SpSUMMA	$\mathcal{O}(\sqrt{p})$	$\mathcal{O}(1/\sqrt{p})$

Table 2

Practical peak performance figures for cuBLAS batched matrix-matrix multiplication in Cuda 6.5 on Nvidia K20. Computed from batches with 64,000 matrix-matrix multiplications, $C_i = \beta C_i + \alpha AB$, $i = 1, 2, \dots, 64000$ with A , B , and C_i being $b \times b$ matrices with $b = 16, 32, 48, 64, 80, 96$. The number of floating point operations is counted as $64000 \times 2b^3$. We note that the figures are up to 40% larger than what we obtained with the benchmark program provided by Nvidia. This is mainly due to the reuse of the A and B matrices for all 64,000 multiplies in the present benchmark. Also, in the present benchmark timers on the GPU were used instead of timers on the CPU combined with synchronization. This results in larger Gflop/s values. Furthermore, switching from Cuda 5.0 to Cuda 6.5 gave performance improvements ranging from 20% for the “double precision, block size 16” case to 310% for the “single precision, block size 96” case, for calculations that were otherwise identical.

Matrix size	16	32	48	64	80	96
Gflop/s (single)	243.1	392.3	276.6	558.4	401.9	628.9
Gflop/s (double)	147.7	210.5	231.3	244.0	263.8	270.5

using the block-sparse leaf matrix library of [Section 4](#). Hereinafter, the number of floating point operations for multiplication of two dense matrices with dimension N is counted as $2N^3$.

6.1. Calculations on cluster of GPU-equipped nodes

We will first present calculations performed on the Erik cluster at the Lunarc computer center, Lund University, using CHT-MPI 1.1 compiled with Open MPI 1.6.5 and gcc 4.8.1, Cuda 6.5 and the Intel Math Kernel Library (MKL) version 11.1. The Erik cluster consists of 24 nodes each with dual 64-bit, 8-core Intel Xeon E5-2650 2.00 GHz processors. The nodes are interconnected with FDR InfiniBand and equipped with Nvidia Tesla K20m GPU cards: 16 nodes with 2 cards, 7 nodes with 4 cards, and 1 node with 8 cards. Leaf matrix submatrix operations were performed with the MKL implementation of BLAS on CPUs and with cuBLAS on the GPUs, as described in [Section 4.1](#).

Our first benchmark measures only the performance of the block-sparse matrix-matrix multiplication used for the leaf multiplications in the Chunks and Tasks matrix library. Thus, Chunks and Tasks is not involved in this benchmark. We perform multiplications with matrix dimension 4096×4096 with varying degree of matrix sparsity. The nonzero submatrix blocks are randomly uniformly distributed over the matrix to get a predetermined fill factor which is the fraction of nonzero matrix elements. Results for double and single precision are shown in [Figs. 5](#) and [6](#), respectively. Practical peak performance values (dashed lines) were calculated with a separate benchmark program measuring the performance of the cuBLAS batched matrix-matrix multiplication, see [Table 2](#). We are not quite reaching up to those peak figures. The main reason is that the benchmark of the leaf matrix library includes data transfers to/from the GPU which are not included in the peak performance figures. The work needed to prepare the list of batches is also included in the measured wall time. However, when the leaf matrix library is used within the Chunks and Tasks matrix library, there will be several threads that need to execute leaf matrix multiplications. This means that both communication to/from the GPU and preparation of batch lists on the CPU can then be overlapped with computation on the GPU as described in [Section 4](#).

This brings us to the next two benchmark figures where the Chunks and Tasks library is used but only on a single computational node, see [Figs. 7](#) and [8](#). The computational node is equipped with 16 CPU cores and 2 GPUs. Considering that the bounded priority queue has length 2 and that there are 2 slots per GPU, this means that up to 6 threads, no processing of batch lists will ever occur on the CPU cores. When 2 threads are executing tasks, there is one GPU dedicated to each thread, similarly to the previous benchmark figures. Therefore, the performance improvement when going from 2 to 6 threads comes solely from overlapping data transfers to/from the GPUs and preparation of batch lists on the CPU with the processing of batch lists on the GPUs. Comparing the practical peak performance limits to the performance for 6 threads in [Figs. 7](#) and [8](#) shows that the performance for 6 threads reaches between 88% and 95% of the practical peak performance values. When increasing the number of threads further, batch lists will also be processed on the CPU cores, according to [Algorithm 3](#). The figures show that we are able to take advantage of both the CPU and GPUs in a load balanced manner. We also note that no parametric models for task execution times on different hardware were required, the load balancing was achieved automatically without any information about the computational power of the devices, other than the assumption

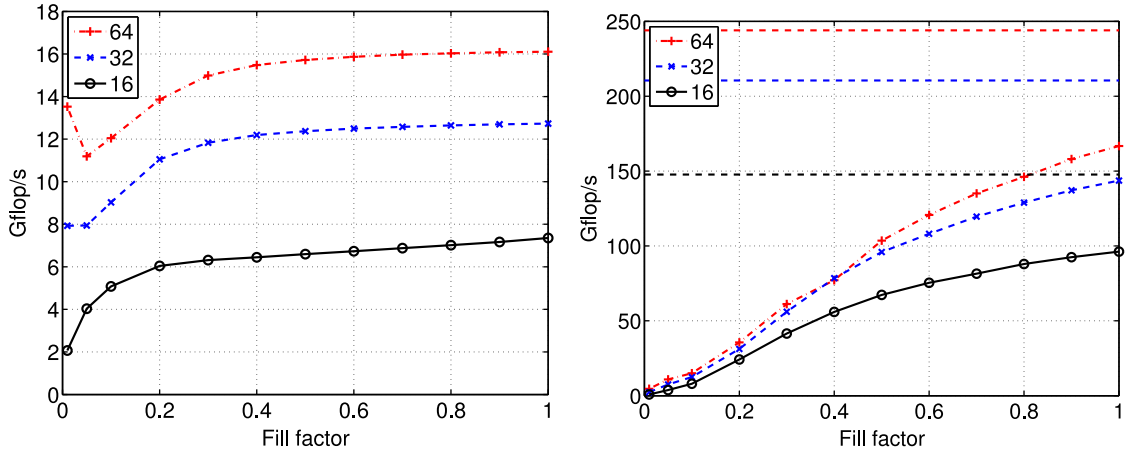


Fig. 5. Results of leaf block-sparse matrix library matrix-matrix multiplication test runs with double precision and matrix dimension 4096×4096 , varying sparsity (fill factor), and blocksizes 16, 32, and 64 on the Erik cluster. The nonzero submatrices are randomly uniformly distributed over the matrix. Left: running on one of the CPU cores. Right: running on one of the CPU cores but processing the list of batches on one of the GPUs. The dashed lines are practical peak performance figures computed from batches with $64,000 \times b$ multiplies with $b = 16, 32, 64$, not including any data transfers, see Table 2.

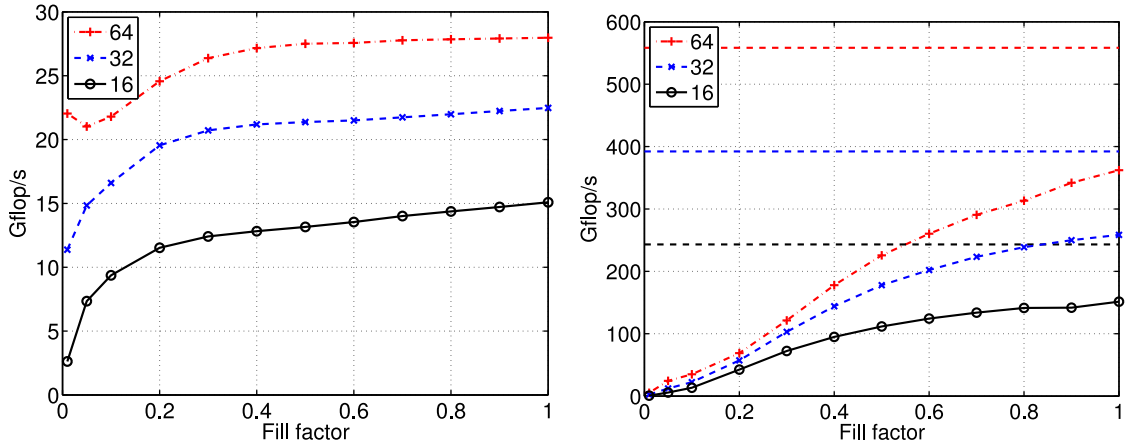


Fig. 6. Test runs for single precision corresponding to the double precision results in Fig. 5. See that figure caption for more information.

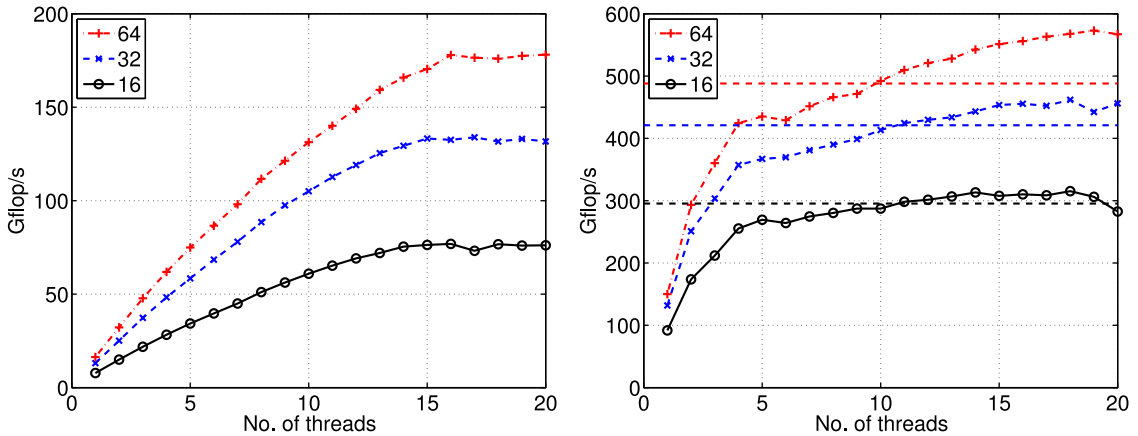


Fig. 7. Results of matrix-matrix multiplication test runs in double precision using the Chunks and Tasks matrix library for dense $25,000 \times 25,000$ matrices and blocksizes 16, 32, and 64 on a single Erik node equipped with two GPUs. A varying number of threads was used by CHT-MPI to execute tasks. The leaf matrix dimension was fixed to 4096×4096 . Left: running on the CPU cores. Right: running on both the CPU cores and the GPUs. The dashed lines indicate practical peak performance figures for the two GPUs, see Table 2. Note that up to 6 threads all batch lists are executed on the GPUs, see the discussion in the text.

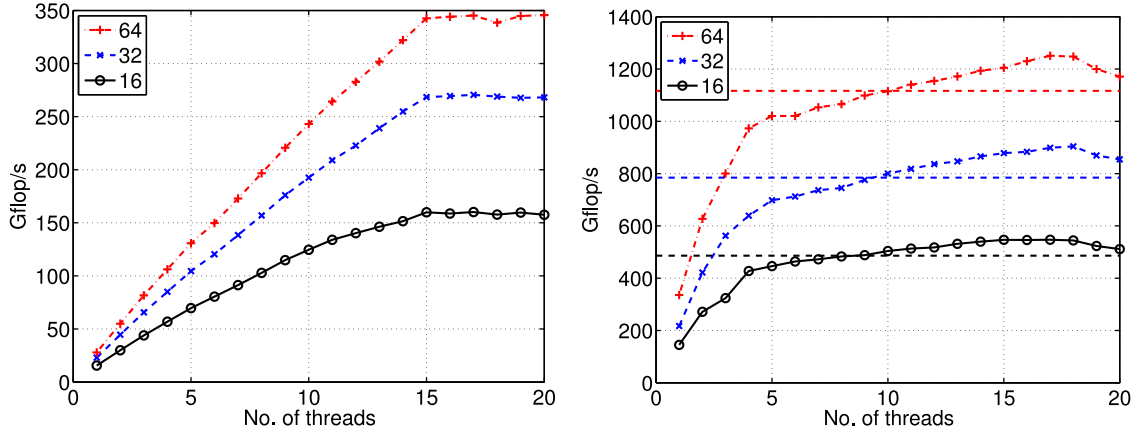


Fig. 8. Test runs for single precision and matrix dimension 40000×40000 , otherwise corresponding to the double precision results in Fig. 7. See that figure caption for more information.

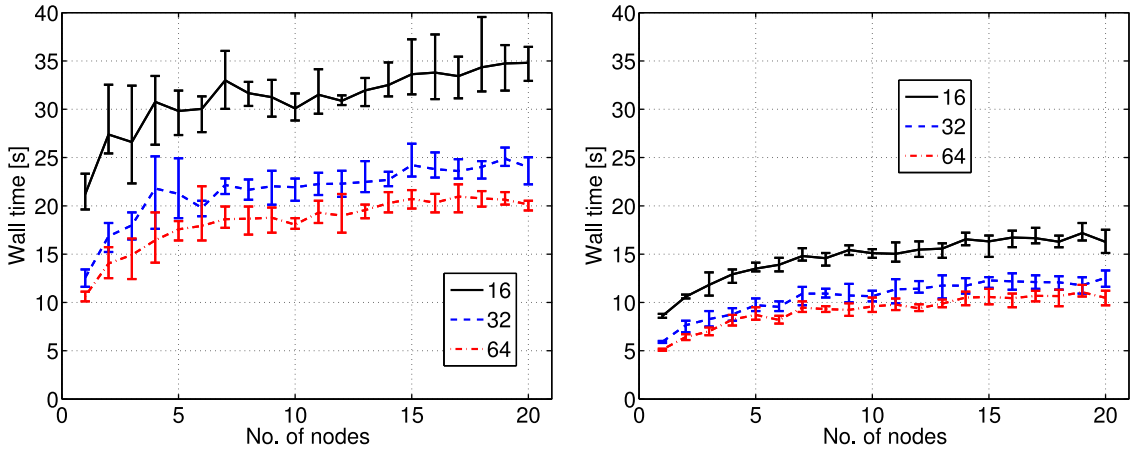


Fig. 9. Weak scaling test for banded matrices with bandwidth $2 \times 4000 + 1$ and matrix dimension $40,000 n \times 40,000 n$, where n is the number of nodes. The calculations were performed in double precision with leaf matrix dimension 4096×4096 for internal leaf matrix block sizes 16, 32, and 64. Left: Regular matrix-matrix multiplication. Right: Symmetric matrix square taking advantage of that the product matrix is symmetric. For each case (no. of nodes and blocksize), the benchmark calculation was repeated 6 times, and we plot the smallest interval containing all 6 wall times. Lines are drawn through the average wall time of the 6 benchmark calculations.

that a GPU is much more powerful than a CPU core. Block-sparse matrix-matrix multiplication using GPUs has also been implemented in the distributed block-compressed sparse row library, but using custom computational kernels rather than the batched kernels in cuBLAS [22]. The performance results in Fig. 7 are comparable to those in Fig. 8 in [22].

In Fig. 9 we investigate the weak scaling of our block-sparse matrix-matrix multiplication for a set of banded matrices with fixed bandwidth but a matrix dimension that is increasing proportionally to the number of computational nodes. Each node used 17 worker threads and a chunk cache size of 5 GB. In this case, we noticed fluctuations in the execution time and have therefore carried out 6 test runs for each case (no. of nodes and blocksize). Note that according to the theoretical weak scaling results of Section 5, we would expect at worst $\mathcal{O}((\log(n))^2)$ scaling of the wall time with increasing number of nodes n . Fig. 9 shows results for both regular matrix-matrix multiplication and the symmetric matrix square operation that assumes upper triangular storage of a symmetric matrix and only computes the upper triangle of the symmetric product. The implementation of the symmetric matrix square operation is straightforward using Chunks and Tasks since all decisions regarding distribution of work and data are handled by the runtime library. The expected speedup of 2 compared to the regular multiplication is achieved.

The effective performance in Gflop/s for the tests in Fig. 9 can be computed based on the number of scalar multiplications and additions for multiplication of banded matrices with bandwidth $2d + 1$:

$$2(N(2d + 1)^2 - (5/3)d(d + 1)(2d + 1)). \quad (16)$$

For the 1-node case in the left panel of Fig. 9 this gives 222, 374, and 441 Gflop/s for block sizes 16, 32, and 64, respectively. For the 20-node case the corresponding numbers are 2933, 4258, and 5065 Gflop/s. These figures can be directly compared to the performance results in the right panel of Fig. 7. In particular, one can compare to the highest performance for each

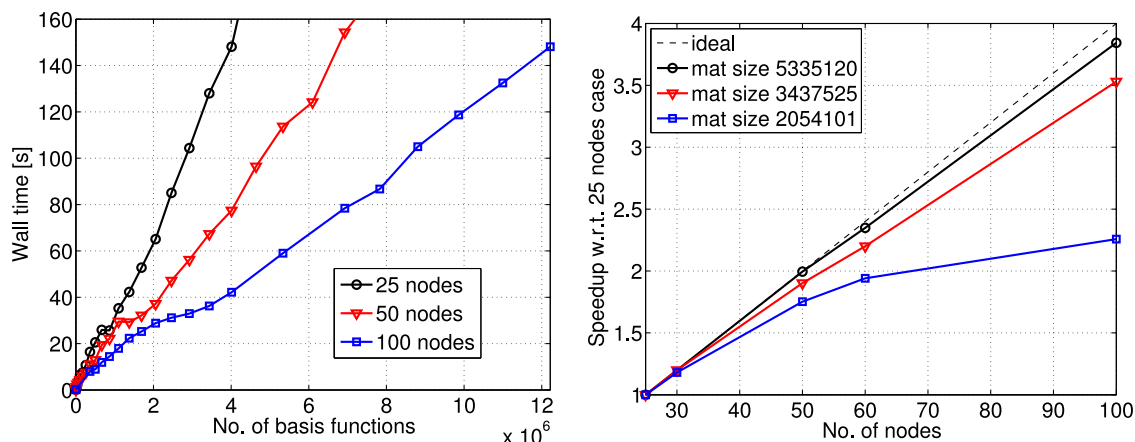


Fig. 10. Timings and scaling for S^2 symmetric matrix square computations on Tintin. Left: Timings for S^2 computations on overlap matrices for water clusters of varying size, using 25, 50, and 100 nodes of the Tintin cluster. Nearly linear system-size scaling is observed. Right: Scaling with respect to number of nodes, for three different matrix sizes. The speedups are relative to the 25 nodes case. We get closer to ideal speedup when the matrix size is increased.

block size. This shows that the banded matrix test runs in Fig. 9 retained 70%, 81%, and 77% of the performance for the 1-node case for block sizes 16, 32, and 64, respectively. For the 20-node case the corresponding numbers are 47%, 46%, and 44% compared to a perfect weak scaling scenario. Since the weak scaling efficiency per node has essentially leveled out at 20 nodes, similar efficiency can be expected also for larger calculations.

6.2. Application to overlap matrix in electronic structure program

To test the applicability of our block-sparse matrix-matrix multiplication in large-scale electronic structure calculations, we have adapted parts of the linear scaling electronic structure code Ergo [6] so that the overlap matrix can be constructed in parallel using Chunks and Tasks. This allows us to test the symmetric matrix square operation for the overlap matrix. See the description of the symmetric matrix square task type in Section 3.3.

The overlap matrix construction was done using a hierarchical representation of the basis set, where each part of the hierarchy contains basis functions located in a particular part of space. At higher levels in the hierarchy, chunk identifiers are stored referring to basis set descriptions at lower levels. Using such a hierarchical basis set description, it is straightforward to implement tasks to compute the overlap matrix.

The basis function ordering, which affects the sparsity pattern of the matrix, was determined based on the spatial coordinates of the basis functions using a recursive divide-space procedure. This is the default ordering used in the Ergo program.

The Ergo overlap matrix test calculations were performed on the Tintin cluster at the UPPMAX computer center, Uppsala University, using CHT-MPI 1.1 compiled with Open MPI 1.8.1 and gcc 4.9.1. The AMD Core Math Library (ACML) version 5.2.0 was used for BLAS operations on submatrices within the block-sparse leaf matrix implementation. The Tintin cluster consists of 160 compute nodes. Each node is a dual AMD Bulldozer compute server with two 8-core Opteron 6220 processors running at 3.0 GHz, with 64 GB of memory. The nodes are interconnected with a 2:1 oversubscribed QDR InfiniBand fabric. CHT-MPI was configured to use 15 threads for executing tasks, leaving one core on each node to handle communication. The chunk cache size was set to 8 GB.

The test molecules were water clusters generated from a molecular dynamics simulation of bulk water at standard temperature and pressure by including all water molecules within spheres of varying radii. The Gaussian basis set STO-3G was used, corresponding to 7 basis functions for each water molecule. The largest test system contained 1,745,413 water molecules, giving 12,217,891 basis functions. The overlap matrix S was truncated so that the Frobenius norm of the error matrix was smaller than 10^{-5} . The calculations were performed in double precision with leaf matrix dimension 4096 and blocksize 16. For the largest test systems this gave a sparsity corresponding to on average 1070 matrix elements per row in S after truncation, and about 7000 matrix elements per row in S^2 .

Timings and scaling for different numbers of compute nodes for the computation of S^2 using the symmetric matrix square operation are shown in Fig. 10. The time scales nearly linearly with the size of the molecular system, and the parallelization speedup improves for larger problem sizes.

Fig. 11 shows memory usage and communication statistics for the same water cluster S^2 calculations. The minimum, maximum, and average values among the 100 compute nodes are shown. Note that since CHT-MPI distributes both work and data dynamically, both the amount of data stored and the amount of communication needed will in general be different among the compute nodes. For the largest water cluster, the S^2 matrix contained about 8.6×10^{10} matrix elements. Since

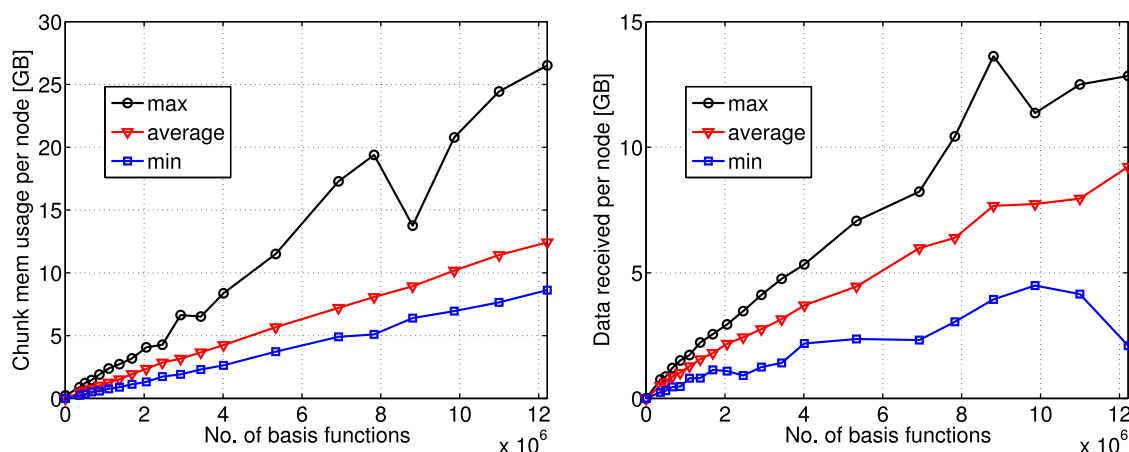


Fig. 11. Memory usage and communication statistics for S^2 symmetric matrix square computations for water clusters of varying size, using 100 nodes of the Tintin cluster. Left: Chunk storage peak memory usage. Only the memory for owned chunks is shown here; chunk cache is not included. Right: Amount of data received by each node during the symmetric matrix square operation.

only the upper triangle was computed and double precision was used, this corresponds to 344 GB of storage for S^2 , or about 3.4 GB per node for the 100 nodes case. The average chunk memory usage shown in the left panel of Fig. 11 is larger, about 12.4 GB per node, as it includes also temporary matrix chunks used during the computation.

Compared to the S^2 test calculations in [24], where plain dense matrix storage with blocksize 500 was used at the lowest level, the results in the present work represent significant improvements. The block-sparse leaf matrix type allows us to exploit sparsity much better, and using the symmetric matrix square operation reduces the computational effort even further. For a given water cluster size and number of compute nodes used, the memory usage for S^2 is reduced by about a factor of 16, and the time for the S^2 computation is reduced by about a factor of 6. Thanks to the reduced memory usage we are able to test significantly larger systems.

The results in this section demonstrate that our block-sparse matrix-matrix multiplication is indeed well suited for applications in large-scale electronic structure calculations; we get the desired linear scaling with respect to the size of the molecular system and reasonable parallel speedup for large enough problems, with dynamic distribution of both work and data. However, there is room for performance improvements. Statistics from the calculations indicate that the worker threads were typically idle more than half of the time, either waiting for data to be fetched from other nodes or because there was not enough remaining work to occupy all worker threads. This can be addressed by improvements within the CHT-MPI implementation, for example by running tasks closer to their input chunks. As seen in the left panel of Fig. 11, a more even distribution of the chunk storage in the CHT-MPI implementation would also be desirable, for example using an upper limit for the chunk storage on each node, and storing chunks elsewhere when that limit is reached. Such improvements could be taken advantage of without changes in the matrix library code, by linking to an improved CHT-MPI or another Chunks and Tasks library.

The compute nodes on the Tintin cluster where the Ergo tests were run are not equipped with GPUs, so the effect of using GPUs was not studied here. However, as seen in Section 6.1, our GPU implementation can provide additional performance in case GPUs are available.

Note that the Ergo test calculations presented here only involved the overlap matrix. Full Hartree-Fock or Kohn-Sham density functional theory calculations require additional parts of the Ergo code, notably the Coulomb and Hartree-Fock exchange matrix construction steps, to be parallelized using the Chunks and Tasks model. When that is done, the matrix library described here will be used to combine the different parts so that full Hartree-Fock and Kohn-Sham density functional theory calculations can be performed. The most performance-critical matrix operations are expected to be the symmetric matrix square operations during density matrix construction. The density matrix in general contains significantly more nonzero elements than the overlap matrix [11]. Therefore, for a given size of the molecular system more work for each matrix-matrix multiplication can be expected compared to the S^2 tests here, especially if larger basis sets are also used.

6.3. Investigation of communication costs

As shown in Section 5, the quadtree representation in principle allows data locality in sparse matrices to be exploited. Here we test this in practice by considering weak scaling using banded matrices similarly to the calculations presented in Fig. 9 with fixed bandwidth but a matrix dimension that is increasing proportionally to the number of worker processes. The average amount of communication for each worker process can then be expected to be constant, provided that the used CHT-MPI library succeeds in distributing the work.

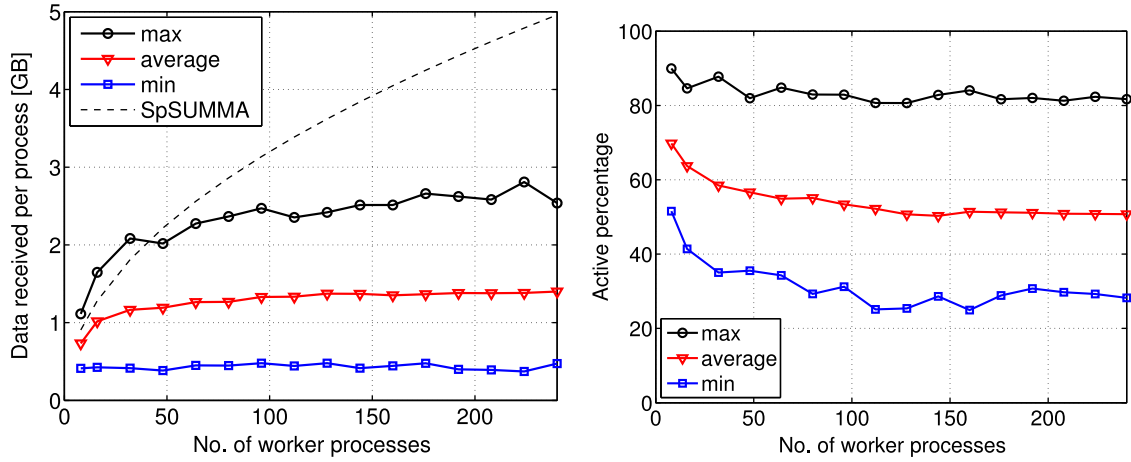


Fig. 12. Weak scaling results for regular matrix-matrix multiplication of banded matrices with bandwidth $2 \times 2000 + 1$ and matrix dimension $5000 p \times 5000 p$, where p is the number of worker processes. Left: Amount of data received by each process during the matrix-matrix multiplication operation. The dashed line indicates the amount of communication that would have resulted if a random reordering and the Sparse SUMMA algorithm had been used, see (17). Right: active percentage: the fraction of the time that worker threads were busy executing tasks.

However, if an approach like [21,22] is used, where a random permutation destroying data locality is employed, the average amount of communication needed will grow, as noted in Section 5.2. In this case, since the matrix dimension is increased together with the number of processes, by (15), the number of matrix elements that each process needs to fetch becomes

$$2mk\sqrt{p} \quad (17)$$

where k is the constant relating N and p in our weak scaling tests such that $N = kp$.

Since we are here interested in how the amount of communication scales for large numbers of processes, we use 8 worker processes per node and one worker thread per process. The test runs used up to 30 nodes of the Tintin cluster, corresponding to up to 240 worker processes. We consider here the total amount of data received by each process from other processes, including both communication between processes on the same node and between processes on different nodes. The chunk cache size for each process was set to 2 GB. The calculations were performed in double precision with leaf matrix dimension 4096 and blocksize 16.

Fig. 12 shows results of our weak scaling tests using the Chunks and Tasks matrix library. The minimum, maximum, and average values among the worker processes are shown. For each case, the plotted numbers are averages from 6 repeated benchmark calculations. For comparison, the amount of communication that would have been needed if a random reordering and the Sparse SUMMA algorithm had been used is also shown. We note that the Chunks and Tasks matrix library, without a priori knowledge about the sparsity structure, is able to take advantage of locality and achieve essentially constant amount of communication per worker process on average, as predicted in Section 5.2. To get an indication of the load balance, the active percentage for the worker threads, defined as the fraction of the time that worker threads were busy executing tasks, is also shown in the figure. When the communication per process no longer increases the active percentage is also stabilized. Note the difference between our locality-aware approach and the Sparse SUMMA algorithm, for which communication would have dominated completely for large enough test cases.

Fig. 13 shows the corresponding results for the symmetric matrix square operation. For large numbers of processes, using the symmetric matrix square operation instead of regular multiplication reduced the average necessary communication per process from about 1.39 GB to 0.76 GB.

Note that although the weak scaling tests described above were performed for the simple case of banded matrices, our Chunks and Tasks matrix library automatically takes advantage of any sparsity and locality that can be exploited by the quadtree structure. As another example, exploitation of data locality in the sparsity pattern improves the scaling of the amount of communication also for the more complex sparsity patterns occurring in electronic structure calculations for three-dimensional molecular systems. This can be seen by comparing some of the overlap matrix tests in Section 6.2. For example, going from 2,463,377 basis functions and 25 nodes to 9,861,383 basis functions and 100 nodes corresponds to a factor of 4 in the number of nodes and a factor of 4.003 in matrix size. The average amount of data received per node increased from 6.0 GB to 7.7 GB, or a factor of 1.28. This is a significant improvement compared to the factor of 2 that would have resulted from a \sqrt{p} behavior. It should be noted that such a comparison of different S^2 calculations does not correspond exactly to a weak scaling study, since the spherical shape of the water cluster systems and the use of Frobenius norm truncation lead to an amount of work that increases slightly more than linearly; in this case the increase in matrix size by a factor of 4 lead to an increase in the number of nonzeros in S^2 by a factor of 4.33.

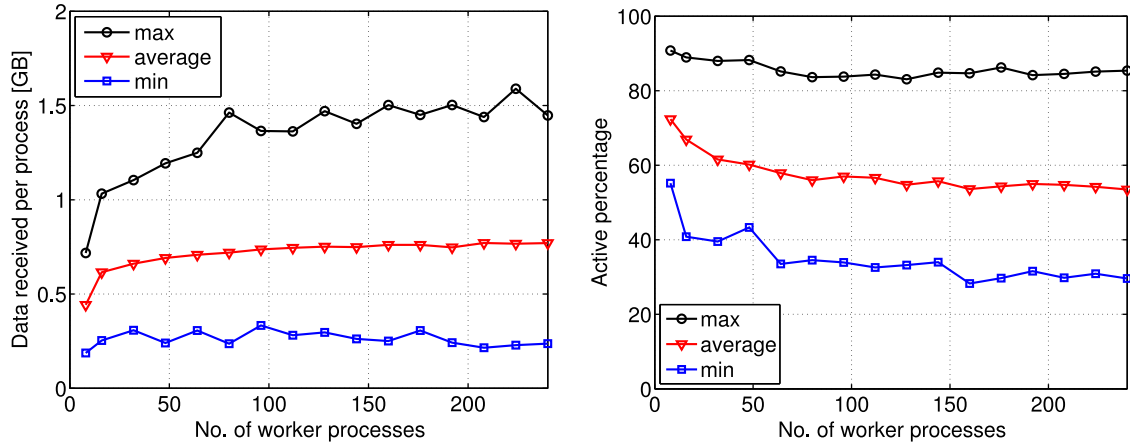


Fig. 13. Weak scaling results for symmetric matrix square computations corresponding to the regular matrix-matrix multiplication results in Fig. 12. See that figure caption for more information.

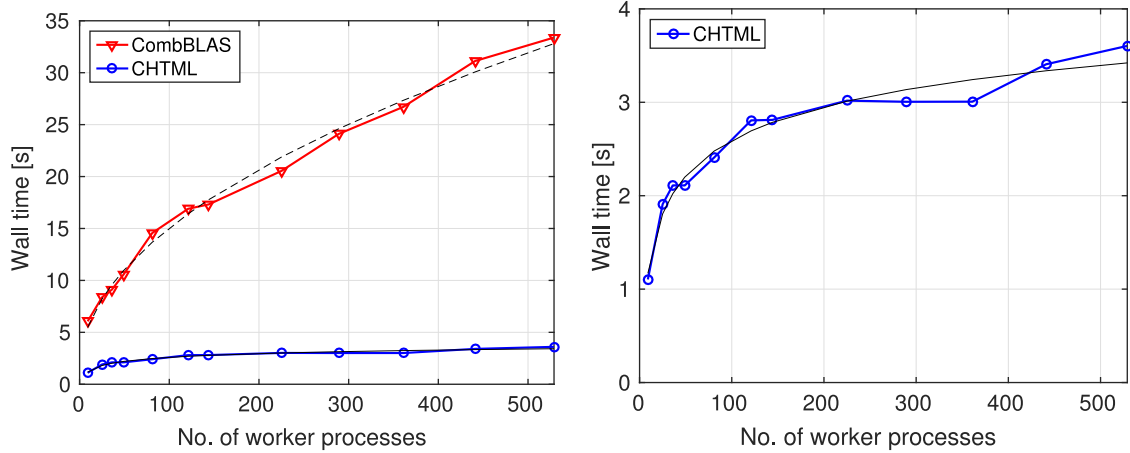


Fig. 14. Timings for weak scaling tests for regular matrix-matrix multiplication of banded matrices with bandwidth $2 \times 20 + 1$ and matrix dimension $20000p \times 20000p$, where p is the number of worker processes. Left: Wall times for Combinatorial BLAS (CombBLAS) compared to the Chunks and Tasks matrix library (CHTML). Right: Closeup of the CHTML timings. The dashed and solid help lines show $c_0 + c_1\sqrt{p}$ and $c_0 + c_1 \log(p) + c_2(\log(p))^2$ least squares fits for CombBLAS and CHTML, respectively.

Fig. 14 shows measured wall times for weak scaling tests on the Tintin cluster using the Combinatorial BLAS SpSUMMA implementation [47] and the here presented matrix library, in order to verify the theoretical results in Section 5.2 regarding execution times. The SpSUMMA tests were performed using the freely available and well organized software package Combinatorial BLAS version 1.4.0 [47]. Both Combinatorial BLAS and our matrix library were compiled with Open MPI 1.8.1 and gcc 4.9.1. The calculations used up to 16 worker processes per node, with the specific number of processes for each case fulfilling the Combinatorial BLAS requirement of a square logical processor grid. For our library, leaf matrix dimension 4096 and blocksize 16 was used and BLAS operations at leaf level were performed using ACML version 5.2.0. The test matrices were chosen so that the asymptotic scaling behavior for each approach can be clearly seen, using banded matrices of size $N = 20000p$ and bandwidth $2 \times 20 + 1$, in double precision. This makes the amount of work per process $\mathcal{O}(d^2N/p)$ relatively small. For SpSUMMA the $\mathcal{O}(\sqrt{p})$ communication cost dominates the calculation. SpSUMMA also involves a cost for processing the data that has been fetched. Owing to the efficient representation of hypersparse matrices provided by the doubly compressed sparse column data structure [48], used in Combinatorial BLAS, this cost is only $\mathcal{O}(\sqrt{p})$ as well. (In contrast, the standard compressed sparse column representation used in e.g. Matlab [49] and Csparse [50] would lead to an $\mathcal{O}(p)$ scaling for the processing of fetched data.) The inherent $\mathcal{O}(\sqrt{p})$ scaling behavior has also been demonstrated for SpSUMMA as implemented in the distributed block-compressed sparse row library, see e.g. Fig. 10 in [22]. For our quadtree based approach Fig. 14 shows a $c_0 + c_1 \log(p) + c_2(\log(p))^2$ least squares fit to the observed timings. We have noted that the c_2 coefficient is very small and even slightly negative, meaning that the $\log(p)$ term dominates in practice for this case. Thus, the observed behavior is slightly better than predicted by theory, see (14). A possible explanation is that although the total number of tasks along the critical path is $\mathcal{O}((\log(p))^2)$, the number of leaf level tasks along the critical path is

$\mathcal{O}(\log(p))$. When a large blocksize is used, leaf level tasks will be more expensive with respect to both computation and communication.

In summary, the results of our numerical experiments are consistent with the theoretical weak scaling results in Section 5.2: we observe constant average communication per process and execution times increasing only with the squared logarithm (or better) for matrices with data locality. In such cases, our approach will thus be increasingly favorable compared to SpSUMMA as the calculation size is scaled up.

7. Concluding remarks

The matrix library presented in this work is based on a quadtree data structure, with the important property that it allows for automatic exploitation of a priori unknown matrix sparsity structure. The hierarchical quadtree representation and associated recursive algorithms have been implemented using the Chunks and Tasks programming model. In this matrix library code, parallelism is exposed to the Chunks and Tasks library by expressing matrices and operations as hierarchies of chunk and task objects. All details regarding message passing and synchronization are left to the Chunks and Tasks library. Thus, the matrix library code can be written without worrying about how many nodes there are, where data is to be sent, etc. Storage and manipulation of matrices at the lowest level of the hierarchy is handled by a separate leaf matrix library. This means that the matrix library code is relieved from the details regarding the best way to store a particular type of submatrix or the best way to perform submatrix-submatrix multiplication on a particular type of hardware. Such modular design is powerful since it allows each part to be developed and optimized separately, and one can easily switch between different implementations of each part. Well designed interfaces between different modules, in our experience, results in both increased programming productivity and improved performance.

For matrices appearing in electronic structure calculations, the basis function ordering plays an important role in determining the sparsity pattern. For the overlap matrix test calculations in the present work, the default ordering in the ERGO program was used. A different ordering, using e.g. space filling curves [16] or network modularity optimization [51,52], could lead to increased data locality and result in improved performance.

In Section 5.2 we showed that for matrices with data locality our quadtree based approach yields superior weak scaling compared to the Sparse SUMMA algorithm [46]. It should be noted that there are algorithms that in theory could provide better scaling than SpSUMMA, see for example the so-called 3D algorithms that were theoretically discussed in [23]. Recently, a practical implementation of such a 3D algorithm was demonstrated [53]. In that work, similarly to SpSUMMA, a random permutation of rows and columns is used to achieve load balancing. However, instead of the two-dimensional $\sqrt{p} \times \sqrt{p}$ process grid used in SpSUMMA the algorithm makes use of a three-dimensional $c \times \sqrt{p/c} \times \sqrt{p/c}$ process grid. Following [23], c should be chosen as $c = \lceil p/m^2 \rceil$ in order to minimize communication, where m is the average number of nonzeros per row. This makes the 3D algorithm equivalent to SpSUMMA ($c = 1$) whenever $p \leq m^2$. This means that for all benchmark calculations presented in this article, where p is always smaller than m^2 , the scaling behavior for the 3D algorithm would in practice be identical to the scaling behavior of SpSUMMA. For example, for our tests in Fig. 14, where $m = 41$, the 3D algorithm would scale as SpSUMMA up to about 1600 processes. Although after this point the scaling behavior is expected to be improved compared to SpSUMMA, the method is still fundamentally different from our approach since it uses a predetermined data distribution and is unable to exploit data locality.

The possibility of using quadtrees in data-parallel or partitioned global address space languages such as High Performance Fortran (HPF) was discussed by Chatterjee et al. in [33]. Chatterjee et al. argued that representation of dense matrices using quadtrees could be incorporated in HPF but that recursion and nested dynamic spawning of computations would be difficult to achieve. In the context of the present work, representation of sparse matrices with a priori unknown sparsity patterns would represent a further obstacle.

This work illustrates the usefulness of programming models allowing dynamic distribution of work and data, which we expect will become increasingly important in the future, as larger compute systems are used. Apart from simplifying the implementation of dynamic algorithms, such models also make it easier to achieve fault resilience. They also facilitate the use of heterogeneous computational resources and allow robustness with respect to varying performance among compute nodes.

In the test calculations in this work, both generation of input matrices and verification of output matrices was performed using Chunks and Tasks programs. Thus, the data distribution of input matrices was a result of the task executions that generated those matrices. The placement of chunks was a result of work stealing as discussed in Section 2.1. Using the Chunks and Tasks matrix library together with other kinds of parallel software would require conversion of the data structures. Task types for carrying out such conversion without need for centralized administration of all data could be provided, although this was not done in the present work.

Finally, we note that the flexible design of the presented Chunks and Tasks matrix library makes it easy to modify and extend the library. For example, inclusion of extra information such as the Frobenius norm of each submatrix in the quadtree as in [54] could be straightforwardly implemented. The library could also be used for elementwise sparse matrices rather than block-sparse matrices by simply switching to a different leaf matrix library, employing for example a compressed sparse row format.

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