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Task-Based Parallel Programming for Scalable Algorithms: application to Matrix Multiplication

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Abstract: Task-based programming models have succeeded in gaining the interest of the high-performance mathematical software community thanks to how they relieve part of the burden of developing and implementing distributed-memory parallel algorithms in an efficient and portable way. In increasingly larger, more heterogeneous clusters of computers, these models appear as a way to maintain and enhance more complex algorithms. However, task-based programming models lack the flexibility and the features that are necessary to express in an elegant and compact way scalable algorithms that rely on advanced communication patterns. We show that the Sequential Task Flow paradigm can be extended to write a compact yet efficient and scalable General Matrix Multiplication. This extension required few modifications to the StarPU runtime system. The final implementation is shown to be competitive up to 32,768 cores with state-of-the-art libraries and may outperform them on some specific problem configurations.

Key-words: Mathematical software, parallelism, task-based programming, scalability, communication patterns, matrix multiplication

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Programmation parallèle à base de tâches pour algorithmes passant à l'échelle: application au produit de matrices

Résumé : Les modèles de programmation à base de tâches ont réussi à susciter l'intérêt de la communauté des logiciels mathématiques de haute performance grâce à la manière dont ils soulagent une partie du fardeau que représentent le développement et la mise en œuvre efficace et portable d'algorithmes parallèles à mémoire distribuée. Dans des grappes d'ordinateurs de plus en plus grandes et hétérogènes, ces modèles apparaissent comme un moyen de développer et maintenir des algorithmes plus complexes. Cependant, les modèles de programmation basés sur les tâches manquent de flexibilité et les caractéristiques nécessaires pour exprimer de manière élégante et compacte des algorithmes passant à l'échelle se basant sur des schémas de communication avancés. Nous montrons que le paradigme de flot de tâches séquentiel (STF) peut être étendu pour écrire une multiplication matricielle passant à l'échelle. Cette extension a nécessité peu de modifications au système d'exécution StarPU. L'implantation finale est compétitive jusqu'à 32 768 cœurs avec les bibliothèques de pointe et peut même les surpasser dans certaines configurations spécifiques.

Mots-clés : Logiciels mathématiques, parallélisme, programmation à base de tâches, passage à l'échelle, schémas de communication, multiplication matricielle

1 Introduction

Recent trends in supercomputer architectures have widened the gap between the speed of computations and the speed of data transfers. Modern supercomputers are composed of fat nodes equipped with many computational cores and specialized processing units (such as long vector units or GPUs) that can process operations at extremely fast rates. On the other hand, the speed of networks that interconnect these nodes has increased by a much smaller factor. For this reason, and more than in the past, a considerable effort is being devoted to the development of dense and sparse linear algebra algorithms that communicate better or communicate less. Although the literature of this subject is extremely vast, many of these scalable algorithms share some features. Computations are commonly arranged in such a way that the use of non-blocking or efficient collective communications is maximized van de Geijn and Watts (1997); Schatz et al. (2016). Data reduction patterns are used to increase parallelism and reduce the critical path of communications Demmel et al. (2012); van de Geijn and Watts (1997); Schatz et al. (2016). In some cases, communications are reduced at the price of a moderate or controllable overhead in terms of memory or operations as in 2.5D or 3D algorithms Schatz et al. (2016); Agarwal et al. (1995); Solomonik and Demmel (2011); Ashcraft (1993); Sao et al. (2019).

With algorithms becoming more and more complex and supercomputers architectures more heterogeneous, the need has raised for programming models that relieve the high performance computing expert from the burden of mixing multiple programming models and interfaces (e.g., MPI, OpenMP and CUDA) and allow them to focus on developing efficient and scalable algorithms. Task-based parallelism is one such model: it allows for a high level description of the workload in the form of a directed acyclic graph (DAG) where nodes correspond to tasks (i.e., elementary operations) and edges the dependencies among them or, equivalently, data transfers. This model has been widely used in the past; one notable example is the MUMPS Amestoy et al. (2001) sparse direct solver that implements task-based parallelism by means of the MPI programming interface. The task-based parallel programming paradigm has recently known a renewed interest thanks to the emergence of novel programming models that allow for a simpler construction of the DAG. Two well known examples are the sequential task flow (STF) where tasks dependencies are inferred from data access modes (more details on this model are provided in section 2.1) and parameterized task graph (PTG) where task dependencies are defined by means of rules provided by the programmer. These programming models are available, through dedicated programming interfaces, in various runtime systems (or, simply, runtimes) whose use in the high performance computational linear algebra domain is increasingly popular. Well known runtimes include StarPU Augonnet et al. (2011) (which provides an interface for the STF programming model), ParSEC Bosilca et al. (2013) (STF and PTG) and OpenMP from version 4.0 (STF). The use of task-based parallelism through modern runtimes has several attractive features. It allows for a relatively easy and portable programming of heterogeneous architectures including distributed memory multinode systems running over multi and manycores as well as accelerators such as GPUs: the runtime takes care of deploying tasks on the available processing units (according to a defined scheduling policy) and of transferring the data required for their execution through the network or a dedicated bus. This approach presents great

modularity. For example, it allows for choosing among pre-defined task scheduling policies or developing new ones; because these are implemented within the runtime, they are not tight to a specific algorithm or application but can be reused transparently. By the same token, as we will show below, it is easy to switch between different communication libraries. Numerous studies exist that demonstrate the effectiveness of this approach for different applications and algorithms mostly on shared memory systems Agullo et al. (2016) but also, and increasingly, on distributed memory ones Herault et al. (2019). Nevertheless, concerns still remain about the expressiveness of task-based parallelism and programming models and their capability to implement distributed memory algorithms with complex features such as those mentioned above. The purpose of our work is to address these concerns and show that, through a suitable extension of the state-of-the-art STF model for distributed memory machines Agullo et al. (2017a), it is possible to implement complex scalable algorithms; the resulting code is easy to maintain and improve yet very efficient and portable. As a reference, we will use the dense generic matrix-matrix product (GEMM) and its distributed memory parallel variants such as the SUMMA van de Geijn and Watts (1997) and 2.5D Solomonik and Demmel (2011) algorithms which possess most of the above mentioned algorithmic features. Experimental results will show that the proposed approach provides equivalent, if not better, performance than reference libraries with a code which is only slightly more complex than the classic three nested loops of a basic, sequential, matrix multiplication code.

2 Background

2.1 The sequential task flow programming interface

As explained above, task-based parallelism can be conveniently implemented using specifically designed programming models such as sequential task flow (STF) or parameterized task graph (PTG). In this document we will focus on the STF one, sometimes also referred to as *superscalar* since it mimics the functioning of superscalar processors where instructions are issued sequentially from a single stream but can actually be executed in a different order and, possibly, in parallel depending on their mutual dependencies. The STF model relies on a task insertion or submission primitive which allows for creating a task. The insertion is non blocking, which means that the control is immediately returned to the caller and the execution of the task is deferred. Upon insertion of a task, the caller must specify the data used by the task and whether the task accesses these data in read (R), write (W) or read-write (RW) mode. Based on the order in which tasks are inserted and their data access modes, dependencies between tasks can be easily determined and the DAG of tasks automatically built.

In the case of a shared memory parallel computer, the STF model commonly relies on the use of a *master* process which is in charge of inserting the tasks and multiple *worker* processes which are in charge of executing them. Multiple types of workers may exist if different processing units are available such as CPU cores and GPUs although in our work we will not cover the case of heterogeneous systems. In the case of a distributed memory machine, multiple masters

exist which communicate by exchanging messages; these communications can be internally implemented by the runtime through the MPI standard but other communication libraries can also be used. These communications essentially correspond to dependencies between tasks that are executed by workers associated with different masters. For this reason, in the most basic use of the STF model, all masters must insert all the tasks of DAG to make sure these communications are correctly detected and executed. This corresponds to the approach based on a concurrent unrolling of the task graph proposed by YarKhan (2012).

This programming model is commonly appreciated because of its simplicity which allows, in a relatively easy way, to transform a sequential code into a parallel one while preserving its readability and maintainability. This advantage must, however, be weighted against potential limitations due to the fact that the DAG must be entirely unrolled by inserting all of its tasks: not only this can be time consuming, but it may require considerable resources for the management of the DAG when it is of large size. Several techniques have been proposed in the literature to alleviate this issue such as pruning of its traversal Agullo et al. (2017a) or hierarchical tasks. The PTG model, on the other hand, has better scalability because the DAG is not explicitly and entirely built but, instead, tasks are efficiently instantiated based on rules defined by the programmer; this, however, comes at the price of a considerably higher programming effort Agullo et al. (2017b).

2.2 Distributed memory scalable GEMM algorithms

The general matrix-matrix multiplication (GEMM) operation, as defined in the BLAS standard, consists in computing

$$C = \alpha \cdot op(A) \cdot op(B) + \beta \cdot C$$

with $C \in \mathbb{K}^{M \times N}$, $op(A) \in \mathbb{K}^{N \times K}$, $op(B) \in \mathbb{K}^{K \times N}$, $op(.)$ being either the identity or the (conjugate) transpose and α and β scalars in \mathbb{K} - either real or complex. Without loss of generality, in the remainder of this paper we drop the $op(.)$ operator (and, thus, assume that neither A nor B are transposed) and assume that both α and β are equal to one.

For the purpose of the parallelization, we will suppose that all matrices are partitioned into blocks of size b and that $m = \lceil M/b \rceil$, $n = \lceil N/b \rceil$ and $k = \lceil K/b \rceil$. This will allow us to use efficient sequential BLAS routines for computations on blocks. Based on this assumption, the sequential matrix multiplication can be simply written as the triply nested loop in Figure 1, where the instruction in the inmost loop computes $C_{i,j} = C_{i,j} + A_{i,l} \times B_{l,j}$. Ignoring the data locality issues in NUMA memory configurations, this code can be trivially parallelized for shared memory parallel computers using, for example, loop parallelism or task-based parallelism (more on this will be said in the next section).

When targeting distributed memory parallel computers, the A , B and C matrices must be distributed among the ranks that participate in the computation. Here, and in the remainder of the article, we use the generic word *rank* to denote processes that communicate by exchanging messages; a rank corresponds to a master process in the STF model or to a MPI process in the widely used MPI programming interface. We will assume that a 2D block-cyclic distribution over a $p \times q$ ranks grid is employed because of its wide use in reference dense linear


```

1  do i=1, m
      do j=1, n
3      do l=1, k
          call gemm(Ai,l, Bl,j, Ci,j)
5      end do
      end do
7  end do

```

Figure 1: Sequential, blocked GEMM

algebra libraries (including ScaLAPACK) and because it complies well with the scalable GEMM algorithms described below; for the sake of simplicity, we will also assume that all matrices are aligned, i.e., have a conforming distribution across the ranks grid.

Despite its large arithmetic intensity, the scalability of the GEMM operation on large size supercomputers can be severely limited by the slowness of network communications and many algorithms have been proposed in the literature to overcome this limitation. The Cannon’s algorithm Cannon (1969), for example, has been proved to minimize both the communication bandwidth and latency Irony et al. (2004); Ballard et al. (2011). Nevertheless, this algorithm only works on square processors meshes and is, therefore, unpractical. SUMMA Schatz et al. (2016); van de Geijn and Watts (1997); Agarwal et al. (1994) overcomes these limitations of the Cannon’s algorithm and has become the most widely adopted algorithm in reference parallel dense linear algebra libraries such as ScaLAPACK Blackford et al. (1997) or PLAPACK van de Geijn (1997). In the SUMMA algorithm, shown in Figure 2, the matrix product is defined as a sequence of outer products where at each iteration $l = 1, \dots, k$ the l -th column of A is multiplied with the l -th row of B and the result added to C . Each $r \times c$ rank computes the contribution for the $C_{i,j}$ blocks it owns and, therefore must receive the corresponding $A_{i,l}$ and $B_{l,j}$ blocks; the outer product formulation allows to transfer these blocks using efficient collective communications: the $A_{i,l}$ block is broadcasted to all the ranks in the r -th grid row and the $B_{l,j}$ block is broadcasted to all the ranks in the c -th grid column. A pipelined version of this algorithm was also proposed van de Geijn and Watts (1997) which further reduces the length of the critical path of the parallel matrix product; however, if non-blocking collective communications are available, the interest of this variant is limited with respect to the basic one.

The SUMMA algorithm presented above is particularly efficient in the case where the C matrix is much larger than A and B because only these two are transferred whereas C stays in place; for this reason we refer to this algorithm as *stationary C* (or stat-C, for short) following the notation proposed by Schatz et al. (2016). Stationary A or stationary B variants can be used in the case where A or B are larger than the other two matrices, respectively; because these two variants behave the same, we only present the first one here. In this algorithm, reported in Figure 3, the matrix-matrix product is defined as a sequence of matrix-panel products where, at each step, the entire A matrix is multiplied by a $B_{*,j}$ block-column producing a $C_{*,j}$ block-column. In this

```

1  do l=1, k
    forall i, i%p==r :
3      bcast( $A_{i,l}$ , to:r×*)
    forall j, j%q==c :
5      bcast( $B_{l,j}$ , to:*×c)
    forall i, i%p==r :
7      forall j, j%q==c :
        call gemm( $A_{i,l}$ ,  $B_{l,j}$ ,  $C_{i,j}$ )
9  end do

```

Figure 2: Stationary-C SUMMA algorithm as executed by the $r \times c$ rank.

```

1  do j=1, n
    forall l, l%p==r, l%q==c :
3      recv( $B_{l,j}$ , from:r×j%q)
    forall l, l%p==c :
5      bcast( $B_{l,j}$ , to:*×c)
    forall i, i%p==r :
7      forall l, j%q==c :
        call gemm( $A_{i,l}$ ,  $B_{l,j}$ ,  $C_{i,j}^h$ )
9      reduce( $C_{i,j}^h$ , to:i%r×j%q)
    end do

```

Figure 3: Stationary-A SUMMA algorithm as executed by the $r \times c$ rank.

case the A matrix stays in place, the B matrix is transferred using efficient collective communications and locally computed contributions to the C matrix (denoted $C_{i,j}^t$) are assembled using reductions. Note that in this algorithm, because of the cyclic data distribution, the `recv` and `bcast` communications in lines 3 and 5 can be more efficiently implemented using scatter and allgather primitives Schatz et al. (2016).

The scalability of the SUMMA algorithm can be further improved using so-called 2.5D or 3D algorithms Georganas et al. (2012); Schatz et al. (2016). In these algorithms we consider the ranks arranged in a three-dimensional grid of size $p \times q \times s$ and the A , B and C matrices initially distributed among the ranks in the lowest level (0) of this grid, i.e., $* \times * \times 0$. The pseudo-code for the 2.5D stat-C case executed by the $r \times c \times h$ rank is reported in figure 4. Here the A and B matrices are partitioned in s parts along the k dimension (columns and rows, respectively) and each part is replicated on one of the higher levels $1, \dots, s-1$ where a partial stat-C matrix product is computed producing local C^h contributions to the final result. The local contributions are finally assembled into the C matrix using reductions. Clearly, equivalent 2.5D algorithms can be formulated for stat-A or stat-B SUMMA; we refer the reader to the paper by Schatz et al. (2016) for the related details.

```

forall j, h*k/s ≤ j < (h+1)*k/s, j%q==c:
2   forall i, i%p==r :
      recv(Ai,j, from:r×c×0)
4
forall i, h*k/s ≤ i < (h+1)*k/s, i%p==r:
6   forall j, j%q==c :
      recv(Bi,j, from:r×c×0)
8
do l=h*k/s, (h+1)*k/s-1
10  forall i, i%p==r :
      bcast(Ai,l, to:r×*×h)
12  forall j, j%q==c :
      bcast(Bl,j, to:*×c×h)
14  forall i, i%p==r :
      forall j, j%q==c :
16      call gemm(Ai,l, Bl,j, Ci,jh)
end do
18
forall i, i%p==r :
20  forall j, j%q==c :
      reduce(Ci,jh, to:r×c×0)

```

Figure 4: 2.5D stationary-C SUMMA algorithm as executed by the $r \times c \times h$ rank.

```

1  do i=1, m
      do j=1, n
3      do l=1, k
            call insert_task(gemm, Ai,l:R, Bl,j:R,
5      Ci,j:RW)
      end do
      end do
7  end do

```

Figure 5: Parallel GEMM using the baseline STF model.

3 STF matrix multiply

3.1 Baseline STF model

The GEMM operation of Figure 1 can be straightforwardly parallelized using the STF model by replacing the calls to the sequential `gemm` on blocks with task insertions through the `insert_task` routine; this delegates the execution of the corresponding operations to the runtime systems. The first argument of the `insert_task` method is the operation to be performed upon task execution and is followed by the list of data used by the task (here, the $A_{i,l}$, $B_{l,j}$ and $C_{i,j}$ blocks). For each data, the corresponding access mode is specified through a colon notation: R and RW denote, read and read-write access modes, respectively. Based on the task insertion order and the tasks data access mode, the runtime can infer the tasks dependencies, build the corresponding DAG and proceed to schedule its tasks on the available processing units.

Thanks to the very high arithmetic intensity of the GEMM operation, the code of Figure 5 can achieve very good performance on shared memory, possibly accelerated (e.g., with GPUs) systems, provided that a suitable block size is chosen which provides a good trade-off between parallelism and efficiency of tasks. Additionally, in order to run this code on distributed memory parallel systems, it is enough to make the runtime aware of the data distribution; for example, in the case of a 2D block-cyclic distribution each (i, j) block of A , B and C is assigned to rank $(i\%p) \times (j\%q)$ of the $p \times q$ ranks grid, where $\%$ is the modulus operator. In this case, the runtime will take care of transferring over the network the blocks needed by a task on the rank where the task is executed. Although this code, based on the baseline model proposed in Agullo et al. (2017a), will be perfectly functional, its performance and scalability can be poor compared with what can be achieved with the algorithms described in section 2.2 for a number of reasons.

First of all, this code will not be able to make use of collective communications. In this baseline STF model, communications are expressed by the edges of the DAG which, essentially, define point-to-point data transfers.

Second, this baseline STF model does not allow any control on the mapping of tasks over the $p \times q$ ranks of the grid. Runtimes implement basic mapping policies where a task is executed on the rank which owns the data that is accessed in read-write mode ($C_{i,j}$, in the case of Figure 5). This can lead to a very large volume of communications, for example, in the case where A and B are much

larger than C and will not allow us to use computing ranks that do not own blocks of the C matrix.

Finally, in this baseline STF model it is not possible to take advantage of the commutativity and associativity of certain operations. In our case, it must be noted that all the summations in tasks of the type $C_{i,j}+ = A_{i,l} \cdot B_{l,j}$ for all l can commute or be grouped in any way. This property can be used to improve parallelism or reduce communications. In the baseline STF model, instead these summations are forced to be executed sequentially: because of the order in which tasks are inserted and of the RW access mode on the $C_{i,j}$ block, the task that computes $C_{i,j}+ = A_{i,l+1} \cdot B_{l+1,j}$ depends on the one computing $C_{i,j}+ = A_{i,l} \cdot B_{l,j}$.

3.2 Improved STF model

The limitations discussed in the previous section, make the baseline STF model Agullo et al. (2017a) unsuitable for implementing scalable algorithms for distributed memory systems such as those presented in section 2.2. It must be noted that it is possible to get around some of these limitations through careful programming. For example, it is possible to take advantage of associativity of some operations by declaring temporary data and explicitly inserting tasks to combine partial results; in other words, this amounts to manually implementing reduction operations. This practice, however, leads to complex code which is poorly portable and hard to maintain which, essentially, defeats the purpose of using a high level task-based parallel programming model. The purpose of this section is to present an extension of the baseline STF model of section 3.1 including features that allow us to implement scalable algorithms. These features extend both the programming interface and the functionality of an STF-based runtime system while preserving the high level expressiveness of the STF model and, ultimately, the portability and maintainability of the code.

Reduction tasks The objective of this feature is to provide a mean of taking advantage of the associativity and commutativity of the block-sum operation to improve parallelism. As explained above, this is not possible in the baseline STF model because for all the tasks that compute a $C_{i,j}+ = A_{i,l} \cdot B_{l,j}$ contribution are inserted with RW (read-write) access mode on the $C_{i,j}$ block; this induces a chain of dependencies on these tasks according to the order in which they have been inserted. One way to overcome this problem is to introduce a new access mode, which we call **REDUX**. With this access mode, each task will assemble its contribution in a temporary block $D_{i,j}^f$, $f = 1, \dots, z$; the runtime system takes care of creating all the z temporary blocks and combining their content through dedicated tasks in a transparent way. This reduction phase is carried asynchronously, the only constraint being that it must be completed prior to any other access to the $C_{i,j}$ block with a different access mode; parallelism is also available in this phase which can be used through suitable reduction trees. For this feature to work, it is necessary that the runtime is informed of how to initialize the temporary blocks and how to combine their values. This can be achieved by declaring to the runtime two methods, called the *initializer* and the *combiner* (to follow the naming in the OpenMP standard). It must be noted that a crucial design choice concerns the number z of temporary copies $D_{i,j}^f$

for each $C_{i,j}$ block. One naive choice could be to have as many copies as the number of tasks that compute a contribution to the $C_{i,j}$ blocks, that is k ; in this case all these tasks will be independent and could potentially be executed in parallel. The downsides of this choice are, first, that it may result in an excessive memory consumption due to the high number of temporary copies and, second, a costly reduction step; besides this may result in an unnecessarily high amount of parallelism. A wiser choice is to have as many temporary copies as the number of workers that participate in the computation of a $C_{i,j}$ block; although to a much lower extent than in the previous option, this choice might still suffer from the same issues considered that current supercomputers have very fat nodes equipped with up to hundreds of cores. In the present work we have chosen to have as many copies as the number of ranks participating in the computation of a $C_{i,j}$ block, which means that all the workers associated with a rank will share the same $D_{i,j}^f$ temporary copy. However, to use additional parallelism, we allow all tasks using the same $D_{i,j}^f$ copy to commute, that is, to execute in any order but not concurrently.

Dynamic collective communications In order to take advantage of efficient and scalable collective communications, whose role is essential in the algorithms presented in section 2.2, we rely on the so-called *dynamic collectives* feature proposed by Denis et al. (2020). This approach consists in automatically detecting that a data must be transmitted from a source rank to multiple destination ranks; when such a pattern is detected, the corresponding transfers are grouped together and achieved through a collective communication. This feature has a number of interesting properties. First of all, it is completely transparent to the user: no change has to be done on the user-level code but the detection and use of collective communications happen in the communication library underlying the runtime system. Second, these collective communications do not rely on the use of subcommunicators which has several advantages as we will explain below. Finally, dynamic collectives are non-blocking which allows for an effective overlapping of communications and computations.

Tasks mapping This feature amounts to binding one task to one rank, which means that it can be executed by any of the workers associated with that rank. This is simply achieved through an additional `ON_RANK` argument to the `insert_task` routine, which defines the identifier of the rank where the task has to be run on. This feature is essential for the stationary-A and 2.5D variants where the placement of tasks is not trivially related to the initial data distribution.

3.3 Scalable GEMM with the extended STF model

Using the improved STF model including the features presented in section 3.2, all the GEMM algorithms of section 2.2 can be conveniently implemented as in the pseudo-code of Figure 6. The `map_and_am` function returns the rank where a (i, j, l) task must be executed and the access mode on the $C_{i,j}$ block for this task: depending on the requested variant, specified by the `stat` and `s` (number of grid levels) arguments, this function will return the corresponding mapping and access mode. It must be noted that the same result will be obtained regardless

```

1  do i=1, m
      do j=1, n
3       do l=1, k
            RANK, ACC = map_and_am(i, j, l, stat, s)
5            call insert_task( gemm,
                        Ai,l:R,
7                        Bl,j:R,
                        Ci,j:ACC,
9                        RANK:ON_RANK)
            end do
11        end do
    end do

```

Figure 6: Parallel GEMM using the improved STF model.

of how the i , j , and l loops are nested because in all cases the DAG of tasks would be, essentially, the same. For the sake of readability, in the pseudo-code of Figure 6 we have omitted the declaration of the initializer and combiner routines for the reductions; these correspond, respectively, to zeroing out all the coefficients of a block and summing two input blocks into an output one.

The central claim of this paper is that the proposed extended STF model allows one to express the three advanced GEMM algorithms (and communication patterns) described in section 2.2 with this extremely compact and simple code (Figure 6), together with an appropriate choice for the mapping and access modes. In the following paragraphs, we present the mapping and access mode corresponding to each of these three GEMM algorithms for completeness.

Stationary-C SUMMA In the stationary-C SUMMA algorithm the rank owning the $C_{i,j}$ block is in charge of all the $C_{i,j} += A_{i,l} \cdot B_{l,j}$ tasks for $l = 1, \dots, k$. Therefore, assuming a 2D block-cyclic distribution on a $p \times q$ grid, the RANK output of the `map_and_am` function will be $(i\%p, j\%q)$. As for the access mode of the $C_{i,j}$ block, this is read-write RW; however, to improve parallelism and take advantage of the fact that each rank has multiple workers, we can make all the tasks related to the same $C_{i,j}$ block commutable. The dynamic collectives feature will transparently group together all the transfers of a $A_{i,l}$ ($B_{l,j}$) along the $i\%p$ ($j\%q$) ranks row (column) and perform them using an efficient collective communication; this corresponds to the broadcast communications in lines 3 and 5 of the pseudo-code in Figure 2.

Stationary-A SUMMA In the stationary-A SUMMA variant, the rank in charge of computing $C_{i,j} += A_{i,l} \cdot B_{l,j}$ is the one that owns the $A_{i,l}$ block; therefore, the RANK returned by the `map_and_am` function is $(i\%p, l\%q)$. As for the access mode for the $C_{i,j}$ block, this has to be REDUX to achieve the reduction on line 9 of the pseudo-code in Figure 3. The dynamic collectives feature will detect that the $B_{l,j}$ block has to be sent to all the ranks in the $l\%q$ grid column and achieve these transfers with an efficient broadcast communication

corresponding to the `recv` and `bcast` communications in lines 3 and 5 of the pseudo-code in Figure 3. It must be noted that in MPI-based implementations, the broadcasting of $B_{l,j}$ to the $l\%q$ grid column must be done in two steps because the rank owning this block does not necessarily belong the $l\%q$ grid column sub-communicator. Because the dynamic collectives feature does not rely on the use of sub-communicators, the `recv` communication in line 3 of Figure 3 is not necessary.

2.5D Stationary-C SUMMA In the 2.5D stationary-C SUMMA variant, the rank selected by the `map_and_am` function for computing the contribution $A_{i,l} \cdot B_{l,j}$ to the $C_{i,j}$ block is $(i\%p, j\%q, l/(k/s))$. The access mode to the $C_{i,j}$ block is `REDUX` to operate the reductions in line 21 of the pseudo-code in Figure 4. The broadcasts in lines 11 and 13 of Figure 4, are performed straight from the lowest grid level through dynamic collectives without the need for the preliminary point-to-point communications of lines 3 and 7. Similarly to what explained above for the stationary-A variant, these point-to-point communications are necessary to move data on a rank belonging to the sub-communicator where the broadcast happens; because in the dynamic collectives feature the scope of a collective communication is arbitrary and not defined by a sub-communicator, these preliminary copies are unnecessary.

4 Implementation

In this section we discuss the practical implementation of the pseudo-code of Figure 6 which we achieved using the StarPU runtime system and its STF programming API within the `qr_mumps` Agullo et al. (2016) library.

4.1 STF advanced features

The proposed implementation of Figure 6 makes use of the features described in section 3.2. In this section we discuss the availability and use of these features in the StarPU runtime and, in the case of the reduction tasks feature, some improvements that we have implemented in order to achieve better performance. The task mapping feature is already available in the latest StarPU releases and will not be discussed any further.

The second feature, the dynamic detection of collective communications, is available, through the NewMadeleine library; the reader is referred to the recent work by Denis et al. (2020) for a thorough discussion of this feature. This mechanism is implemented such that the communication pattern used to achieve collective communications can be chosen at run time through an environment variable. In all our experiments we have used a binomial tree. It must be noted, though, that the use of a chain (where each rank forwards the message to only one other rank) essentially leads to an asynchronous implementation of the pipelined SUMMA algorithm (van de Geijn and Watts, 1997, sec. 5.2). We reserve the analysis of this approach for future work.

Regarding the third feature, advanced reduction patterns, recent official releases of StarPU provide the `REDUX` access mode to data. Every worker executing a task on a data provided with this access mode allocates and modifies a private copy of it; as soon as another task is submitted that accesses the same data with

a different access mode, the runtime transparently creates tasks that perform a reduction to merge all the private copies into the original one. This design maximizes parallelism but may lead to an excessive memory consumption and time consuming reductions when the number of workers is high. Additionally, the reduction step is performed in a sequential fashion (i.e., with a flat tree). We implemented the `RANK_REDUX` access mode in StarPU; here only one private copy of the data is created per rank and, therefore, shared by all the workers associated with the rank. In order to compensate for the loss of parallelism, we allow all the tasks mapped on the same rank to commute. In our implementation multiple reduction tree shapes can be used which can be chosen at run time; in all our experiments a binary tree was used.

4.2 Handling the general case

The general matrix-matrix multiplication operation includes multiplications by the α and β scalars and the possibility of transposing the A and B matrices as explained in section 2.2. Additionally, in a completely general setting, the matrices may not be aligned on the ranks grid and possibly they can be distributed over different, non-overlapping, sets of ranks.

4.2.1 Transposition and alignment

In MPI the scope of a collective communication is defined by a (sub)communicator. This does not represent a problem in the case where the A , B and C matrices have a conforming distribution over ranks and neither A nor B must be transposed: all blocks of A (respectively, B) already belong to the row (column) subcommunicators where the broadcasts happen in the stat-C SUMMA algorithm (similar observations can be made for the stat-A and B algorithms). In the opposite case, however, a block of A (respectively, B) might reside on a rank which does not belong in the same row (column) subcommunicator as where the broadcast happens. Handling this case requires additional communications and code, as it is the case, for example, in ScaLAPACK. In our approach, though, handling misaligned distributions and matrix transpositions does not require any special care because dynamic collectives do not rely on the use of subcommunicators but are constructed on the fly for any arbitrary set of ranks.

4.2.2 Scaling

Scaling by the α scalar does not require any special handling. Scaling of the C matrix by the β scalar, instead, might be handled in such a way to achieve better efficiency and parallelism. Let's assume $k = 2$ in the pseudo-code of Figure 6; this implies that each $C_{i,j}$ block is concerned by the two tasks

$$\begin{aligned} \text{task1} : C_{i,j} &= \alpha A_{i,1} B_{1,j} + \beta C_{i,j} \\ \text{task2} : C_{i,j} &= \alpha A_{i,2} B_{1,2} + C_{i,j} \end{aligned}$$

These two tasks, which can be computed using the BLAS GEMM routine, do not commute because of the multiplication by β ; this means that the second task can only be performed after the first even if all the data it needs is already available on the computing rank. In our implementation, the scaling by β is

performed beforehand through dedicated tasks

```

task1 :  $C_{i,j} = \beta C_{i,j}$ 
task2 :  $C_{i,j} = \alpha A_{i,1} B_{1,j} + C_{i,j}$ 
task3 :  $C_{i,j} = \alpha A_{i,2} B_{2,j} + C_{i,j}$ 

```

Here, the first task is relatively cheap and does not require communications and the two other tasks are commutative; this allows for a faster start of computations on all the ranks which might lead to significant performance improvements especially for small size problems.

4.3 Scalability of the STF model

In STF, all the tasks must be created sequentially in order to ensure the dependencies are correctly detected. Although this has some favorable implications (for example it can be used to reliably control the memory consumption of a parallel execution Agullo et al. (2016)), because of this the STF model is commonly regarded as less scalable than other task-based parallel programming models such as PTG, that is, less capable of handling large workloads on large parallel systems. Nevertheless, with some care in the programming and some appropriate techniques, it is possible to overcome this limitation even for very large workloads and computers, as shown by the experimental results in the next section.

4.3.1 DAG pruning

Our implementation employs a DAG pruning technique Agullo et al. (2017a) to prevent each rank from creating all the tasks in the DAG as in the basic concurrent unrolling YarKhan (2012) (see the discussion in section 2.1). Each rank, instead, will create a part of the DAG containing only local tasks (i.e., the tasks it has to execute), remote tasks that use data it owns and remote tasks that produce data needed by local tasks to ensure dependencies are correctly detected at a global scale. In the case of the stat-C SUMMA algorithm, for example this means that a rank has to create a task only if it owns one of the three blocks involved from A , B and C , respectively; in this case the local size of the DAG is $(mnk)/(pq)$ if the A , B and C matrices are aligned over the ranks grid.

4.3.2 Efficient submission of tasks

It must be noted that all the possible nesting orders for the loops of Figure 6 will lead to equivalent DAGs where collective communications and reductions are correctly detected and executed for all the presented variants. Nevertheless, if the nesting order is appropriately chosen, it is possible to ensure some properties of the execution that can be exploited, for example, to control the memory consumption (more on this will be said in section 5.2.3). For the stat-A, stat-B and stat-C variants, the nesting order used in our implementation is, respectively, (n, m, k) , (m, n, k) and (k, m, n) . Although it is still possible to implement the three variants in a single code by using suitable iterators, this would inevitably render the code less readable and, most importantly, the creation of tasks less efficient. Preliminary experimental results revealed that the

StarPU runtime is particularly sensitive to the efficiency of tasks creation and, for this reason, we decided to have three separate codes for the stat-A, stat-B and stat-C algorithms and their 2.5D variants.

5 Experiments

In this section we report experimental results that aim at assessing the effectiveness of the proposed approach. Parallel GEMM is an extremely important numerical kernel and a subject that has been the object of numerous research works. As a result, many different implementations exist in many software packages. Among the most recent efforts, we can cite the work by Herault et al. (2019) where remarkable performance is achieved on large, GPU-based systems, with a task-based parallel approach relying on the PaRSEC runtime system and its PTG programming model. An exhaustive comparison with other software packages is out of the scope of this paper. Rather, the main objective of this experimental analysis is to show that the proposed approach can achieve performance that is on par with reference implementations, despite the use of a very high level parallel programming model and the fact that most of the complex work is delegated to a runtime system. For this reason we have chosen to compare with libraries that either are well known references to which most other works compare, or share some features with our approach; these are

- ScaLAPACK (version 2.0.2): this is the *de facto* standard in parallel dense linear algebra. This library implements the stat-A, B and C 2D SUMMA algorithms using MPI; shared memory parallelism is achieved through the use of multithreaded BLAS routines.
- Slate (commit bb599ae4¹): this recent effort has the objective of producing a ScaLAPACK replacement for modern multicore and accelerator based supercomputers. It implements the stat-A and C 2D SUMMA variants using a hybrid MPI+OpenMP approach and employs a lookahead method to achieve better efficiency by overlapping communications and computations.
- Chameleon (commit 9825fbf1²): this library provides parallel dense and data-sparse linear algebra subroutines using task-based parallelism through different runtimes. It implements the pipelined stat-C 2D SUMMA algorithm using the baseline STF model (see section 3.1); therefore, it does not make use of the extended features described in the present work but, rather, communications are explicitly handled through data copies into temporary matrices. It uses a lookahead mechanism to overlap computations and communications. The runtime chosen for our experiments is StarPU.

5.1 Experimental setup

Our experiments were run on two different partitions of the Joliot-Curie supercomputer of the French Très Grand Centre de Calcul (TGCC) supercomputing center:

¹<https://bitbucket.org/icl/slate/src/bb599ae4/>

²<https://gitlab.inria.fr/solverstack/chameleon/-/tree/9594f9af>

- *Skylake*. A partition of 1,656 nodes equipped with two Intel Skylake 8168 @ 2.7 GHz processors, 24 cores each, and 192 GB of DDR4 memory. This partition uses an Infiniband EDR interconnect through a 4x links pruned fat tree topology.
- *Rome*. A partition of 2,292 nodes equipped with two AMD Rome (Epyc) @ 2.6 GHz, 64 cores each, and 256 GB of DDR4 memory. This partition uses an Infiniband HDR100 interconnect through a dragonfly topology using 5 cells. Each cell is interconnected through a fat tree topology.

On both computers and for all software packages we used the BLAS routines provided by the Intel MKL (version 21.3.0) library and the GNU (version 9.3.0) compilers suite. OpenMPI (version 4.0.5) was used for ScaLAPACK and Slate whereas for `qr_mumps` and Chameleon we use StarPU (commit 1afdaeb09 in the `nmad-coop-mcast` branch³) with the NewMadeleine communication backend (commit fdec689ab⁴) which, in the case of `qr_mumps`, provides support for the dynamic collective communications. For Slate, Chameleon and `qr_mumps` experiments were run with one rank per node using as many threads/workers as the available cores. Although for these three packages slightly better performance could be achieved with multiple ranks per node, this requires a careful placement of processes and threads and a fine tuning of these parameters is out of the scope of this experimental analysis. In the case of ScaLAPACK multiple ranks (MPI processes) per node were used (24 and 64 for Skylake and Rome, respectively) each using two threads as this configuration resulted in the best performance.

In all the packages it is possible to tune the block size to achieve a favorable trade-off between parallelism and efficiency of local computations. We have chosen to run all experiments using multiple block sizes, namely, 256, 512 and 1024, and report the best results. Smaller and larger block sizes were found to be suboptimal on all the tested configuration either because of an excessively small granularity of computations or because of an insufficient amount of parallelism.

For the `qr_mumps` tests, because the GEMM routines are benchmarked alone and not as part of a larger application, we enriched the DAG with artificial tasks to make sure that the dynamic collective communications are correctly detected and the reduction operations entirely executed.

Finally, multiple runs were executed for each experiment, including a warm-up run which is not taken into account in the performance measurement; median performance across these runs is reported in the following sections.

5.2 Experimental results

We have chosen 3 matrix problem types ($m = n = k$, $m = n = 8k$, $m = 8n = k$) and, for each type, three sizes of increasing value. For each type and size we have conducted experiments using 16, 64 and 256 nodes (i.e., up to 12,288 and 32,768 cores on Skylake and Rome, respectively). For the 2D variants, Chameleon, Slate and `qr_mumps` use square grids with $(4 \times 4, 8 \times 8, 16 \times 16)$ ranks, one rank per node each using all the available cores; ScaLAPACK uses grids with $(24 \times 16, 48 \times 32, 96 \times 64)$ and $(32 \times 32, 64 \times 64, 128 \times 128)$ ranks

³<https://gitlab.inria.fr/starpu/starpu/tree/1afdaeb09>

⁴<https://gitlab.inria.fr/pm2/pm2/tree/fdec689ab>

using two cores each on Skylake and Rome, respectively. When using a 2.5D variant in `qr_mumps`, we used rank grids with 4 layers ($4 \times 4 \times 4$, $8 \times 8 \times 4$) on 64 and 256 nodes.

5.2.1 Stationary-C variant

We have evaluated the effectiveness of the stat-C variant on two different problems: the first one uses square matrices ($m = n = k$) and the second one only keeps the C matrix square with A and B being smaller, i.e., ($m = n = 8k$). The first problem is often used in the literature for comparing parallel GEMM algorithms and implementations. In this case, where all matrices are of comparable size, all 2D variants are likely to achieve comparable performance although the stat-C one can be preferred due to its relative simplicity. The stat-C is still the best suited variant for the second problem despite k being small. Although, in this case, a considerable amount of parallelism might still be available when m and n are large, the latency to exchange blocks of A and B is critical to achieve high execution speed because the number of outer products is reduced.

For these problem types, we can observe in figures 7 and 8 that our method is competitive with all the libraries: it obtains the best median across several configurations. When it is not the most efficient method, our method is able to be on par with other ones as it proves strongly scalable. This is likely due to the use of asynchronous collective communications and the ability of the runtime to take advantage of the commutativity of the tasks that contribute to the same C_{ij} block. Notably, our implementation is able to achieve over 500 Tflop/s on the largest size of the $m = n = 8k$ problem on the Rome partition.

For square matrices, we compared the 2D and 2.5D stat-C variants of `qr_mumps`. The 2D variant is better than the 2.5D in all tests except on Rome for the smallest problem and largest grid. In this case the 2.5D variant achieves better performance than the 2D one thanks to its ability to achieve better parallelism without using an excessively small block size. This is not inconsistent with results presented in the literature related to 2.5D matrix multiplication Schatz et al. (2016); Georganas et al. (2012); Demmel et al. (2012) because our approach heavily relies on multithreading for using the cores of each node rather than message passing and employs non-blocking collective communications which were not available in MPI at the time of those works.

5.2.2 A-stationary algorithms

In order to evaluate the effectiveness of the stat-A variant, we have chosen problems where the size of the A matrix is much larger than that of the B and C ones, namely $m = 8n = k$ with $m = \{65,536, 131,072, 262,144\}$. For `qr_mumps` and `Slate` the stat-A and stat-C routines are directly callable and, thus, we include results for both of them; for `ScaLAPACK` it is only possible to call the generic GEMM routine which, internally, chooses the most appropriate variant (which ended up being stat-A for the chosen problem sizes).

For this problem type, we can observe on Figure 9 that our method is significantly better than all the libraries. This is likely due to the use of non-blocking collective communications and the runtime leveraging commutativity of local tasks as well as the reduction patterns.

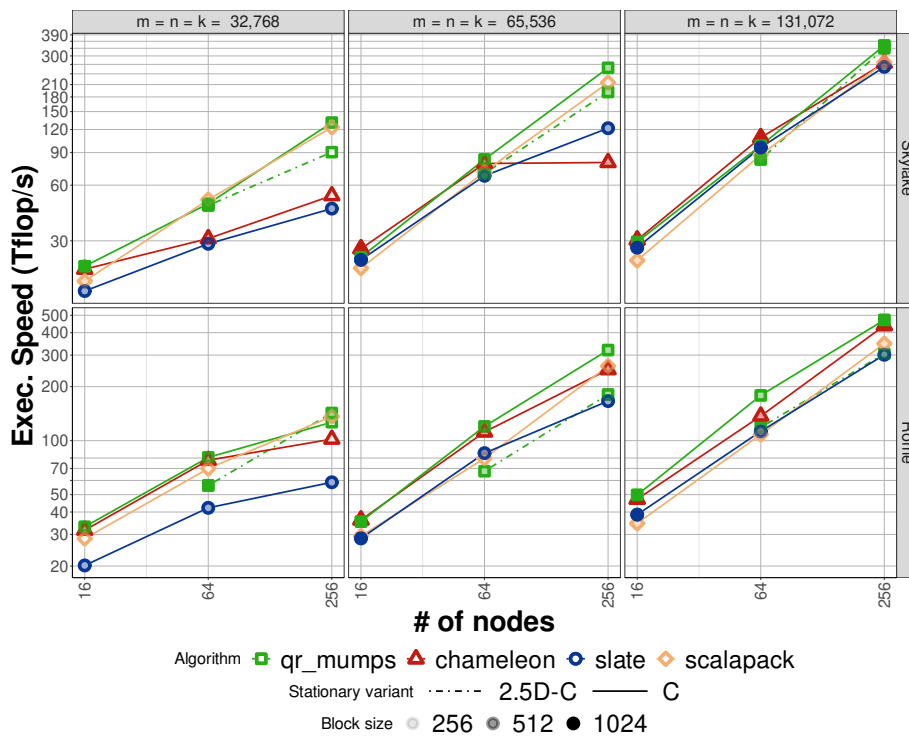


Figure 7: Comparisons of ScaLAPACK, Slate, Chameleon and qr_mumps DGEMM on square matrices ($m = n = k$) and an increasing number of nodes. Markers correspond to the best median across runs using a given blocking for an algorithm.

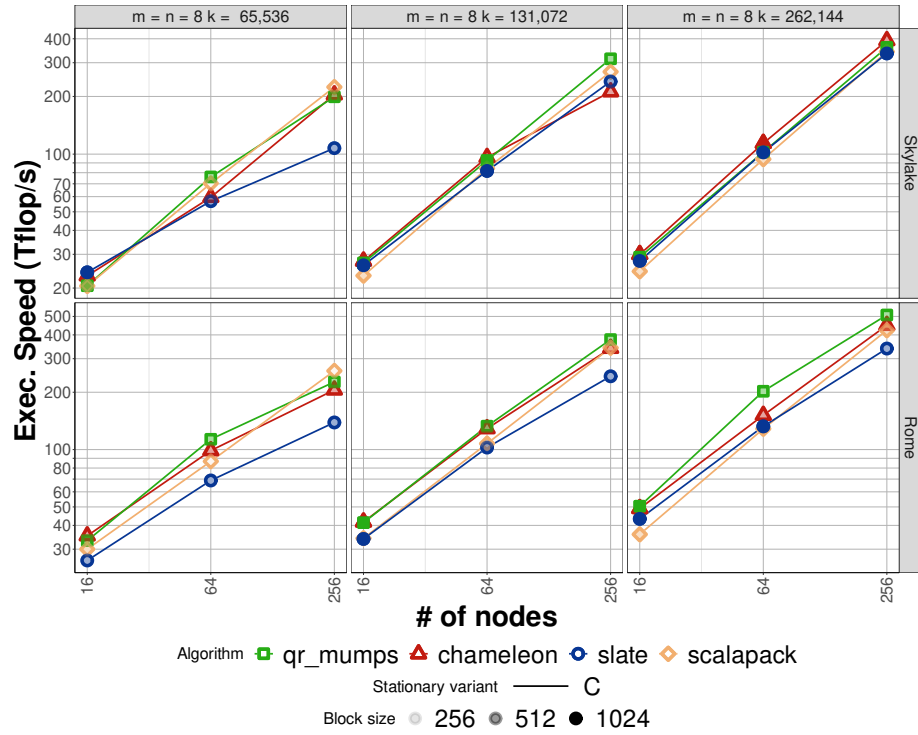


Figure 8: Comparisons of ScaLAPACK, Slate, Chameleon and qr_mumps DGEMM involving a large C matrix ($m = n = 8k$) on an increasing number of nodes. Markers correspond to the best median across runs using a given blocking for an algorithm.

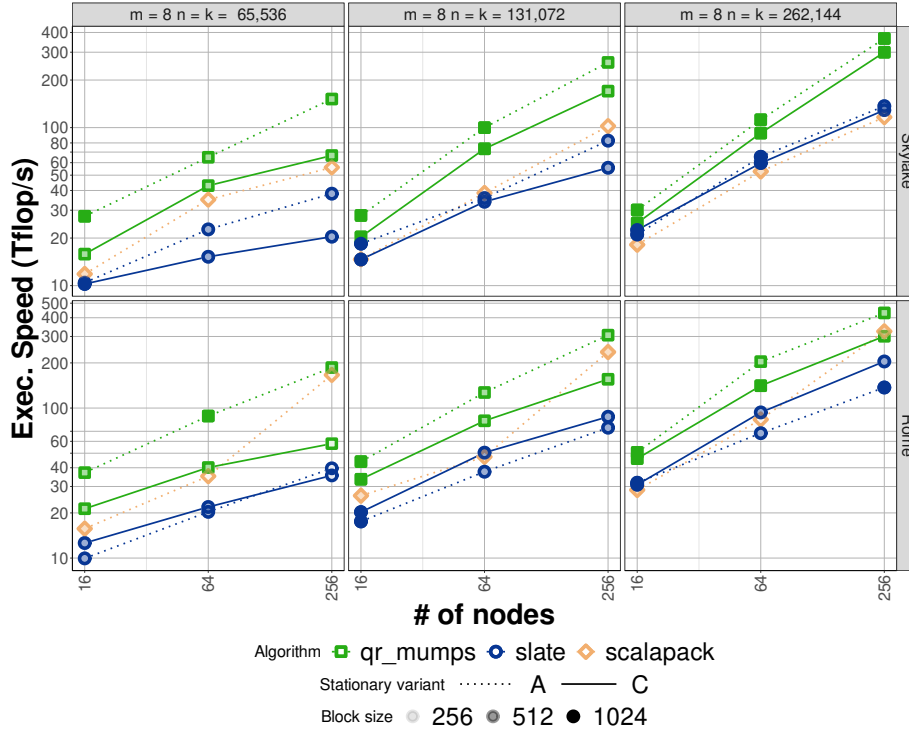


Figure 9: Comparisons of ScaLAPACK, Slate, Chameleon and qr_mumps DGEMM involving a large A matrix ($m = 8n = k$) on an increasing number of nodes. Markers correspond to the best median across runs using a given blocking for an algorithm.

Stat-A variants are significantly better than their stat-C counterparts especially on small size problems and large size grids where the execution time is dominated by communications. As the problem size increases, this difference is less remarkable because there's more opportunity for overlapping communications and computations.

5.2.3 Controlling the memory consumption

Task-based parallel runtimes in general, and StarPU in particular, commonly rely on an eager scheduling: as soon as a task becomes ready, it is scheduled for execution. In the case of StarPU, this also concerns communications because they are fulfilled by dedicated tasks which are automatically and transparently created by the runtime. When the GEMM routine is not evaluated as part of a larger application where the matrices are produced by other operations, all the communication tasks are immediately ready and scheduled for execution and potentially all executed in the very early stages of the matrix product. This has two effects. First it might cause an excessive memory consumption because StarPU must allocate communication buffers for blocks that are received much earlier than when they are actually used. Second, it may reduce performance

because of the high contention that it generates on the network.

StarPU does not currently offer a proper feature to control the execution of communication tasks. Nevertheless, it is possible to control the execution of communications indirectly by retarding the submission of tasks. Note that this relies on a fundamental property of the STF programming model: tasks are created sequentially, that is, in exactly the same order in which the corresponding operations would be executed in a sequential code. Based on this assumption it is possible to use a feature of StarPU that allows one to cap the number of submitted tasks through a sliding window mechanism (this feature is already discussed in related work by Agullo et al. (2017a)). This feature provides two environment variables that can be used to define the maximum and minimum number of submitted tasks: when the maximum is reached, the tasks creation is suspended (the task submission routine becomes blocking) and is resumed when, upon execution of already created tasks, the number falls below the minimum. We used this feature to implement a lookahead mechanism in our implementation. The maximum is set to be the number of tasks in a prescribed number of iterations of the outer loop of the product which, essentially, corresponds to the lookahead depth.

The results obtained with this approach on the stat-C variant are presented in Figure 10 (similar results were obtained on the stat-A one). This figure shows the maximum memory consumption over all ranks, including the initial A , B and C matrices (represented by the gray dotted line), with respect to performance for multiple values of the lookahead depth compared to Slate and Chameleon (for which the default lookahead of 1 was used). The figure clearly shows that when no memory control mechanism is used in `qr_mumps` (which corresponds to an infinite depth lookahead), the memory consumption is excessively high and much higher than the other packages. When a fixed-depth lookahead is used, instead, not only the memory consumption becomes comparable to that of Chameleon and Slate, but performance is improved thanks to a reduced pressure on the communication layer.

This analysis suggests that the memory consumption could be reliably controlled through an analogous feature that allows for capping the maximum size of communication buffers rather than the number of tasks. The sequential tasks submission order will ensure that no deadlocking occurs provided that a sufficient amount of memory can be used Agullo et al. (2016). We reserve the implementation and study of such a feature for future work.

6 Conclusions and future work

The main conclusion of this work is that it is possible to design state-of-the-art matrix multiplication algorithms for distributed memory machines through a very compact sequential-like code. The programmer can be relieved from the burden of writing low-level complex communication schemes as it the case for instance with MPI-based codes such as ScaLAPACK. This can be achieved through the use of advanced features of the STF, task-based, parallel programming model. In this enhanced STF model we have proposed, we have indeed shown that efficient communication patterns can be effectively inferred from an appropriate choice of the mapping and data access modes. The second main conclusion is that, as of 2022, the software ecosystem is mature enough to en-

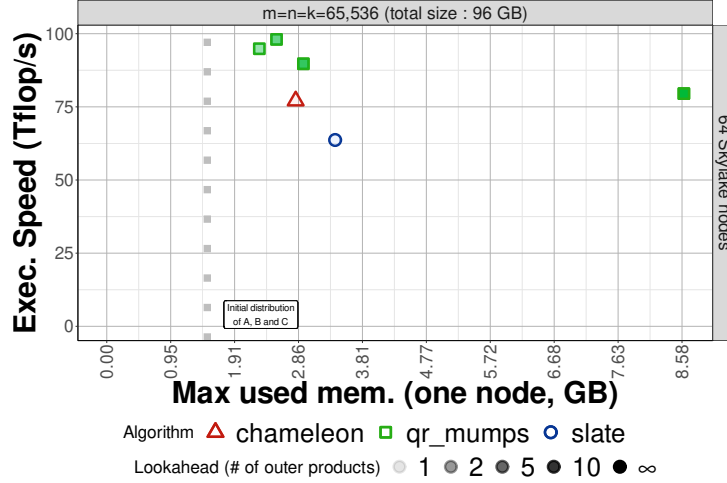


Figure 10: Comparisons of Slate, Chameleon and qr_mumps DGEMM on square matrices of size 65,536 using 64 nodes and a blocking of size 512. qr_mumps uses different tasks window mechanisms.

sure that the resulting code may be competitive against state-of-the-art, finely hand-tuned libraries.

We believe that these conclusions shall motivate the community to further consider task-based programming models, and the STF one in particular, for designing scalable algorithms, while ensuring an enhanced portability and maintenance. On our side, we plan to assess the suitability of the model proposed in this paper to design dense and sparse direct methods.

While the performance results are already remarkable, we believe that the proposed programming model still offers additional opportunities for further optimization. Indeed, while state-of-the-art MPI-based libraries rely on communicators to ensure collective communications, on the contrary, multiple collective communications can (be inferred and) progress concurrently Denis et al. (2020) in our model. In particular, a careful study of their impact on state-of-the-art distributed memory matrix multiplication algorithms remains to be conducted.

While we have shown that the memory footprint can be kept under control without performance penalty (or even improving it), the employed technique – a sliding window mechanism – has required some manual tuning regarding the window size. We plan to make it automatic with respect to a prescribed memory footprint, similarly to Agullo et al. (2016).

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