

ExaFMM: a high-performance fast multipole method library with C++ and Python interfaces

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Summary

ExaFMM is an open-source library for fast multipole algorithms, providing high-performance evaluation of N-body problems in three dimensions, with C++ and Python interfaces. This new implementation is the most recent of many across a decade of work in our research group. Our goal for all these years has been to produce reusable, standard code for this intricate algorithm. The new header-only C/C++ implementation is easier to extend, still competitive with state-of-the-art codes, and includes a pybind11 Python interface (Jakob et al., 2017).

The fast multipole method (FMM) was introduced more than 30 years ago (Greengard & Rokhlin, 1987) as a means of reducing the complexity of N-body problems from $\mathcal{O}(N^2)$ to $\mathcal{O}(N)$ using hierarchical approximations of long-range interactions. Two major variants of hierarchical N-body algorithms exist: treecodes and FMM. (Algebraic analogues also exist, such as H-matrix methods.) Both were originally developed for fast evaluation of the gravitational potential field, but now have found many applications in different fields. For example, the integral formulation of problems modeled by elliptic partial differential equations can be reinterpreted as N-body interactions. In this way, N-body algorithms are applicable to acoustics, electromagenetics, fluid dynamics, and more. The present version of ExaFMM implements the so-called kernel-independent variant of FMM (Ying et al., 2004). It supports computing both the potential and forces of Laplace, low-frequency Helmholtz and modified Helmholtz (a.k.a. Yukawa) kernels. Users can add other non-oscillatory kernels with modest programming effort.

Statement of Need

Over the past three decades, a plethora of fast N-body implementations have emerged. We will mention a few notable ones for context. Bonsai (Bédorf et al., 2012) is a gravitational treecode that runs entirely on GPU hardware. ChaNGa (Jetley et al., 2008) is also a treecode that uses Charm++ to automate dynamic load balancing. ScalFMM (Agullo et al., 2014) implements the black-box FMM, a kernel-independent variant based on interpolation. It comes with an option to use StarPU runtime system to handle heterogeneous task scheduling. TBFMM (Bramas, 2020) is a task-based FMM library that features a generic C++ design to support various types of tree structures and kernels, through heavy use of C++ templates. PVFMM (Malhotra & Biros, 2015) can compute both particle and volume potentials using a kernel-independent FMM, KIFMM (Ying et al., 2004).

The first version of ExaFMM focused on low-accuracy optimizations and implemented a dual-tree traversal (Barba & Yokota, 2012; Yokota, 2013; Yokota & Barba, 2012). It was GPU-enabled using CUDA, parallel with MPI and exploited multithreading using OpenMP. Despite all these efforts, it has remained a challenge in the FMM community to have a well-established



open-source software package, analogous to FFTW for the fast Fourier transform, delivering compelling performance with a standard and easy-to-use interface.

The "alpha" version of ExaFMM is long and complex, and hard to maintain. Its length is partly due to a focus on fast execution, which led to specialized treatment of low-p evaluations and extensive hand-coded optimizations. This emphasis on achieving high performance led to poor reusability and maintainability. The current version uses the kernel-independent variant of the method for higher performance at high accuracy (higher truncation order p). Keeping focus on maintainability, it stays close to standards, with clean function interfaces, shallow inheritance and conservative use of classes. Instead of a fixation with being the fastest of all, we focus on "ballpark" competitive performance. Above all, the Python API and ability to call it from Jupyter notebooks should make this software valuable for many applications.

Features of the software design

exafmm—t is designed to be standard and lean. First, it only uses C++ STL containers and depends on mature math libraries: BLAS, LAPACK and FFTW3. Second, exafmm—t is moderately object-oriented, namely, the usage of encapsulation, inheritance and polymorphism is conservative or even minimal in the code. The core library consists of around 6,000 lines of code, which is an order of magnitude shorter than many other FMM packages.

exafmm—t is concise but highly optimized. To achieve competitive performance, our work combines techniques and optimizations from several past efforts. On top of multi-threading using OpenMP, we further speed up the P2P operator (near-range interactions) using SIMD vectorization with SSE/AVX/AVX-512 compatibility; we apply the cache optimization proposed in PVFMM (Malhotra & Biros, 2015) to improve the performance of M2L operators (far-range interactions). In addition, exafmm—t also allows users to pre-compute and store translation operators, which benefits applications that require iterative FMM evaluations. The single-node performance of exafmm—t is on par with the state-of-the-art packages that we mentioned above. We ran a benchmark that solves a Laplace N-body problem with 1 million randomly distributed particles on a workstation with a 14-core Intel i9-7940X CPU. It took 0.95 and 1.48 seconds to obtain 7 and 10 digits of accuracy on the potential, respectively.

exafmm—t is also relatively easy to extend. Adding a new kernel only requires users to create a derived FMM class and provide the kernel function. Last but not least, it offers high-level Python APIs to support Python applications. Thanks to pybind11, most STL containers can be automatically converted to Python-native data structures. Since Python uses duck typing, we have to expose overloaded functions to different Python objects. To avoid naming collisions and keep a clean interface, we chose to create a Python module for each kernel under exafmm—t's Python package, instead of adding suffixes to function and class names to identify types.

Application

We have recently integrated exafmm-t with Bempp-cl, an open-source boundary element method (BEM) package in Python (Betcke & Scroggs, 2021), whose predecessor, BEM++ (Śmigaj et al., 2015), has enabled many acoustic and electromagnetic applications. In BEM applications, computations are dominated by the dense matrix-vector multiplication (matvec) in each iteration. exafmm-t reduces both time and memory cost of mat-vec to a linear complexity, thus makes Bempp-cl feasible to solve large-scale problems. In an upcoming paper, we demonstrate the capabilities and performance of Bempp-Exafmm on biomolecular electrostatics simulations, including solving problems at the scale of a virus (Wang et al., 2021). The showcase calculation in that paper (submitted) obtains the surface electrostatic



potential of a Zika virus, modeled with 1.6 million atoms, 10 million boundary elements (30M points), at a runtime of 1.5 hours on 1 CPU node.

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