### Making PyExaFMM Performant

Srinath Kailasa

Department of Mathematics University College London

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Fast Python
Is Numba Up To It?
Language Bottlenecks for Speed

Software Design Separating Concerns Designing Around Data

Accelerating M2L Randomised SVD Recap Compressing the M2L Gram Matrix Implicitly Unexpected Problems



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### Is Numba Up To It?

Problem: Laplace Kernel applied to interactions between 20,000 particles to find Gram Matrix

■ Numba: ~ 296ms

■ Rust: ~ 155ms

So at first glance, yes - Numba appears to be useful.



## Is Numba Up To It?

Important caveats for benchmark,

- Numba targeting all CPUs via prange operator
- Data is already arranged 'nicely'



### Language Bottlenecks For Speed

- Native Python objects are slow for our purposes, with important implications for data access
- Python's numeric libraries (Numpy/Numba) etc haven't focussed on distributing data as much as computation. There are moves in this direction (Legate)



### Language Bottlenecks For Speed

#### Problems faced by PyExaFMM:

- How do we store tree efficiently if we want to avoid native PyObjects?
- How do we access precomputed operators and coordinate data without dictionaries, or access to pointers?
- How can we accelerate computations on the tree structure?



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### Separating Concerns

- Separate trees into their own module as a dependency -AdaptOctree
- Separate precomputations into script, with CLI for interactive programming
- Minimal API, consists of a single object containing FMM loop and required data
- Separate compute 'backend' (e.g. choice between Numba, Cython etc) and 'frontend' consisting of the API



## Separating Concerns

#### AdaptOctree is:

- Accelerated with Numba
- Tree construction is multithreaded, balancing and interaction list computation are single-threaded
- Benchmark: 1e6 particles distributed randomly, build time 244  $\pm$  6 ms, balance time 97.5  $\pm$  3 ms, interaction lists (slow) 5.81  $\pm$  0.07s



### Separating Concerns

#### We use a single HDF5 file to store:

- Precomputed FMM operators
- Precomputed tree stored as a single vector, and tree parameters (physical center, radius etc)
- Precomputed index pointers tying together tree index and particle indices
- Precomputed interaction lists
- Benchmark: 1e6 randomly distributed particles, order 5 multipole expansion, order 6 local expansion, max 100 particles per node, target rank 1 in SVD, takes 9s.



### Designing Around Data

Avoiding array creation/destruction leads to use of index pointers (linking indices between two arrays) e.g. between array of tree nodes, and their corresponding particles.

- Prioritise simple data structures, arrays
- Avoid sets and dicts if possible, these are partially supported by Numba - but add complications in terms of allowed return types, instatiation time
- All computational methods stripped down to bare loops to help Numba, and operator in place on data - looks UnPythonic and more C-like.



### Designing Around Data

#### Numba Gotchas

- Numba threading on the face of it simple: range → prange for a given for loop, picks up backend (TBB, OMP etc).
- Can lead to oversubscription of threads (silently) killing performance
- JIT'ing doesn't change allocation cost, so have to be careful to operate in place, this then feels like you are programming to an (unspecified) framework!
- Sometimes multithreading doesn't work as expected, especially when performing recuctions - rewriting to ensure that all operations are in place may help (i.e. avoiding allocations within prange loop).
- Numba works 'too well' fails silently, and always produces output!



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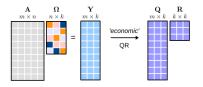
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### Accelerating M2L

Offers a case study in utility of Numba for scientific computing, but also the pitfalls.



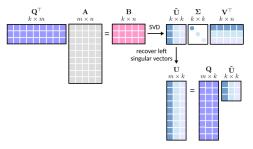
# Randomised SVD Recap



Adapted from [1]



# Randomised SVD Recap



Adapted from [1]



# Compressing the M2L Gram Matrix Implicitly

- 'A' corresponds to Gram Matrix for unique M2L interactions for a given target node and up to 316 source nodes at each level
- We just store coordinates, and compute matrix elements implicitly, saving memory
- Use Cupy and Numba interoprability to transfer data to device, and compute SVD/QR algorithms - easy!
- Only compute unique interactions for a given level, referencing these with a 'transfer vector' [2]



### **Unexpected Problems**

- Need to compute a hash corresponding to a given transfer vector on the fly, can't lookup using coordinate vector as these can't be used as keys in HDF5 (or PyDict).
- How does one efficiently hash a vector? Does Numba support hashes? Is it fast?
- Inability to treat functions as first class objects (for interoprability with Cupy functions) leads to massive code duplication for different kernels.
- At this point, the code is already fairly UnPythonic (no objects for encapsulating behaviour, no enforced interfaces) - why are we still using Python?



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### References I

- N. Benjamin Erichson, Sergey Voronin, Steven L. Brunton, and J. Nathan Kutz. Randomized matrix decompositions using r. Journal of Statistical Software, 89(11), 2019.
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