

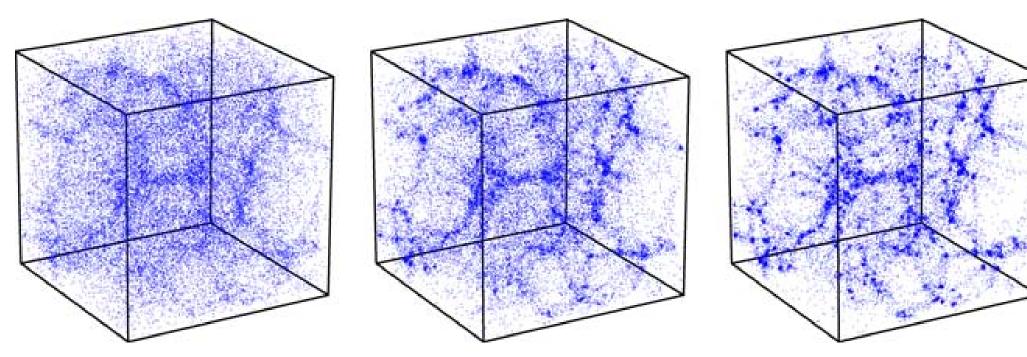
SAKURA: RECURSIVE CONSTRUCTION AND TRAVERSAL OF FMM SKELETON

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Motivation

- Tree building and graph traversal for interaction challenge parallel architectures, slowing down parallel FMM execution.
- Updating the tree and interaction lists is critical in time-dependent computations.
- Infrequent updates lead to loss of accuracy.

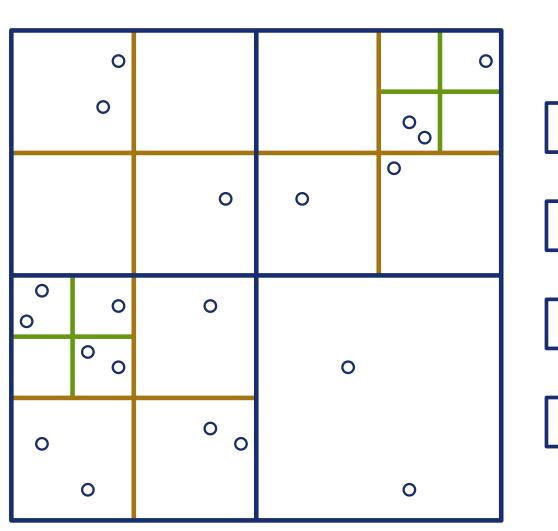


Evolution of galaxy distribution in Astrophysics simulation; figure taken from [2].

→ Sakura takes a fully recursive approach for processing the FMM skeleton (tree partitions + interaction lists), ensuring the efficient storage and traversal of the data structure.

Adaptive Tree Building

- Fast spatial partitioning of source and target sets via adaptive geometric binning.
- Hierarchically local mapping of particle records to memory.



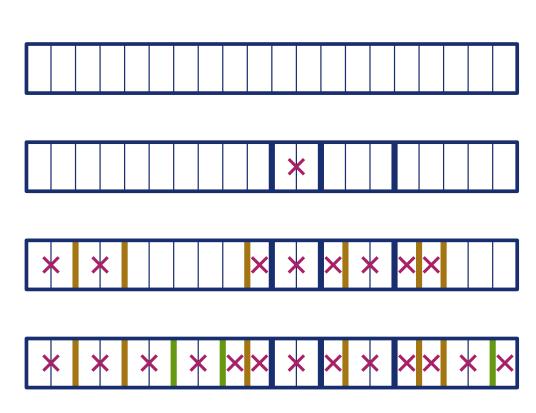


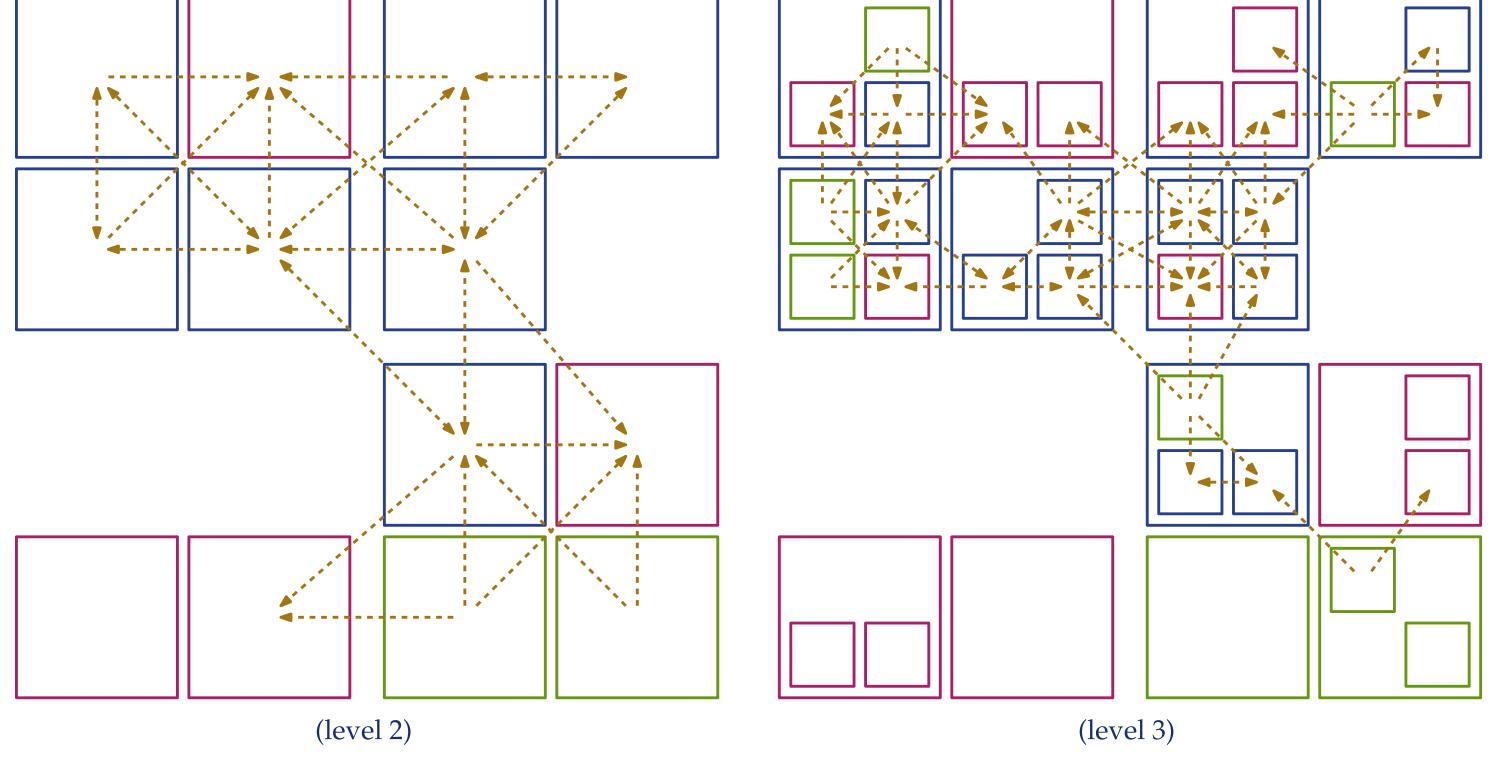
Illustration of geometric partitioning and corresponding memory rearrangement of a 2D particle-set. The particle records are recursively binned in memory, following the spatial partition hierarchy. The recursion terminates for partitions/bins whose population is below a threshold, while empty partitions/bins are discarded.

References

- [1] L. F. Greengard and V. Rokhlin. A fast algorithm for particle simulations. *Journal of Computational Physics*, 73(2):325–348, 1987.
- [2] E. Platen, R. van de Weygaert, and B. J. T. Jones. A cosmic watershed: the WVF void detection technique. *Monthly Notices of the Royal Astronomical Society*, 380(2):551–570, Sept. 2007.
- [3] H. C. Plummer. On the problem of distribution in globular star clusters. *Monthly Notices of the Royal Astronomical Society*, 71(1):460–470, 1911.
- [4] R. Yokota. An FMM based on dual tree traversal for many-core architectures. *Journal of Algorithms and Computational Technology*, 7(3):301–324, 2013.

FMM Interaction Lists

- ► Explicit formation of interaction lists
- 1. Fast counting of interaction links (dual-tree traversal; pass 1)
- 2. Tight memory allocation for all links with a single OS call
- 3. Instantiation of interaction lists (dual-tree traversal and memory stuffing; pass 2)
- Minimal number of node-pair visits during the counting and instantiation passes.

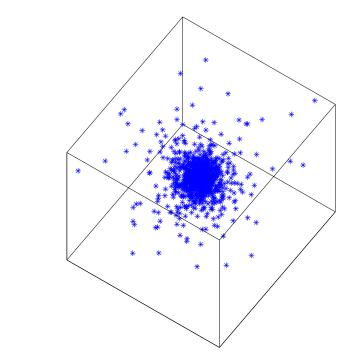


Near-neighbor links in two resolution levels of a sample dual tree (source + target trees). **Green** boxes contain target particles, **magenta** boxes contain source particles, while **blue** boxes contain both target and source particles. **Orange** arrows indicate target-to-source links. Interaction lists (not shown) are induced by multi-level neighboring relationships; hence, only near-neighboring pairs of nodes need to be visited for interaction list formation.

- Original, compressed, and cross-level interaction lists are supported.
- ► Implicit-list interactions
 - The fast traversal algorithm is also applicable to implicit interaction list approaches.

Experimental Set-up

- Particle distributions for simulations:



• Low- and high-accuracy simulations:

- expansion order p = 4 and p = 9

uniform distribution on sphere octant surface

Plummer distribution [3]

• Simulation platform:

| | | | | | | | Cache levels | | | |
|---------------------|-------------|-------|------|-----------|------|-------|--------------|--|--|--|
| CPU Type | CPUs | Cores | Thr. | CPU clock | L1 | L2 | L3 | | | |
| Xeon E5-2650 | 2 | 8 | 16 | 2.4GHz | 32KB | 256KB | 30MB | | | |

Experiments

Sakura performance (comparison with the widely-used ExaFMM package [4]):

ratio of combinatorial construction to total FMM execution time (t_c/t_c+t_n)

| | | <i>p</i> = | = 4 | | p=9 | | | | |
|-----|--------|------------|-------|-----|--------|-----|-------|-----|--|
| | Octant | | Plum. | | Octant | | Plum. | | |
| N | skr | exa | skr | exa | skr | exa | skr | exa | |
| 10 | 20 | 48 | 22 | 38 | 6 | 16 | 4 | 11 | |
| 20 | 18 | 50 | 22 | 43 | 5 | 18 | 4 | 12 | |
| 40 | 18 | 53 | 22 | 45 | 5 | 21 | 4 | 13 | |
| 80 | 18 | 50 | 20 | 45 | 5 | 18 | 3 | 14 | |
| 160 | 18 | 54 | 21 | 50 | 5 | 19 | 3 | 15 | |
| 320 | 19 | 54 | 21 | 52 | 5 | 19 | 3 | 31 | |
| | | | | | | | | | |

total FMM execution time in seconds $(t_c + t_n)$

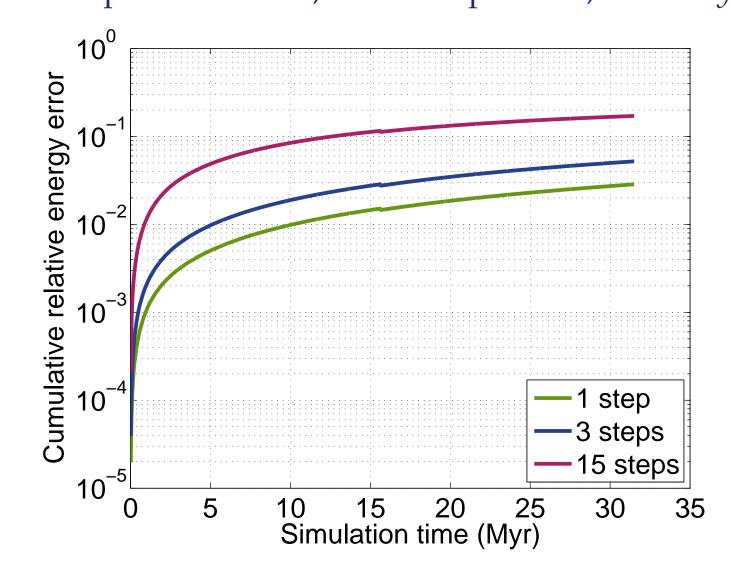
| | p=4 | | | | p=9 | | | |
|-----|--------|-----|-------|-----|--------|-----|-------|-----|
| | Octant | | Plum. | | Octant | | Plum. | |
| N | skr | exa | skr | exa | skr | exa | skr | exa |
| 10 | 2 | 3 | 5 | 6 | 5 | 7 | 10 | 14 |
| 20 | 4 | 6 | 10 | 13 | 11 | 13 | 20 | 27 |
| 40 | 8 | 13 | 19 | 25 | 21 | 26 | 40 | 52 |
| 80 | 14 | 23 | 30 | 43 | 40 | 51 | 75 | 101 |
| 160 | 28 | 49 | 58 | 90 | 82 | 102 | 151 | 201 |
| 320 | 49 | 103 | 107 | 189 | 159 | 214 | 299 | 502 |

N: # of particles (M); **skr**: Sakura execution; **exa**: ExaFMM execution t_c : time for combinatorial construction; t_n : time for numerical evaluation of FMM

- FMM skeleton construction cost ratio reduceed by $2.5–3.5 \times$
- total FMM time reduced up to $2\times$ for larger data-sets

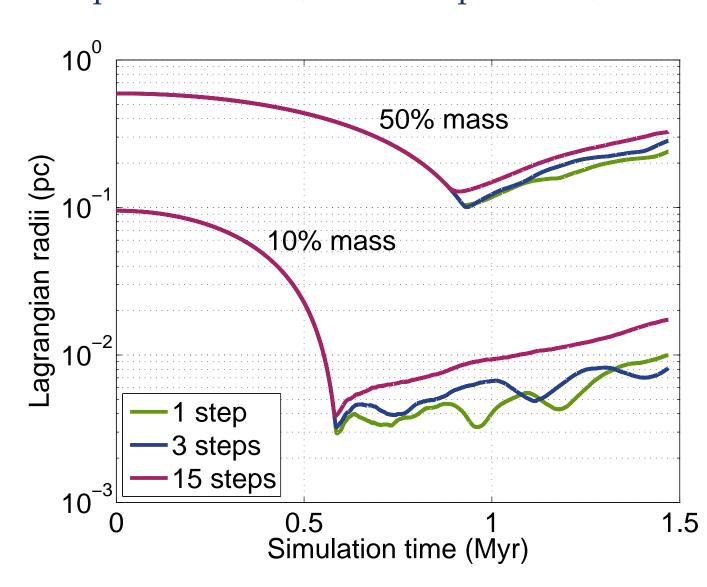
Plummer sphere evolution

- cumulative relative energy error: $\Delta E_t = \frac{E_t E_0}{E_0}$
- E_t : total (kinetic + potential) energy at time t
- # particles: 10M; # time steps: 3000; $\Delta t = 1kyr$



Plummer sphere core collapse

- Lagrangian radii: minimal radii which enclose a certain portion of the total system mass
- # particles: 10M; # time steps: 25000; $\Delta t = 100yr$



- error increases with less frequent FMM skeleton updates
- execution time: 3-step updates with Sakura \approx 15-step updates with ExaFMM

Strong thread scalability

- # particles: 160M
- p = 4
- FMM skeleton construction scales just as well as the FMM computations
- both scale almost ideally up to 16 threads (# of cores)

