



VELOCIraptor: A MPI+OpenMP data analysis tool for cosmological simulations

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Outline

- Introduction to Numerical Cosmology & Structure Finding
- VELOCIraptor
- MPI-raptors
- OpenMP-raptors



Cosmological Simulations

- Common approach to understanding the largescale universe and the objects that form within it, like galaxies, is to run high-resolution, largevolume cosmological simulations.
 - Hundreds of billions of elements, 10s to 100 Setonix nodes. Necessitates MPI
- Critical step in extracting information from sophisticated, multi-billion particle simulations is the non-trivial identification of cosmic structures, like dark matter halos and synthetic galaxies.
 - Need to load data rapidly to quickly analyse 100s of snapshots. Needs 10s to 100 Setonix nodes. Necessitates MPI.





VELOCIraptor

- A phase-space [multi-dimensional] (sub)halo finder capable of identifying dark matter halos and galaxies
- Open source <u>https://github.com/pelahi/VELOCIraptor-STF.git</u> & documentation <u>http://velociraptor-stf.readthedocs.io/en/latest</u>
- MPI + OpenMP, C++17, CMAKE build system,, Parallel IO via HDF5 (ADIOS in works)



Halo and Substructure Identification from Simulation



VELOCIraptor

Configuration

• Read configuration & init deployment setup (MPI, OpenMP)

Input

- Read input file domain setup MPI decomposition
- Fully read data in parallel
- Can have different # of tasks reading input file(s)

Field Search

- Each MPI process searches for field halos (parent clusters) locally and across neighbouring MPI domains.
- MPI process can locally search with OpenMP
- Localise all field halos to single MPI domain, adjust MPI decomposition

Substructure

- MPI localised field halos searched for (sub)substructures in all (sub)structures.
- OpenMP enabled

Analysis

- Calculate properties for all objects.
- OpenMP enabled

Output

- Output fields containing object properties and particle associations in parallel.
- Can gather MPI process to write to single file or many.

MPI Input & Decomposition

- Input can single file or split between many files
- Number of MPI no relation to number of input files
- Parallel read where number of read tasks need not be MPI_COMM_WORLD.
 - Read ranks are spread across, maximising node spread of reading ranks.
 - Can have tasks just waiting for input.
- Communication dominated by asynchronous point-to-point
- To ensure MPI not flooded, communication sent in chunks.
- Input data highly clustered, load balancing is critical. Not just about memory want to reducing amount of point-to-point communication.
- Use a mesh placed across the input domain and then assigns cells in the mesh to a given MPI rank using a space-filling Z-curve.



MPI Search & Communication

- MPI will local search is particles for those that meet linking criteria. Iterate across MPI domains till no new links are found.
- Now localise clusters to MPI domain with fewest particles.
 - Each rank does not know a priori what other ranks contain desired particles and what other ranks require particles from it.
 - MPI rank able to determine if local particle's search window intersects the volume of a cell that belongs to other MPI ranks using a fast binary tree



Example particle distribution with single cluster (left) Resulting initial 3 Rank MPI decomposition (middle): Volume fraction = [0.56, 0.1, 04], Particle fraction = [0.36, 0.35, 0.28]. Particles coloured by rank. Final MPI decomposition after linking where single object now on Rank = 2. Mesh now has 19 cells shared between at least 2 ranks.

- Communication dominated by asynchronous pointto-point.
- To reduce wait times and congestion use communication work queue containing all communicating pairs

```
//NProcs = total number of Process in MPI_COMM_WORLD
std::vector<tuple<int, int>> MPIGenerateCommPairs(unsigned long long
    *send info)
ſ
   std::vector<std::tuple<int, int>> commpair;
   for(auto task1 = 0; task1 < NProcs; task1++)</pre>
       for(auto task2 = task1+1; task2 < NProcs; task2++)</pre>
           if (send_info[task1 * NProcs + task2] == 0 && send_info[task2
               * NProcs + task1] == 0) continue;
           commpair.push_back(make_tuple(task1, task2));
       3
   // ensure rank = 0 doesn't dominate communication pairs by
        randomizing
   unsigned seed = 4322;
   std::shuffle(commpair.begin(), commpair.end(),
        std::default_random_engine(seed));
   return commpair;
```

- Use HDF5 library to produce parallel IO file(s).
- Includes a test to check HDF5 Parallel IO performance. Expanding this to determine performance
- Output is aggregated using MPI communicators, one for each file.
 - MPI ranks writing to same file are close in rank, thereby reducing internode communication when coordinating file writes.

Reading: Example of ranks reading single input file. Here, nodes have 9 cores, 3 nodes, and total number of MPI ranks reading is 5

Writing: Example MPI ranks colour coded according to file they write to. Here, nodes have 9 cores, 3 nodes, and total number of MPI ranks per file is maximum of 4.





MPI Future Work



- Compute and IO decomposition does make some assumptions about process thread affinity and topology but this need not be correct.
- Further progress to generate rank ordering based on core affinity, numa regions and node placement (using numactl, lscpu, hostname, etc to gather initial setup)

Summary



- MPI + OpenMP, C++17, CMAKE build system, Parallel IO via HDF5 capable of running on thousands of cores.
- Adding better OpenMP task parallels, GPU parallelism, MPI decomposition tied to numa topology, ADIOS IO for improved parallel IO and streaming.



Halo and Substructure Identification from Simulation

Questions?





