

A Classical MD Primer

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Classical Molecular Dynamics

- Multiple ECP Projects do this: EXAALT, COPA
 - Widely used codes: LAMMPS, GROMACS, AMBER, NAMD
- Bio-simulations:
 - relatively restricted set of physics models
 - Short Range LennardJones+Coulomb
 - Long Range Coulomb
 - Bonded interactions (bonds/angles)
 - Implicit solvent
 - Open Codes: LAMMPS, GROMACS, AMBER, NAMD
 - Most MD simulations by cycles are Bio-simulations
 - Important in the pharmaceutical industry
- Material simulations
 - Hundreds of physics models
 - LAMMPS by far the dominant code, but many physics-specific codes exist

Problem Sizes

- Total number of atoms: 100k – 10B
 - Sometimes parameter study with thousands of such systems
- Neighbors per atom: ~10-100 without Coulomb; <1000 with Coulomb
- Memory: 0.1-1kB per particle
 - 50-90% in Neighbor Lists
- Important: total number of timesteps high (10^8 - 10^{10} ...)
 - Targeting 100-10000 timesteps per second
 - Time per Kernel (parallel loop) ~ 0.1-10ms
- Exception: Some folks (LANL, LLNL) do giant short time sims
 - Think $10^{11/12}$ atoms for $<1^6$ timesteps -> memory constraint

The Kernels

- **Long Range Coulomb** (not used or ~50% of time):
 - Discretize Charge Distribution: Scatter Add onto a lattice
 - **Global 3D FFT**: this is the scaling bottleneck in most cases
- **Bonded interactions** (~5% - 20% of time):

```
for(int a=0; a<num_angles; a++) { // Loop over all angles
    int j1 = angles(a,0); // What are the atoms in this angle
    int j2 = angles(a,1);
    int j3 = angles(a,3);
    double3 f1,f2,f3;
    // Compute the actual forces on j1,j2,j3
    // Involves many gathers (positions, force parameters)
    compute_angle(j1,j2,j3,f1,f2,f3);
    // Scatter Add forces back to the three atoms of the angle
    force(j1) += f1;
    force(j2) += f3;
    force(j3) += f3;
}
```

Kernels 2

- **Short Range Force Kernel:** usually 50%-95% of total time

```
for( int i=0; i<natoms; i++) { // Loop over num_atoms 10k-10B
    double3 pos_i = pos(i); // Load positions of i
    int type_i = types(i); // Load type of i

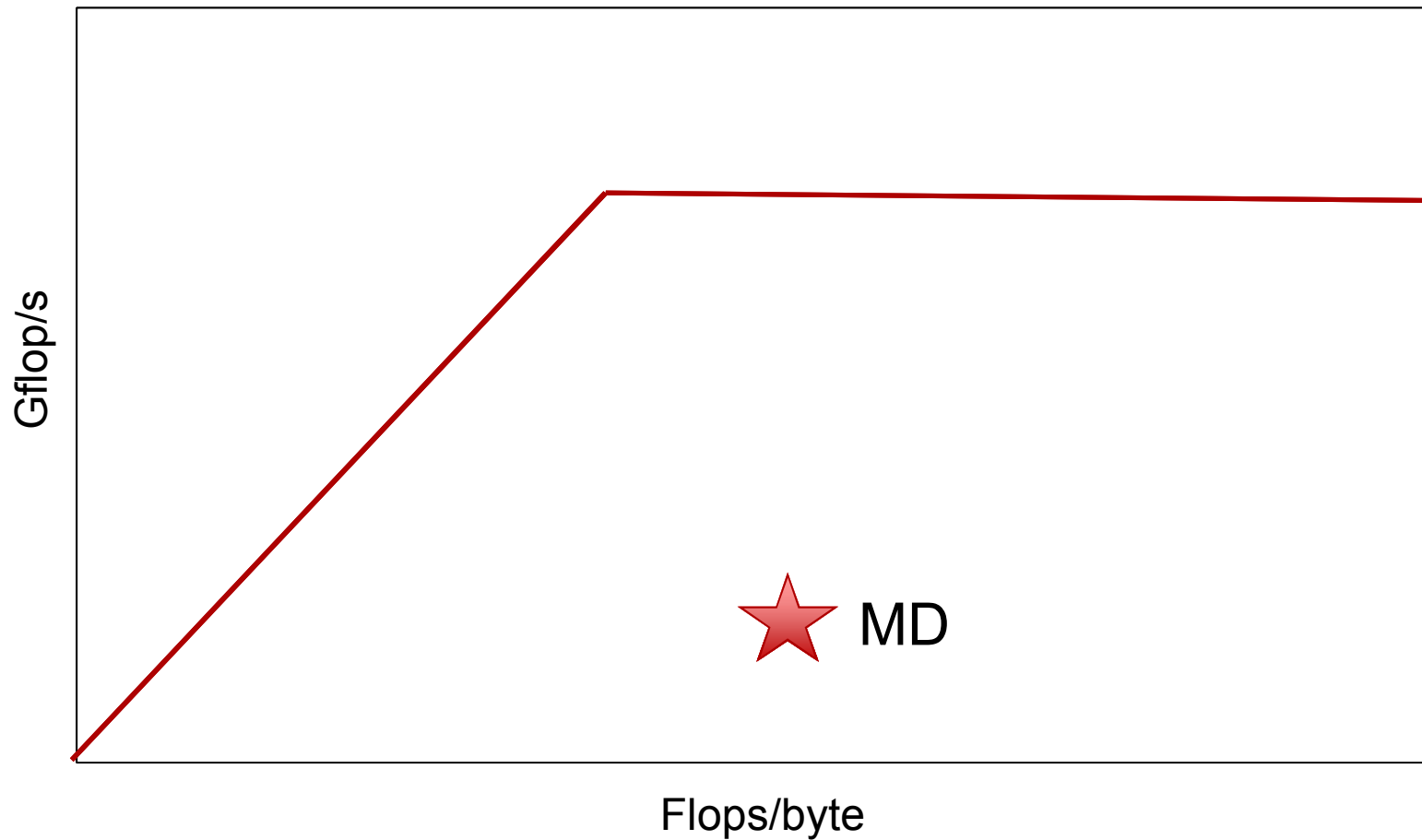
    for(int j=0; j<num_neighs(i); j++) { // Loop over neighbors 10-1000
        int j = neighs(i,j); // This is coming from HBM
        double3 pos_j = pos(j); // Gather from cache (L1) (spatial locality)
        int type_ij = type_i * ntypes + type(j);
        // type_ij specific gathers:
        double3 f_ij;
        if(rsq < cutoff[type_ij]) {
            // Compute actual force
            // 10 – 200 flops, 2-20 more type_ij loads
        }
        force_i += f_ij;
        force(j) -= f_ij;
    }
    force(i) += f_ij;
}
```

- Manybody potentials: loop over neighbors of neighbors

Stuff which is very similar

- Astro Physics:
 - like the large scale MD simulations
 - many atoms,
 - few timesteps
 - Specialised long range solvers for gravity
- Finite Element Codes
 - Explicit codes very similar: loop over elements compute something based on neighbors
 - Implicit codes reflect this in matrix assembly phase
 - Physics often pretty complex

Roofline Model





kokkos is a C++ framework for parallel programming

<http://www.github.com/kokkos>