Continued Effort on Porting LAMMPS to Aurora

The Science

- Molecular Dynamics (MD) is awesome
- LAMMPS is extremely general and extensible software for largescale molecular simulations (MD and beyond).
- Forcefields describe how particles interact:
 - Bonded interactions, dispersion, electrostatics (short + long ranged), granular, nonreactive vs. reactive
- LAMMPS on Aurora supported via two routes
 - Kokkos package: goal of being gpu-resident
 - Many capabilities still need to be ported
 - GPU package: only pair calculation off-loaded
 - Relies heavily on CPU parallelization

The Challenge

- A large number of different forcefield models and capabilities need to be evaluated for functionality, correctness, and performance.
- Aurora will be available to users soon and need to ensure key capabilities in LAMMPS are supported for users to achieve their science goals
- Support for long-range KSPACE electrostatics with Kokkos/SYCL was still missing.



Solvated proton in water and rhodopsin protein embedded in membrane (Image credit: Christopher Knight (ANL)).

Relevant publication(s) on science background: <u>A.P. Thompson</u>, et.al., *Comput. Physics. Commun.*, **271**, 108171 (2022).

https://www.lammps.org/ https://github.com/lammps/lammps

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The Accomplishment

- Added support for offloading 1D FFTs via oneMKL when using the SYCL backend of Kokkos, thus enabling support for multiple capabilities in LAMMPS
 - Long-range electrostatic solvers
 - 17 additional pair potentials to utilize GPUs on Aurora
 - Initial support for other packages: ELECTRODE, DIELECTRIC, and AMOEBA (more testing needed)
- Competitive performance achieved with Kokkos??
 - Issue with NeighborBuild needs resolution
 - Waiting on new Intel Compiler drop to evaluate how broadly beneficial some optimizations are for indexing into multi-dimensional Kokkos views.
- CMake build issues for oneAPI and Kokkos/SYCL were resolved

The Impact

- Users getting access to Aurora in CY25 more likely to have success running their LAMMPS simulations
 - This directly supports one 2024 Aurora INCITE allocation.
- The Makefiles and CMake scripts provided also helps users with building LAMMPS on local Intel GPU resources.
- This also potentially helps with ongoing INCITE review and understanding the readiness of LAMMPS for specific workloads.



LAMMPS-NB Z NeighborBuild

Relative runtimes of the rhodopsin benchmark on an NVIDIA A100 and Intel PVC GPU (1- and 2-tiles). Runtimes are scaled relative to A100 runtime with the NeighborBuild (NB) time separated. (Image credit: Christopher Knight (ANL))

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Publication/PR for this work: 9/3/24 https://github.com/lammps/lammps/pull/4313

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LAMMPS Rhodopsin Benchmark

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