

Streamlining Machine Learning for Molecular Dynamics by Interfacing Python with LAMMPS

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Motivation

- LAMMPS is a gold-standard molecular dynamics simulation program written in C++
- The ML-IAP (Machine-Learning Interatomic Potential) package provides a simplified interface for connecting external C++ codes into LAMMPS
- However, many advanced ML packages are written in Python
- While LAMMPS allows users to couple Python models with their interatomic potentials through its Python package, this is meant for prototype codes and only allows for simplified potentials
- LAMMPS lacks a direct interface to arbitrary Python models**

The ML-IAP Unified Interface

- The new “ML-IAP Unified” Interface unifies both model and descriptor functionalities into a single streamlined Python interface
 - Descriptors give features but neural net finds own
- This allows users to construct elegant Python models that seamlessly interface with LAMMPS using familiar methods**
- Enables computational scientists to more quickly and easily construct Python models using GPU-accelerated machine-learning interatomic potentials for use with LAMMPS
- Allows for easier usage of PyTorch-based machine learning models such as HIP-NN with LAMMPS**

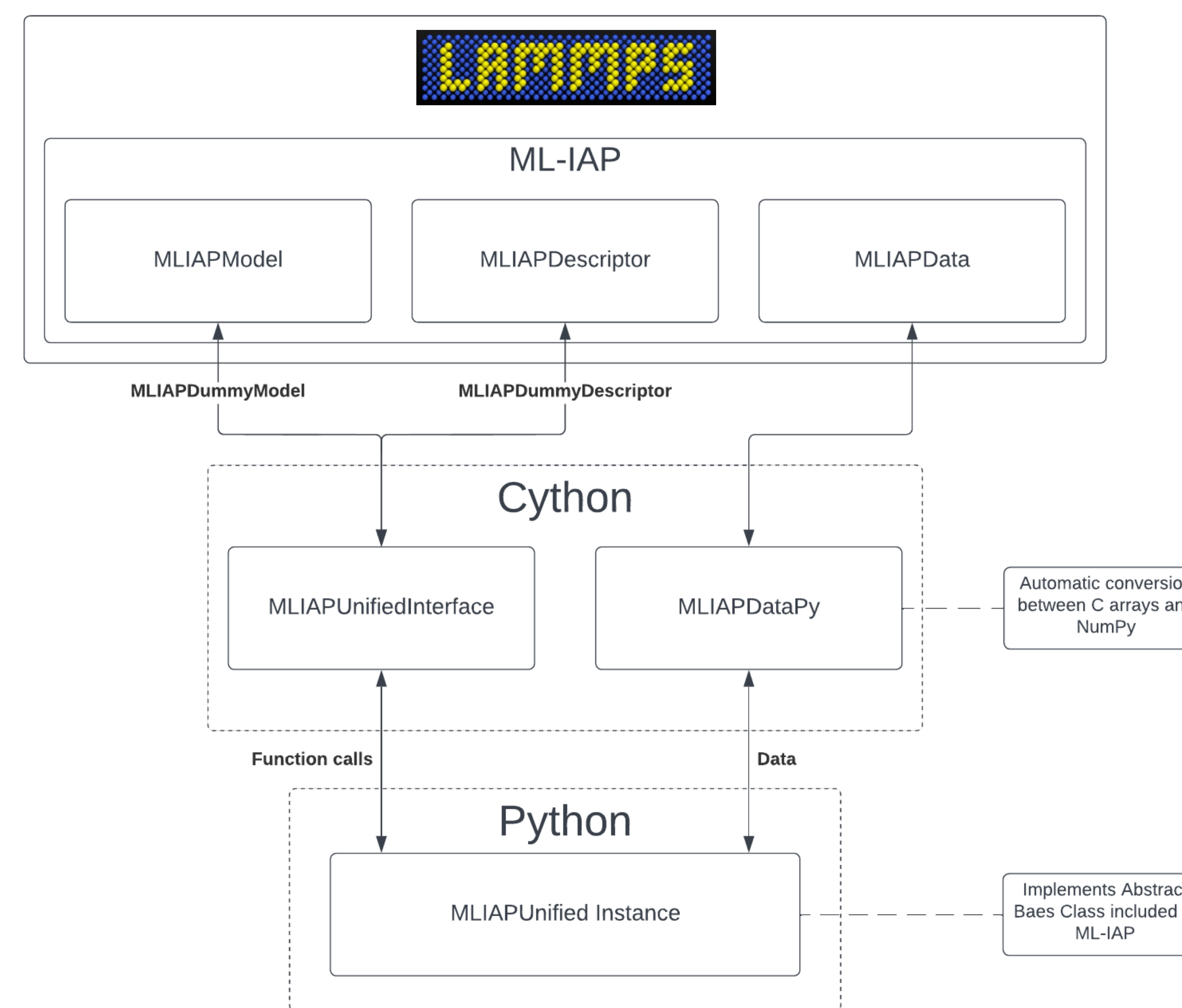
Background: HIP-NN

- Convolutional neural network designed for interatomic potentials
- Highly-tunable parameters, performance
- Implemented in Python using PyTorch

Challenges Encountered

- Creating direct, two-way Cython interface
- Implementing automatic data conversion between C++ and NumPy
- Supporting modified atom neighbor lists for models that require them
- Supporting multiple atom types (see InP Example)
- Identifying and addressing Python bottlenecks

Unified Interface Architecture



InP Example: Unified HIP-NN vs SNAP

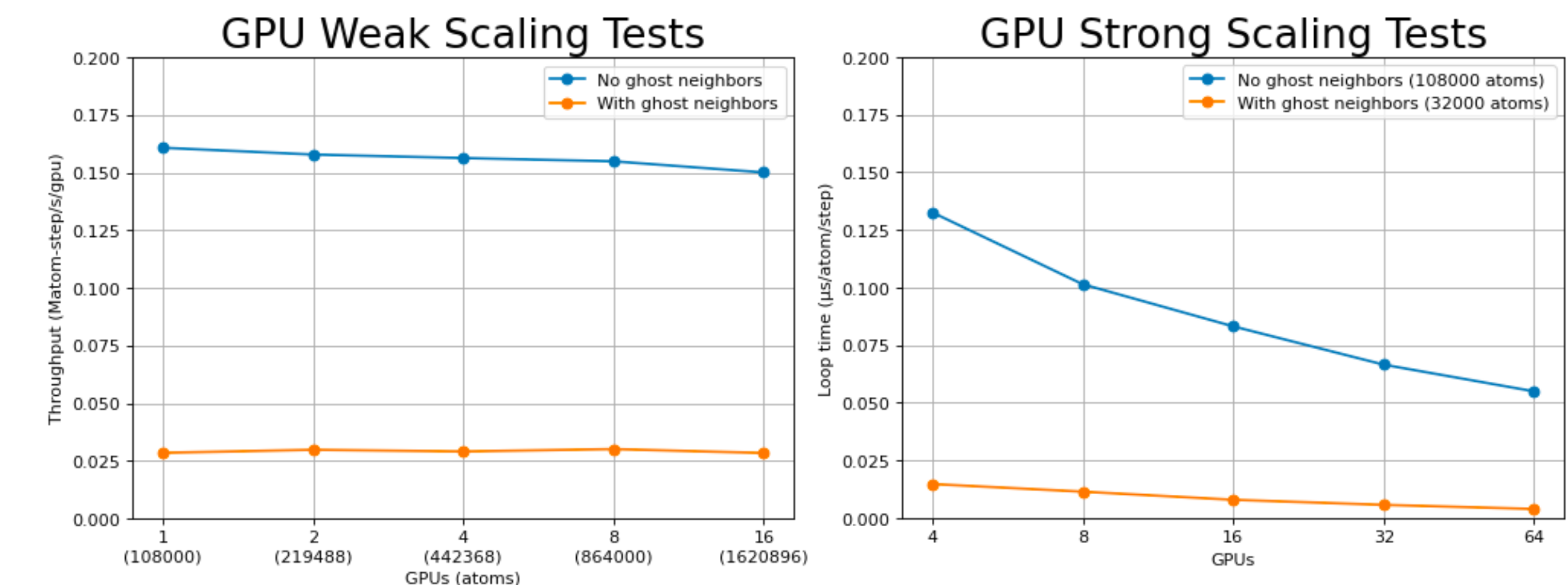
- SNAP: machine-learning interatomic potential that uses bispectrum components
- Unified HIP-NN has approx. 9X shorter loop time
- Energy and pressure differ considerably, but neither model considered ground truth

Unified HIPNN										SNAP									
in.mliap.unified.hippyinn.InP										in.snap.InP.JCPA2020									
Per MPI rank	memory allocation (min/avg/max)	= 996.7		996.7		Mbytes				Per MPI rank	memory allocation (min/avg/max)	= 6.027		6.027		Mbytes			
Step	Temp	E_pair	E_mol	TotEng	Press	E_pair	E_mol	TotEng	Press	Step	Temp	E_pair	E_mol	TotEng	Press	E_pair	E_mol	TotEng	Press
0	300	-4.8412739	0	-4.8025716	-2305.4876	-4.8025716	0	-3.4418771	1353.5968	0	300	-3.4805794	0	-3.4418771	1353.5968	-3.4805794	0	-3.4418771	1353.5968
10	289.4449	-4.8399121	0	-4.8025714	-2253.981	-4.8025714	0	-3.4418766	1611.7131	10	285.84677	-3.4787531	0	-3.4418766	1611.7131	-3.4787531	0	-3.4418766	1611.7131
20	259.98539	-4.836111	0	-4.8025709	-2005.1126	-4.8025709	0	-3.4418756	2312.0308	20	248.14649	-3.4738884	0	-3.4418756	2312.0308	-3.4738884	0	-3.4418756	2312.0308
30	218.06383	-4.8307021	0	-4.8025701	-1448.3016	-4.8025701	0	-3.4675394	3168.1943	30	198.94136	-3.4675394	0	-3.4675394	3168.1943	-3.4675394	0	-3.4675394	3168.1943
40	170.34948	-4.8249327	0	-4.8025693	-686.48167	-4.8025693	0	-3.4615791	3903.5749	40	152.74831	-3.4615791	0	-3.4615791	3903.5749	-3.4615791	0	-3.4615791	3903.5749
50	136.32353	-4.8201554	0	-4.8025686	13.292264	-4.8025686	0	-3.4566863	4387.1254	50	121.9796	-3.4566863	0	-3.4566863	4387.1254	-3.4566863	0	-3.4566863	4387.1254
60	114.93675	-4.8173961	0	-4.8025684	410.54349	-4.8025684	0	-3.4418729	4556.3803	60	113.27555	-3.4566863	0	-3.4418729	4556.3803	-3.4418729	0	-3.4418729	4556.3803
70	111.96058	-4.8170124	0	-4.8025686	404.16934	-4.8025686	0	-3.4418735	4431.2883	70	125.68089	-3.4580873	0	-3.4418735	4431.2883	-3.4580873	0	-3.4418735	4431.2883
80	124.36461	-4.8186132	0	-4.8025693	58.262582	-4.8025693	0	-3.4614159	4107.2369	80	151.47475	-3.4614159	0	-3.4614159	4107.2369	-3.4614159	0	-3.4614159	4107.2369
90	144.90639	-4.821264	0	-4.80257	-422.85179	-4.80257	0	-3.4418754	3739.5881	90	179.18708	-3.4649919	0	-3.4418754	3739.5881	-3.4649919	0	-3.4418754	3739.5881
100	164.9825	-4.8238546	0	-4.8025705	-800.75432	-4.8025705	0	-3.441876	3492.7778	100	197.50662	-3.4673559	0	-3.441876	3492.7778	-3.4673559	0	-3.441876	3492.7778

Loop time of 2.42707 on 1 procs for 100 steps with 512 atoms
Performance: 1.780 ns/day, 13.484 hours/ns, 41.202 timesteps/s
99.4% CPU use with 1 MPI tasks x 1 OpenMP threads

Loop time of 21.9953 on 1 procs for 100 steps with 512 atoms
Performance: 0.196 ns/day, 122.196 hours/ns, 4.546 timesteps/s
99.8% CPU use with 1 MPI tasks x 1 OpenMP threads

Unified HIP-NN GPU Scaling Tests



- Tested on Chicoma with NVIDIA A100 GPUs
 - 1 dual-socket CPU per node
 - Weak scaling: 1 GPU per node
 - Strong scaling: 4 GPUs per node
 - 1 process per GPU
 - 16 OpenMP threads per process
- Higher is better; flat, horizontal line is ideal**
- Efficient scaling with modified (ghost) neighbor lists
- HIP-NN highly competitive with other ML interatomic potentials such as SNAP and ASE

Conclusion

- LAMMPS enhanced with support for arbitrary, easy-to-make Python models
- ML and GPU acceleration provide LAMMPS Python models with major performance advantage**
- ML-IAP Unified + HIP-NN highly competitive with other machine-learning interatomic potentials
- ML driving HPC into the future!**

Future Work

- Successfully merge code into LAMMPS GitHub repo
- Produce thorough documentation of each Unified Interface component and usage
- Perform scaling tests with more than 64 GPUs
- Implement other ML interatomic potential models

