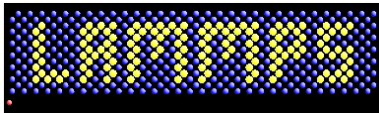


Lecture #3 – Styles, styles, styles

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Goals for this lecture

Teach you ...

- What styles are in LAMMPS
- The most versatile styles and what they do
- How styles let you tailor your LAMMPS simulation

What is a style in LAMMPS lingo

- A command with many variants
 - `pair_style` \Rightarrow `pair_style lj/cut`, `pair_style eam`, etc
 - `fix` \Rightarrow `fix nve`, `fix rigid/small`, etc
- Why not command names = `pair` or `fix_style` ?
 - single instance: command name includes **style**
 - many instances: command name w/out **style**
- A virtual parent class with many derived child classes
 - parent = `src/pair.cpp` or `src/fix.cpp`
 - children = `pair_lj_cut.cpp`, `pair_eam.cpp`, etc
 - children = `fix_nve.cpp`, `fix_rigid_small.cpp`, etc

20 different styles in current LAMMPS

Most are invoked as input script commands, some are internal

atom	25	sets of per-atom properties
pair	230	pairwise and manybody potentials
bond/angle/dihedral	15/20/15	intra-molecular interactions
kspace	15	long-range Coulombic solvers
fix	225	operations while timestepping
compute	140	diagnostic calculations
region	8	geometric regions
dump	25	output of simulation snapshots
integrate	2	Verlet or rRESPA algorithms
minimize	9	energy minimization algorithms
command	45	added input script commands create_box, create_atoms, run, etc

Pair styles

- LAMMPS lingo for interatomic potentials or force fields
- Define how atoms interact each other
 - short-range portion only
- Critical choice to make for your model and material
- Trade-offs in accuracy vs computational cost
- A pair style can be **pair-wise** or **many-body**
 - LJ, Coulombic, Buckingham, Morse, Yukawa, ...
 - EAM, Tersoff, REBO, ReaxFF, ...
- **Short-range Coulombics** included in pair style
 - lj/cut, lj/cut/coul/cut, lj/cut/coul/long, lj/cut/coul/wolf
 - done to optimize inner loop

Categories of pair styles for different materials

- Solids
 - eam, eim, meam, adp, etc
- Bio and polymers
 - charmm, class2, gromacs, dreiding, etc
- Reactive or bond-order
 - tersoff, bop, airebo, comb, reaxff, etc
- Coarse-grained
 - dpd, granular, sph, peri, colloid, lubricate, brownian, etc
- Aspherical
 - gayberne, resquared, line, tri, etc

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- **Pair table** for tabulation of any pair-wise interaction
- **Pair hybrid** enables modeling of hybrid systems
 - polymers on metal
 - CNTs in water
 - solid-solid interface between 2 materials

List of pair styles available natively in LAMMPS

See website [Commands](#) ⇒ [Pair Styles](#)

Annotated with (gikot) for 5 accelerated variants

5.8. Pair_style potentials

All LAMMPS [pair_style](#) commands. Some styles have accelerated versions. This is indicated by additional letters in parenthesis: g = GPU, i = INTEL, k = KOKKOS, o = OPENMP, t = OPT.

none	zero	hybrid (k)	hybrid/overlay (k)
hybrid/scaled	kim	list	
adp (o)	agni (o)	airebo (io)	airebo/morse (io)
atm	awpmd/cut	beck (go)	body/nparticle
body/rounded/polygon	body/rounded/polyhedron	bop	born (go)
born/coul/dsf	born/coul/dsf/cs	born/coul/long (go)	born/coul/long/cs (g)

- LAMMPS [pair kim](#) = interface to OpenKIM repository
- Ellad Tadmor [talk: Thurs early session](#)
- OpenKIM [breakout: Fri](#)

One-line descriptions of pair styles

See [pair_style.html](#) doc page

- [none](#) - turn off pairwise interactions
- [hybrid](#) - multiple styles of pairwise interactions
- [hybrid/overlay](#) - multiple styles of superposed pairwise interactions
- [hybrid/scaled](#) - multiple styles of scaled superposed pairwise interactions
- [zero](#) - neighbor list but no interactions
- [adp](#) - angular dependent potential (ADP) of Mishin
- [agni](#) - AGNI machine-learning potential
- [airebo](#) - AIREBO potential of Stuart
- [airebo/morse](#) - AIREBO with Morse instead of LJ
- [atm](#) - Axilrod-Teller-Muto potential
- [awpmd/cut](#) - Antisymmetrized Wave Packet MD potential for atoms and electrons
- [beck](#) - Beck potential
- [body/nparticle](#) - interactions between body particles
- [body/rounded/polygon](#) - granular-style 2d polygon potential
- [body/rounded/polyhedron](#) - granular-style 3d polyhedron potential
- [bop](#) - BOP potential of Pettifor

Relative CPU cost of different potentials is dramatic

See website [Benchmark](#) \Rightarrow [Interatomic potential](#) for details

Useful to estimate how long your simulation will run

Potential	System	# Atoms	Timestep	Neighs/atom	Memory	CPU	LJ Ratio
Granular	chute flow	32000	0.0001 tau	7.2	33 Mb	2.08e-7	0.26x
FENE bead/spring	polymer melt	32000	0.012 tau	9.7	8.4 Mb	2.86e-7	0.36x
Lennard-Jones	LJ liquid	32000	0.005 tau	76.9	12 Mb	8.01e-7	1.0x
DPD	pure solvent	32000	0.04 tau	41.3	9.4 Mb	1.22e-6	1.53x
EAM	bulk Cu	32000	5 fmsec	75.5	13 Mb	1.87e-6	2.34x
REBO	polyethylene	32640	0.5 fmsec	149	33 Mb	3.18e-6	3.97x
Stillinger-Weber	bulk Si	32000	1 fmsec	30.0	11 Mb	3.28e-6	4.10x
Tersoff	bulk Si	32000	1 fmsec	16.6	9.2 Mb	3.74e-6	4.67x

AIREBO	polyethylene	32640	0.5 fmsec	681	101 Mb	3.25e-5	40.6x
ReaxFF/C	PETN crystal	32480	0.1 fmsec	667	976 Mb	1.09e-4	136x
COMB	crystalline SiO2	32400	0.2 fmsec	572	85 Mb	2.00e-4	250x
eFF	H plasma	32000	0.001 fmsec	5066	365 Mb	2.16e-4	270x
ReaxFF	PETN crystal	16240	0.1 fmsec	667	425 Mb	2.84e-4	354x

Bond styles (also angle, dihedral, improper)

LAMMPS lingo for intra-molecular 2,3,4-body interactions

- Used for molecules with **fixed covalent bonds**
 - Fix bond/react can form and break them
 - Jake Gissinger **talk: Thurs late session**
- To learn what bond styles LAMMPS has ... where to look?

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- Website **Commands** \Rightarrow **Bond Styles** or **bond_style.html** doc page

5.9. Bond_style potentials

All LAMMPS **bond_style** commands. Some styles have accelerated versions. This is indicated by additional letters in parenthesis: g = GPU, i = INTEL, k = KOKKOS, o = OPENMP, t = OPT.

none	zero	hybrid	
class2 (ko)	fene (iko)	fene/expand (o)	gaussian
gromos (o)	harmonic (iko)	harmonic/shift (o)	harmonic/shift/cut (o)
mm3	morse (o)	nonlinear (o)	oxdna/fene
oxdna2/fene	oxrna2/fene	quartic (o)	special
table (o)			

- **none** - turn off bonded interactions
- **zero** - topology but no interactions
- **hybrid** - define multiple styles of bond interactions
- **class2** - COMPASS (class 2) bond
- **fene** - FENE (finite-extensible non-linear elastic) bond
- **fene/expand** - FENE bonds with variable size particles
- **gaussian** - multicentered Gaussian-based bond potential
- **gromos** - GROMOS force field bond
- **harmonic** - harmonic bond
- **harmonic/shift** - shifted harmonic bond

KSpace styles

LAMMPS lingo for long-range Coulombic solvers

Web **Commands** \Rightarrow **Kspace Styles** or **kspace_style.html** doc page

- Options:
 - traditional **Ewald**, scales as $O(N^{3/2})$
 - **PPPM** (like PME), scales as $O(N \log(N))$
 - **MSM**, scales as $O(N)$
- Additional options:
 - non-periodic: PPPM (z) or MSM (xyz)
 - long-range dispersion (LJ) or dipolar

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- Additional options:
 - non-periodic: PPPM (z) or MSM (xyz)
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- **PPPM is fastest** choice for most systems
 - FFTs can scale poorly for large processor counts
- **MSM can be faster** for low-accuracy or large proc counts
- Pay attention to cutoff & accuracy settings
 - adjusts Real-space versus KSpace work
 - can affect performance (see logfile timings)

Fix styles

Most **flexible feature** in LAMMPS

Fixes enable control of **what** happens **when** in a timestep

Internal flags determine when different methods of fix are invoked

communicate ghost atoms

build neighbor list (once in a while)

compute forces

communicate ghost forces

output to screen and files

Fix styles

Most **flexible feature** in LAMMPS

Fixes enable control of **what** happens **when** in a timestep

Internal flags determine when different methods of fix are invoked

Loop over timesteps:

fix initial NVE, NVT, NPT, rigid-body integration

communicate ghost atoms

fix neighbor insert particles

build neighbor list (once in a while)

compute forces

communicate ghost forces

fix force SHAKE, langevin drag, wall, spring, gravity

fix final NVE, NVT, NPT, rigid-body integration

fix end volume & T rescaling, diagnostics

output to screen and files

More on fix styles

- Assign each fix a unique alphanumeric ID
- Choose what group of atoms to apply fix to

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- Choose what group of atoms to apply fix to
- Examples of groups in obstacle script:
 - fix 1 **all** nve
 - fix 2 **flow** temp/rescale 200 1.0 1.0 0.02 1.0
 - fix 5 **upper** aveforce 0.0 -0.5 0.0
 - fix 6 **flow** addforce 1.0 0.0 0.0

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 - fix 6 **flow** addforce 1.0 0.0 0.0
- An input script may use dozens of fixes
 - same fix multiple times, on different groups of atoms
- Fixes can store **output** or **persistent info**
 - thermostat energy, forces on wall, time-zero atom coords
 - accessible by other commands or variables or log/dump output

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If familiarize yourself with fixes,

you'll know many things LAMMPS can do

Website **Commands** \Rightarrow **Fix Styles** or **fix.html** doc page

Compute styles

LAMMPS lingo for a **diagnostic calculation**

- Calculate some property of the system
- Always for the **current timestep**
- **Defining** a compute in an input script does not **invoke** it
- Fixes or output commands (thermo, dump) invoke computes, only on timesteps when needed

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Examples:

- Thermostat fixes: **compute** temp, temp/asphere
- Time averaging **fixes**:
fix ave/time, ave/chunk (spatial), ave/atom, ave/histo
- Thermo output to logfile: **compute** temp, pe, press
- Dump files: **compute** coord/atom, cna/atom, voronoi/atom

More on compute styles

- All computes **store output**:
 - global vs per-atom vs local info
 - compute with **/atom** suffix produces per-atom info
 - compute with **/local** suffix produces local info
 - scalar vs vector vs array data structure
 - accessible by other commands or variables or log/dump output
- **Examples**:
 - temp & pressure = global scalar or vector
 - pe/atom = potential energy per atom (vector)
 - displace/atom = displacement per atom (array)
 - pair/local & bond/local = per-neighbor or per-bond info

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To learn what compute styles LAMMPS has ... where to look?

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