

Molecular Dynamics in Dyalog APL

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- What is Molecular Dynamics?
- In what ways is APL useful?
 - Teaching MD
 - Developing MD

What is MD?

- Modelling atomic and molecular systems using classical (Newtonian) mechanics

$$\vec{F} = m\vec{a}$$

$$F = - \frac{dV(x)}{dx}$$

Numerical Integration

- Insoluble ODEs
- Step from state to state using numerical approximations
- Velocity Verlet

What MD codes are there?

- LAMMPS
- GROMACS
- HOOMD
- + many more...

<https://lammps.sandia.gov/other.html>

APL for MD

- Teaching MD
 - Explanation
 - Visualisation

The Lennard-Jones fluid

Link to Jupyter notebook in the description ↓

<https://nbviewer.jupyter.org/github/rikedyp/APLPhys/blob/master/Jupyter/LJmelt.ipynb>

$$V_{LJ} = 4\epsilon \left[\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^6 \right]$$

LJ ← {4×-/ω°. *⁻¹² -⁻⁶}

The Lennard-Jones fluid

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$$\mathbf{F} = -\frac{1}{r} \nabla V(\mathbf{r})$$

$$\mathbf{F} = -\frac{1}{r} \nabla V_{LJ}(\mathbf{r}) = -\frac{1}{r} \frac{dV_{LJ}}{dr} = -24 \left[2 \left(\frac{\sigma}{\mathbf{r}} \right)^{14} - \left(\frac{\sigma}{\mathbf{r}} \right)^8 \right]$$

Thermostats: TempRescale

$$\Delta T = \frac{1}{2} \sum_{i=1}^2 \frac{m_i (\lambda v_i)^2}{N_{df} k_B} - \frac{1}{2} \sum_{i=1}^2 \frac{m_i v_i^2}{N_{df} k_B}$$

$$\Delta T = (\lambda^2 - 1) T(t)$$

$$\lambda = \sqrt{T_0 / T(t)}$$

`{omega * 0.5 * ~fixtemp ÷ alpha}`



Thermostats: Berendsen

$$T = T_0 - Ce^{-t/\tau}$$

$$\lambda^2 = 1 + \frac{\Delta t}{\tau_T} \left(\frac{T_0}{T} - 1 \right)$$

APLPhys

<https://github.com/rikyedyp/APLPhys>

- Jupyter notebooks
- Key  and rank 
- MiServer GUI (MiPhys)

Future

<https://github.com/rikydyp/APLPhys>

- Scalability: Isolates
- More thermostats
- More models
- Graphs on MiPhys