A lightning-fast introduction to molecular <u>dynamics</u>

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What are MD simulations?

- Basic idea: simulate molecules in time by calculating forces at each instant and applying Newton's second law
- Do this over and over...usually 10⁶ to 10¹² steps
- Underlying assumption: $\overline{\Theta}_t = \langle \Theta \rangle$ (ergodic hypothesis)

A basic MD integration scheme

Supposing we have initial coordinates and velocities:

$$\left\{\vec{x}_i(t_0)\right\}$$
 and $\left\{\vec{v}_i(t_0)\right\}$

Integration scheme (how to propagate in time):

$$\vec{x}_{i}(t_{0} + \Delta t) = \vec{x}_{i}(t_{0}) + \frac{\partial \vec{x}_{i}}{\partial t}\Big|_{t_{0}} \Delta t + \frac{1}{2} \frac{\partial^{2} \vec{x}_{i}}{\partial t^{2}}\Big|_{t_{0}} \Delta t^{2} + \dots = \vec{x}_{i}(t_{0}) + \vec{v}_{i}(t_{0}) \Delta t + \frac{1}{2} \vec{a}_{i}(t_{0}) \Delta t^{2}$$

$$\vec{a}_i(t_0) = \frac{1}{m_i} \vec{F}_i(t_0) = -\frac{1}{m_i} (\nabla U)_i$$

...and then increment by another Δt

...and some fancier things

- Suppose we want to carry out a simulation at constant temperature (e.g., NVT): <u>thermostats</u> can modify atomic velocities accordingly
- Suppose we want to carry out a simulation at constant pressure (e.g., NPT): <u>barostats</u> can modify system volumes accordingly

The bare minimum to run MD

- Initial coordinates of atoms (3D structure)
- <u>Topology</u> (which atoms are connected to each other and how)
- Force field (to calculate energies/forces)
- Integrator (to propagate in time)

A word on MD time steps (1/2)

- How big is Δt ? Typically 1 fs (due to X-H bond vibration frequencies).
- Want a larger Δt ?
 - Constrain X-H bonds: 2 fs
 - Repartition mass from heavy atoms to H atoms: 4 fs
- Moral of the story: For all-atom MD simulations, we're stuck using atomic timescales.

A word on MD time steps (2/2)

- If you really want to speed up sampling in your simulations, you'll need an *enhanced sampling* method (e.g., replica exchange MD, accelerated MD, or metadynamics).
- Additionally, there are multiple time step methods (e.g., Margul and Tuckerman, JCTC 2016) that can effectively lengthen Δt by 1-2 orders of magnitude.

Diagram of classes in OpenMM 6.0

