Molecular Dynamics Primer

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Simulation vs. Experiment

Experiment

- Setup experimental conditions
- Measure quantity of interest
- Analyze apply theory

Simulation

- Setup Model and approximations
- Calculate subject setup to conditions
- Analyze - measure,match, possibly predict.

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What is Molecular Dynamics?

Molecular Dynamics(MD) is classical!

Think Of Pool!

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What is Molecular Dynamics? Molecular Dynamics(MD) is classical! \vec{F} Force on an atom = Mass of an atom Acceleration of an atom m \vec{a} MD As Hard-Spheres

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MD is the application of classical mechanics to treat the evolution of atoms interacting via a determined potential at nanometer length and picosecond time scales.

Adapted from www.wag.caltech.edu

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What Do Atoms Look Like in MD

QM vs. MD Atom

The Approach

- Nuclei and Electrons are not explicitly treated
- Describe atoms with an interatomic potential(i.e. Model)
- **•** Given Newtons laws of motion evolve a system
- Modify equations to include temperature,pressure,etc.

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Limitations of MD

- Time and length scale: nanoseconds and nanometers
- Interatomic potentials dictate potential energy surface
- Chemistry should be carefully interpreted
- Know the limitations of the MD model

Should You Use Other Methods

- Does your phenomena fall outside the range of time and length scale of MD: Spatial - Dislocation Dynamics Time - Diffusion
- Do you need the effects of electrons -; QM calculations

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Classifying Materials

Since we have eliminated the direct interaction of electrons which are responsible for chemistry (i.e. bonding) are models need to same something about the bonding.

- Dispersion forces
- Ionic bonding
- **•** Covalent bonding
- Metallic bonding

Adapted from http://www.abc.net.au

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Cohesive Energy

- Potential energy is the sum of all interacting atoms
- Cohesive energy is the energy need to move all the atoms infinitely apart

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Pair Potentials

- Energy and force are sum of atom pair interactions
- Only dependent on separation distance
- • Some examples are: Lennard-Jones, Buckingham, and Morse

Adapted from Callister, William D. ,Jr. Materials Science And Engineering An Introduction, 7th ed.

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Dispersion forces

- No directionality to bond
- **•** Pairwise interaction between atoms
- Repulsive nature due need for electron orthogonality (i.e. Pauli exclusion principle)
- **•** Attractive nature from instantaneous electron fluctuations around nuclei

Adapted from LeSar, Richard. Introduction to Computational Materials Science: Fundamentals to Applications, 1st ed.

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Lennard-Jones

Very common simple interatomic potential good for noble gases and molecular interactions.

 ϵ : potential well depth

σ: distance were energy is zero

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Adapted from <http://chemwiki.ucdavis.edu>

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How Well Does LJ Do!

Other Simple Potentials

- Mie Potential adds two more variables for fitting
- Morse Potential approach from diatomic bond potentials

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Ionic Bonding

Coulombic In Nature

- **•** Atoms in ionic materials behave like point charges
- Simple to describe by Coulomb potential
- Very long-range interactions

Terms Added For Correction

- Usually a repulsive "Wall" is needed
- Also Dipole-Dipole or Multipole expansion added: $\frac{C}{r_{ij}^6}$
- Referred to as Born or Born-Huggins-Meyer potential

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Pair Potential Limitations

- **Pair Potentials only function of distance**
- For pure solids will always produce simple structures (e.g FCC)
- Pair potentials obey Cauchy relations: $c_{12} = c_{44}$
- **•** Predominately make use of empirical data

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Many-body Potentials

Types of Systems

Primarily used to describe metallic and covalent materials. Fit using empirical and ab-initio data.

Metals

Embedded atom method/Modified embedded atom method:

- **•** Environment dependent interatomic potentials
- • Glue Model/Effective medium theory

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Covalent Materials

- Stillinger-Weber: 2-body plus 3-body
- **•** Tersoff (Bond-Order)
- REBO/AIREBO adds hydrocarbon descriptions

Example: Tersoff bond-order term $E_i = f_R(r_{ij}) + \widehat{b_{ij}} \widehat{f}_A(r_{ij}); b_{ij}(r_{ij}, r_{ik}, \theta_{ijk})$ Si_n 0.95 0.9 0.85 0.8 b_{ii} 0.75 0.7 0.65 0.6 0.55 120 140 160 180 100 θ_{kij}

Umeno, Y., et al. Ideal Shear Strength under Compression and Tension in C, Si, Ge, and Cubic SiC: An Ab Initio Density Functional Theory Study. J. Phys.: Condens. Matter 2011, 23

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Other Approaches

Force Fields

- Used predominately is chemistry and biology
- Force Field descriptions:

$$
E = E_{bond} + E_{angle} + E_{dihedral} + E_{coul} + E_{vdw}
$$

- **•** Force Fields use fixed description of bonds i.e. single, double, triple
- Allows for simulation of mixed interactions

Adapted from http://mmb.irbbarcelona.org

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Advanced Potentials

- **Reactive Force Fields**
- Multi-component systems
- On-The-fly determination of bond-order
- Also includes charge transfer (IMPORTANT to define chemistry)
- Parametrized from large ab-initio data sets
- 10-100x more computationally expensive

Adapted from http://www.rxffconsulting.com : battery electrolyte with graphite electrodes

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Integration of Newtons Equations

Verlet Algorithm

Taylor expansion of position in time:

$$
r(t + \Delta t) = r(t) + v(t)\Delta t + \frac{1}{2!}a(t)\Delta t^{2} + \cdots
$$

$$
r(t + \Delta t) = r(t) - v(t)\Delta t + \frac{1}{2!}a(t)\Delta t^{2} - \cdots
$$

$$
r(t + \Delta t) = r(t) + v(t)\Delta t + \frac{a(t)}{2}\Delta t^{2}
$$

- Accurate $(O(\Delta t^4))$ and simple
- No need to calculate velocity
- Unfortunately need to calculate velocity indirectly:

$$
v(t) = \frac{r(t + \Delta t) - r(t - \Delta t)}{2\Delta t}
$$

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Velocity Verlet Algorithm

- Most widely used
- Makes use of half time steps:

$$
v\left(t+\frac{\Delta t}{2}\right)
$$

Calculates positions from velocity, then updates velocity:

$$
r(t + \Delta t) = r(t) + v\left(t + \frac{\Delta t}{2}\right)\Delta t
$$

$$
v(t + \Delta t) = v\left(t + \frac{\Delta t}{2}\right) + \frac{1}{2}a(t + \Delta t)\Delta t
$$

- Simple to implement and stable
- **•** Time reversible as well

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High-Order Algorithm

- Predictor-corrector algorithm
- Measures error in acceleration and correct
- **Gear Predictor-corrector even more accurate**
- typical timestep for all algorithms is 1-5 fs

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Potential Cutoffs & Neighbor Lists

Techniques

- Most computationally expensive part of MD is calculating energy and forces.
- \bullet Use cutoffs to reduce computing time, r_c
	- \bullet FAM: 5 \AA
	- Tersoff: 3-5 \AA
	- Coulombic: long-range summation needed
	- Pair potentials: $10-15$ \AA
- Populate a list of neighbors every ∆t

Adapted from http://catalyst.blogs.rice.edu/

- Verlet list
- Link-cell list

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Boundary Conditions

- **•** Periodic Boundary Conditions make use of images
- **•** Free Surface Boundary Conditions vacuum above surface

Adapted from http://catalyst.blogs.rice.edu/

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Position and Velocities

- MD requires some knowledge of initial structure
- We can prescribe velocities using a distribution corresponding to a desired temperature

Defining Temperature & Pressure

• Use equipartition theorem:

$$
\langle E_{ke} \rangle = \frac{3}{2} N k_b T = \langle \sum_i \frac{1}{2} m v_i^2 \rangle
$$

$$
T(t) = \frac{1}{3Nk_b} \sum_i mv_i^2
$$

• Virial pressure:

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Ensembles

- Atoms in a simulation box during MD run sample many microstates
- Thermodynamic limit at equilibrium all possible configurations sampled
- We can impose external conditions on ensembles (i.e. Temperature, pressure, volume)

List of Ensembles

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Visual Aid

Adapted from Lee, June Gunn. Computational Materials Science: An Introduction; CRC Press, 2012.

Implementation Example: Thermostats

- **Berendsen rescale velocities**
- Nose-Hoover modify the Hamiltonian of the system
- Langevin force damping
- **•** Barostats are carried out in a similar fashion

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Analysis

Some Key Aspects

- When running NVE, system must conserve energy!
- **•** The net linear momentum of the simulation cell should be zero
- **•** Try to identify if in a local minimum
- Check for numerical errors: bad dynamics, too large Δt ,etc.

Time-Averaging

Since the thermodynamic quantities we calculate are instantaneous we take the time average at equilibrium.

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Visualization

Inspect For Issues

- Visualize trajectories of atoms
- Color code according to properties (e.g. energy per atom)
- Check for any mistakes in initial structure

Simply Look!

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Equilibrium

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Properties

- Bulk modulus: $B = -V \left(\frac{\partial P}{\partial V} \right)$ $\frac{\partial P}{\partial V}$) nvt
- Lattice parameter, a
- Thermal expansion coefficient: $\alpha = \frac{1}{V}$ $\frac{1}{V} \left(\frac{\partial V}{\partial T} \right)$ $\frac{\partial V}{\partial \mathsf{T}})_{NPH}$
- **•** Heat Capacity
- **•** Elastic Constants

Structural Information

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Other Properties

Viscosity: Green-Kubo Method

Thermal Conductivity: Green-Kubo Method

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Go To Texts

Frenkel & Smit, Understanding Molecular Simulation: From Algorithms to Applications.Probably most through text on MD.

Allen & Tildesley, Computer Simulation of Liquids. Great MD intro and algorithms.

Rapaport, The Art of Molecular Dynamics Simulation. MD cookbook text, ideal for developing your own code.

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User Texts

J. Fan, Multiscale Analysis of Deformation and Failure of Materials. Nice section on using various codes.

J.G. Lee, Computational Materials Science: An Introduction. Ideal for quickly using LAMMPS and VASP.

R. Lesar, Introduction to Computational Materials Science: Fundamentals to Applications.Great multiscale introductory with strong emphasis on understanding and using.

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