#### **ME 517: Micro- and Nanoscale Processes**

## Lecture 28: Molecular Dynamics - I

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## Molecular Interaction Forces: Lennard-Jones 6-12 Potential

$$V_{ij}(r) = 4\varepsilon \left[ c_{ij} \left( \frac{r}{\sigma} \right)^{-12} - d_{ij} \left( \frac{r}{\sigma} \right)^{-6} \right]$$

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$$F_{ij}(r) = -\frac{\partial V_{ij}}{\partial r} = \frac{48\varepsilon}{\sigma} \left[ c_{ij} \left( \frac{r}{\sigma} \right)^{-13} - \frac{d_{ij}}{2} \left( \frac{r}{\sigma} \right)^{-7} \right]$$

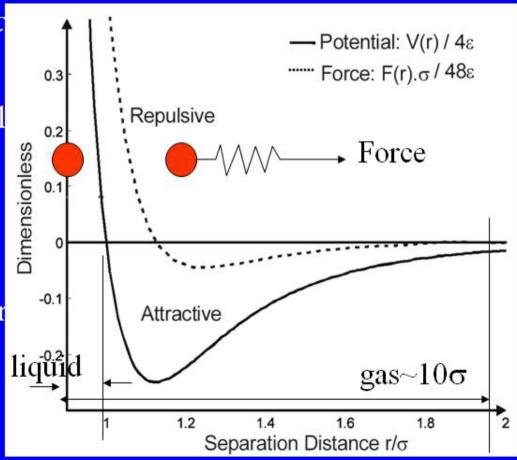
V<sub>ii</sub>=potential energy between t molecules i and j

F<sub>ii</sub>=force between two molecul i and i

c<sub>ij</sub> and d<sub>ij</sub> are parameters for chosen molecules

ε,σ are characteristic energy ar length scales respectively

r is the separation distance



## **Lennard Jones Constants**

$\varepsilon/K(K)$	$\sigma$ (nm)
97	0.362
91.5	0.368
190	0.400
113	0.343
124	0.342
	97 91.5 190 113

Phases	Intermolecular Forces	Ratio of Thermal Vibration Amplitude Compared to $\sigma$	Approach Needed
Solid	Strong	<b>«</b> 1	Quantum
Liquid	Moderate	~ 1	Quantum/classical
Gas	Weak	» 1	Classical

# Molecular Dynamics Governing Equations

$$m\frac{d^{2}\mathbf{r}_{i}}{dt^{2}} = \sum_{j \neq i} \frac{\partial V_{ij}}{\partial \mathbf{r}_{i}} - \frac{m}{\tau} \frac{d\mathbf{r}_{i}}{dt} + \eta_{i}$$

$$\tau = \sqrt{\frac{\sigma^{2}}{\varepsilon}}$$

$$\tau = \sqrt{\frac{\sigma^2 m}{\varepsilon}}$$

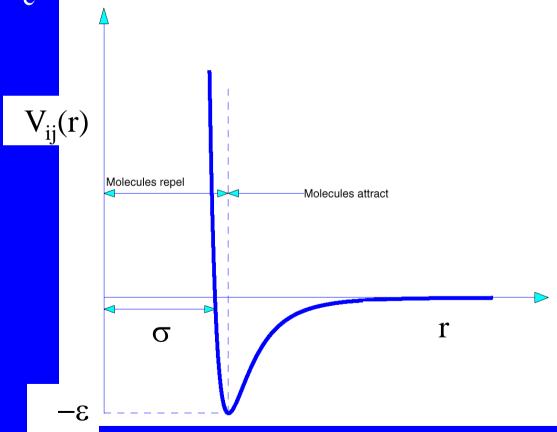


- where  $\mathbf{r_i}$  is the position vector,  $V_{ij}$  is the potential energy between any two molecules, τ is characteristic time scale, m is atomic mass
- Last two terms on RHS couple the particle dynamics with thermodynamics
  - Velocity term governs heat exchange with reservoir
  - η<sub>i</sub> term is a Gaussian stochastic force with variance  $2mk_b/\tau$
- For liquid argon,  $\tau=2.2^{-12}$  sec
- Evolve the position of every molecule forward in time using Newton's 2<sup>nd</sup> Law

#### Shifted Lennard-Jones Potential

$$V_{ij}(r) = 4\varepsilon \left[ c_{ij} \left( \frac{r}{\sigma} \right)^{-12} - d_{ij} \left( \frac{r}{\sigma} \right)^{-6} - \left( c_{ij} \left( \frac{r_c}{\sigma} \right)^{-12} - d_{ij} \left( \frac{r_c}{\sigma} \right)^{-6} \right) \right]$$

• r<sub>c</sub> is cut-off radius, typically 2.2σ<r<sub>c</sub><2.5σ



#### Lennard-Jones Potential

- Works reasonably well for electrically neutral,
   non polarizable, spherical molecules
  - Dynamics for immiscible liquids  $(d_{12}=d_{21})$ 
    - d<sub>12</sub>=0 implies pure short-range repulsion
    - d<sub>12</sub>=1 implies symmetric interaction -> single species
    - d<sub>12</sub>> implies enhanced attraction
  - Dynamics of wall boundaries can be simulated with same L-J approach but different constants
  - Complicated molecules require complicated potentials, e.g. polymer chains can have a potential between all monomers plus a strongly attractive potential when all monomers in neighboring molecules line up

Micro/Nanoscale Physical Processes

#### Temporal Evolution

- Equations of motions can be integrated forward in time by typical predictor-corrector scheme
  - typical time step size  $\Delta t=0.005\tau$
  - for liquid Ar,  $\Delta t=1.1e-14$  sec
- Another commonly used discretization called Verlet integration rule is

$$\boldsymbol{r}^{n+1} = 2\boldsymbol{r}^n - \boldsymbol{r}^{n-1} + \Delta t^2 \boldsymbol{a}(t) + \mathcal{O}(\Delta t^4)$$

#### Imposing External Forces

- Most flows driven by some external force
  - e.g. vibrating wall, pressure gradient, body force
- Need some way to couple in those forces
- Eulerian velocity computed as time average of N<sub>i</sub> molecules according to

$$oldsymbol{v}(oldsymbol{x}) = rac{1}{N_i} < \sum_j rac{doldsymbol{x}_j}{dt} >$$

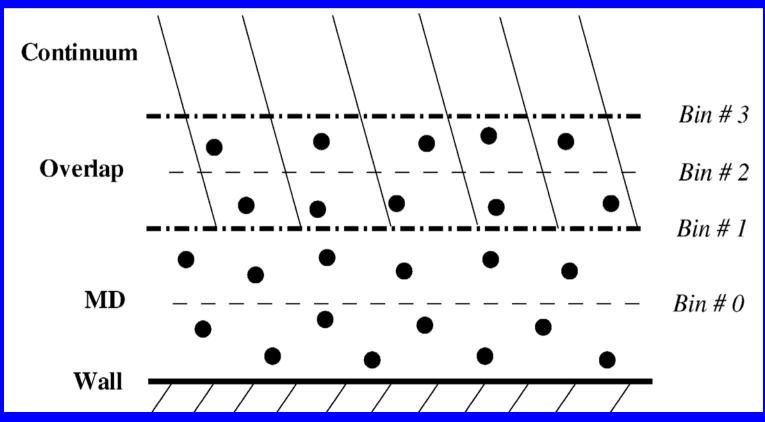
$$au(oldsymbol{x}) = rac{1}{V_i} < \sum_j \left[ rac{doldsymbol{x}_j}{dt} - oldsymbol{v}(oldsymbol{x}) 
ight] \left[ rac{doldsymbol{x}_j}{dt} - oldsymbol{v}(oldsymbol{x}) 
ight] + \sum_{j < i} oldsymbol{r}_{ij} oldsymbol{f}_{ij} >$$

#### Computational Complexity

- Have to sum over all pairs of molecules so order N<sup>2</sup>
- Cut-off distance reduces computational complexity somewhat
- Pairs of interacting molecules stored in Verlet list
  - For each molecule a=1,2,...N create a list of neighbors that are within a distance  $r_c+r_s$  where  $r_c$  is the cut-off distance and  $r_s$  is called a *skin thickness*
  - $r_s$  is chosen such that in a time interval of S~20  $\Delta t$ , no molecules from outside the skin enter the interaction range of molecule a
- Computational intensity reduced somewhat to  $(order\ N) + S^{-1} (order\ N^2)$

#### MD/Continuum Approach

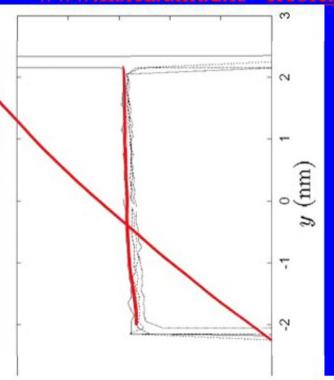
- Occasionally desirable to combine MD and Continuum approach (e.g. N-S)
  - For example external flow over a body



#### MD: Water Flow between Graphite Sheets

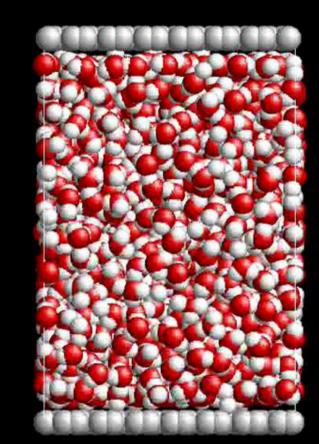
- ETH-Zurich simulated flows in and around CNTs and graphite sheets (Nanotech 2003)
- Exploring validity of no-slip assumption

www.fisica.uniud.it/~ercolessi/md/; www.icos.ethz.ch.

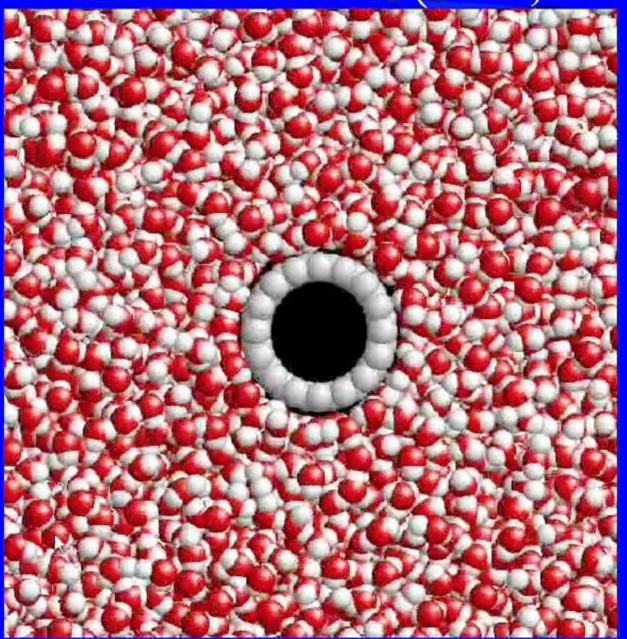


Slip lengths of 14-63 nm

→No-slip violated



# Flow Around CNTs (ETHZ)



#### Flow Around CNTs (ETHZ)

- Flow agrees quite well with continuum theory
- Slip length less than a single molecular diameter
- Considerable variations in fluid density near CNT

