





Molecular Dynamics: an outlook

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Molecular Dynamics: an outlook

- Where from MD simulations?
- What is MD?
- MD(or, for that, MC) as Equilibrium Statistical Mechanics
- A challenging case: Rare Events and Free Energies
- MD as Non-Equilibrium StatMech (→ multiscale etc)
- Hydrodynamics from an atomistic viewpoint if time permits
- Holonomic constraints and Blue Moon







The fundamental law

In principle, the behavior of a piece of matter in ordinary conditions comes out of *t*-dependent Schroedinger Equation and QStatMech with [Relativistic Quantum Field Theory not needed for that!]

$$\mathcal{H}(r, R; p, P) = K_N(P) + \underbrace{K_e(p) + \underbrace{V(r, R)}_{\text{Coulomb}}}_{H_e(r, p|R)}$$







Born-Oppenheimer approximation

since $m_N \gg m_e$

$$H_e(r, p | R)\Phi_s(r | R) = E_s(R)\Phi_s(r | R)$$

and $|\nabla_R \Phi_s| \ll |\nabla_r \Phi_s|$

$$\Psi(r, R; t) = \sum_{s} \chi_s(R; t) \Phi_s(r|R) \simeq \chi_0(R; t) \Phi_0(r|R)$$

i.e. the (often valid) adiabatic approximation where







 $\chi_0(R;t)$ is given by

$$i\hbar \frac{\partial}{\partial t} \chi_0(R;t) = \mathcal{H}_N(R,P) \chi_0(R;t)$$

$$\equiv [K_N(P) + E_0(R)] \chi_0(R;t)$$

the strict adiabatic approximation (no electronic jumps allowed)

the dynamics of the nuclei, apparently independent from the electrons, is driven by $E_0(R)$ as interaction potential (a mean field, modelizable, no more Coulomb!)







nuclei are heavy enough

when
$$\begin{cases}
\text{flucter are neavy enough} \\
\text{temperature is high enough so that} \\
\Lambda = \frac{h}{\sqrt{m_N k_B T}} << \text{internuclear } r
\end{cases}$$

Dynamics, no more quantum, is Newton:

$$m_N \ddot{R} = -\nabla E_0(R)$$







to sum up:

- lacktriangle a classical system of particles interacting via an effective interaction potential, E_0
- ♦ E_0 can be obtained *ab initio* (AIMD) or by some suitable fitting procedure→phenomenological model, e.g. with a Pairwise Additive Potential, $V(R) = \sum_{i < j} v(R_{ij})$
- ◆ if PAP, the equations of motion are numerically integrable for a number of particles finite but large enough to study, by statistical approach, the thermal properties of matter







Equilibrium (classical) Statistical Mechanics (1)

- a closed system evolving in time under timeindependent forces will reach a STATIONARY state
- the microscopic properties are 'irrelevant' while the statistical (or macroscopic) are stable and interesting (THERMODYNAMICS).

They can be computed by time or ensemble averages:







Equilibrium (classical) Statistical Mechanics (2)

$$O = \overline{\hat{O}(R, P)} \equiv \lim_{\tau \to \infty} \frac{1}{\tau} \int_0^{\tau} dt \, \hat{O}(R(t), P(t))$$

$$= \frac{1}{\mathcal{N}} \sum_{i=1}^{\mathcal{N}} \hat{O}\left(R(ih), P(ih)\right) = \sum_{l \in \text{visited states}} \left[\frac{n_l}{\mathcal{N}}\right] \hat{O}_l$$

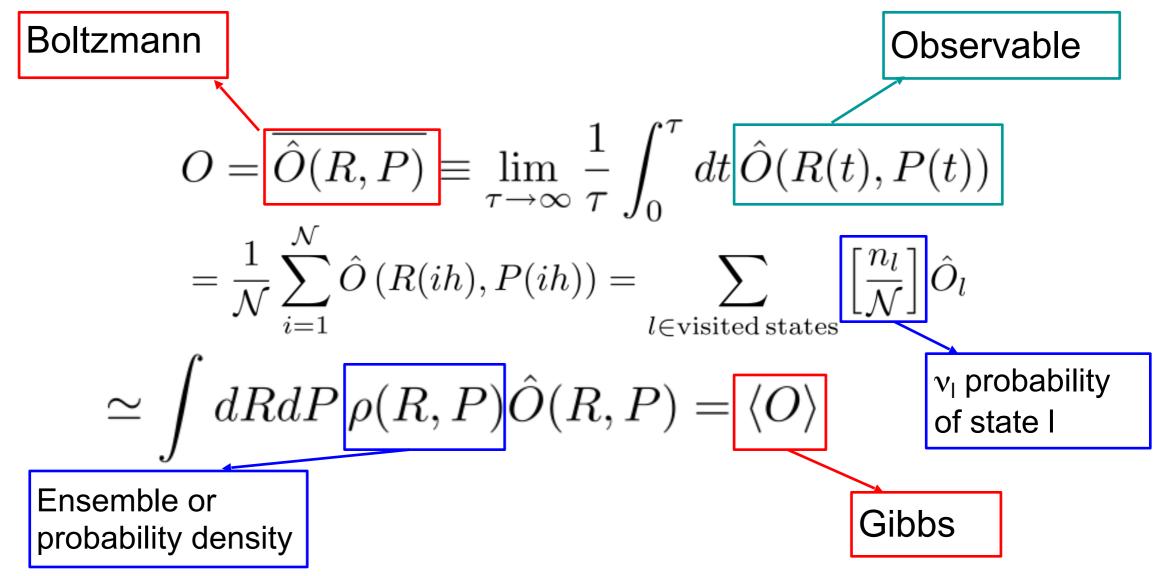
$$\simeq \int dR dP \, \rho(R, P) \hat{O}(R, P) = \langle \hat{O}(R, P) \rangle$$







Equilibrium (classical) Statistical Mechanics (3)



- ◆ properties coming from an observable → mechanical (e.g., pressure)
- \rightarrow properties coming from n_l or ρ , i.e. probability \rightarrow thermal (e.g., free energy)







(BO originated) Classical Stat. Mech. Model

Theoretically:

- $N \to \infty$ atoms / molecules \Longrightarrow point particles (p.p.), (r_1,\ldots,r_N) / connected sets of p.p., $(\{r_1\}\ldots\{r_N\})$ Interactions between p.p., $V_N (\{r_1\}\ldots\{r_N\})$
- · Boundary Conditions (compulsory, no BC's no equilibrium)
- · Initial Conditions (necessary to start although irrelevant for macroscopic behavior. However, they can be a headache!)
- Evolution laws: Newton equations and Laplace deterministic dream, $\{r(t; r_0, p_0), p(t; r_0, p_0)\}_{t \in (0, \tau \to \infty)}$







What is MD?

Computationally:

$$N \sim 32 \div 10^6 \, (10^9), \quad n = \frac{N}{V}$$

Boundary Conditions: Periodic (PBC)

for thermodynamic limit: min(S/V) effect

Initial Conditions: positions, regular lattice; velocities, maxwellian

 V_N : simple pairwise additive $\left(\sum_{i < j} v_{ij}; \mathcal{O}(N^2)\right)$; short range (MIC) extensions: \rightarrow long range (Coulomb) by Ewald sums

- → n-body potentials $\mathcal{O}(N^n)$ but glue potential $V_N(\alpha)$ with $\alpha = \sum_{i < j} \varphi_{ij} \dots$
- → stiff intramolecular potentials:

Constraints: $\sigma(r)=0; \ m\ddot{r}=F-\lambda\nabla\sigma$: SHAKE Multiple Time-Step (Martyna, Tuckerman, Berne) : RESPA

Integration Algorithms: robust, time reversible, symplectic e.g. velocity Verlet

Various ensembles (thermostats, barostats ...):
extended variables simulations (Andersen, Nosé, Hoover,...

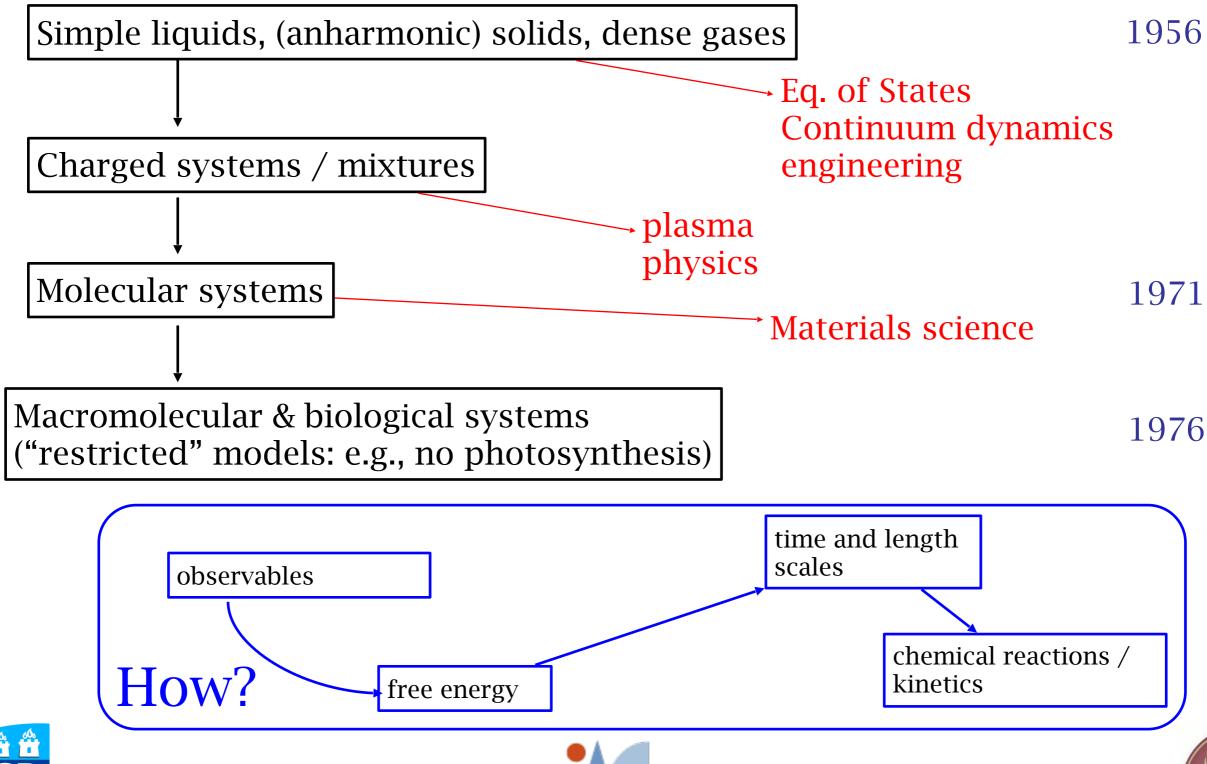




What is MD?

APPLICATIONS

What can be done?







Probabilistic interpretation of the thermal properties

• Entropy (Boltzmann)

$$S(E) = k_B ln \Sigma(E) \equiv -k_B ln \frac{1}{\Sigma(E)} = -k_B ln P_E(\Gamma)$$

Similarly in general ensembles

$$F(\cdots) = -k_B T \ln \mathcal{Z}(\cdots) \approx k_B T \ln [f(\Gamma)]$$

where $f(\Gamma)$ is the probability density function of the given ensemble







Mechanical vs thermal properties

$$\mathcal{O} = \lim_{\tau \to \infty} \frac{1}{\tau} \int_0^{\tau} dt \hat{\mathcal{O}}(\Gamma(t)) = \sum_s P_s \mathcal{O}_s = \\ = \int d\Gamma \underbrace{\left(\begin{array}{c} \exp[-\beta S(E)] \\ 0 \end{array} \right)}_{\frac{\delta(H(\Gamma) - E)}{\int d\Gamma \delta(H(\Gamma) - E)}} \hat{\mathcal{O}}(\Gamma) = \\ = \int d\mathcal{O}^* \mathcal{O}^* \left\{ \int d\Gamma \delta(\hat{\mathcal{O}}(\Gamma) - \mathcal{O}^*) \frac{\delta(H(\Gamma) - E)}{\int \delta(H(\Gamma) - E)} \right\} = \\ = \int d\mathcal{O}^* \mathcal{O}^* P_{\hat{\mathcal{O}}}(\mathcal{O}^*)$$







Free energy of collective variables

• Given a collective variable (i.e. a function of the configuration space) $\hat{\mathcal{O}}(x)$, the free energy associated with its probability density function is

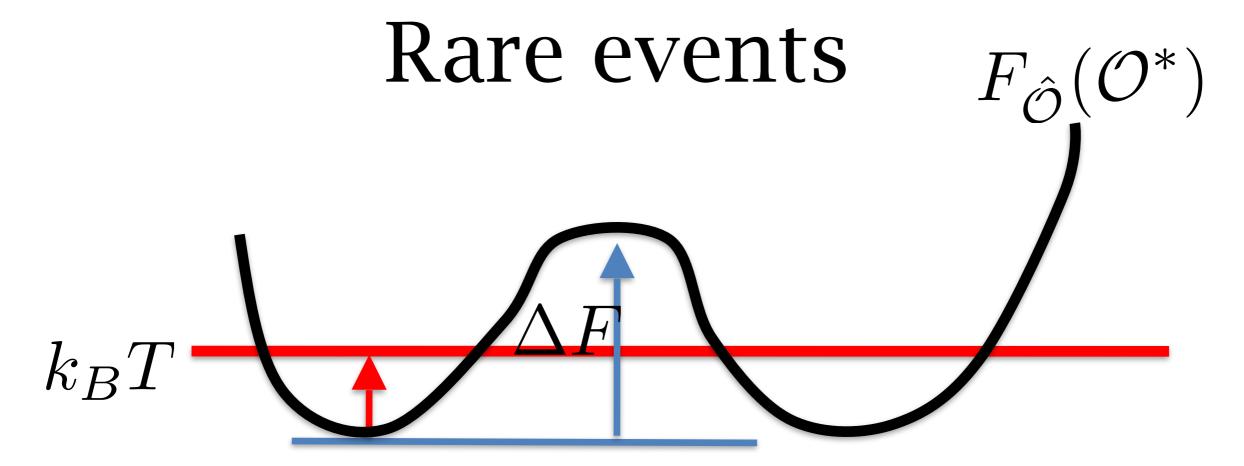
$$-\frac{F_{\hat{\mathcal{O}}}(\mathcal{O}^*)}{T} = k_B ln P_{\hat{\mathcal{O}}}(\mathcal{O}^*) =$$

$$k_B \ln \frac{\int d\Gamma f(\Gamma) \delta(\hat{\mathcal{O}}(x) - \mathcal{O}^*)}{\mathcal{Z}}$$









$$au \propto \exp\left[rac{\Delta F}{k_B T}
ight]$$

• If $\Delta F >> k_B T$ then $\tau >> t_{max}$







TA-MD/MC (<u>Temperature Accelerated Molecular Dynamics/Monte Carlo</u>)

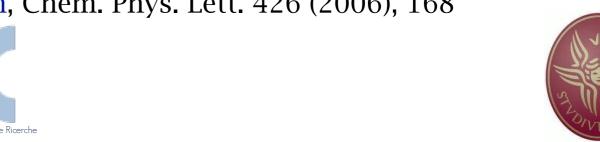
Accelerating the sampling of the collective coordinates so as to sample $P_{\hat{\mathcal{O}}}(\mathcal{O}^*)$, including the low probability regions

- Extended (adiabatically separated) molecular dynamics
 - atomic degrees of freedom (\mathcal{X})
 - Extra degrees of freedom connected to the collective variables (\mathcal{Z})
 - Coupling potential term between $\mathcal X$ and $\mathcal Z$: $\frac{k}{2} \left(\mathcal O(x) z \right)^2$

$$\begin{cases} m\ddot{x} = -\nabla V(x) - k(\hat{\mathcal{O}}(x) - z)\nabla\hat{\mathcal{O}}(x) + thermo(T) \\ \mu \ddot{z} = -k(z - \hat{\mathcal{O}}(x)) + thermo(\bar{T}) \end{cases}$$



L. Maragliano and E. Vanden-Eijnden, Chem. Phys. Lett. 426 (2006), 168



TAMD: adiabaticity

- x are much faster than z
- \longrightarrow z moves according to the effective force

$$-\frac{1}{\tau} \int dt k(z - \mathcal{O}(x(t))) \stackrel{E.H.}{=} \frac{\int dx \ k(z - \mathcal{O}(x)) \exp\left[-\beta(V(x) + \frac{k}{2}(z - \mathcal{O}(x))^{2})\right]}{\int dx \ k \exp\left[-\beta(V(x) + \frac{k}{2}(z - \mathcal{O}(x))^{2})\right]}$$

(we have assumed that, apart for the $\mathcal{Z}\,$, the remaining degrees of freedom of the system are ergodic)







TAMD: the strong coupling limit

Interpretation of the effective force as mean force

$$\frac{\int dx \ k(z - \mathcal{O}(x)) \exp\left[-\beta(V(x) + \frac{k}{2}(z - \mathcal{O}(x))^2)\right] / \mathcal{Z}}{\mathcal{Z}_k(z) / \mathcal{Z}}$$

$$k \to \infty$$

$$\exp[-\beta \frac{k}{2} (\mathcal{O} - z)^2] \to \delta(\mathcal{O} - z)$$

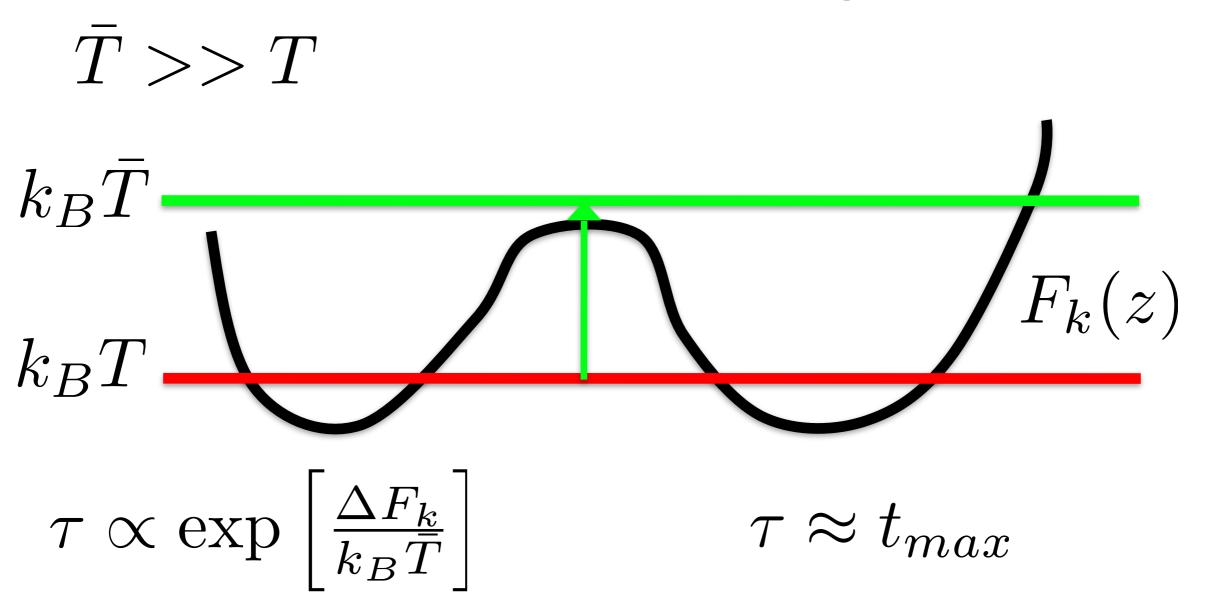
$$-\int dx \ k(\mathcal{O}(x)-z)\frac{P(x,z)}{P(z)} \qquad \frac{d}{dz}\beta^{-1}\ln\frac{\mathcal{Z}_k(z)}{\mathcal{Z}} \xrightarrow[k\to\infty]{} \frac{d}{dz}\beta^{-1}\ln\frac{\mathcal{Z}(z)}{\mathcal{Z}}$$







TAMD: collective variable at high temperature



Suitable interpolation algorithms (e.g. Single Sweep) can be used to reconstruct from these data the free energy surface







TAMC: the problem of non-analytical Collective Variables

• In TAMD nuclei evolve under the action of:

$$-\nabla V(x) - k(\mathcal{O}(x) - z)\nabla \mathcal{O}(x)$$

- TAMD (but also Metadynamics, Adiabatic Dynamics, ...) can be used only if the collective variable is an explicit-analytic function of the atomic positions
- In TAMC nuclei are evolved by MC instead than by MD according to the accelerated probability density function while the \mathcal{Z} 's are still evolved by MD. The adiabaticity conditions are easy to generalise so that we have a more powerful tool







Where is TAMC extension important?

- Classical cases
 - Nucleation
 - Rigorous collective variable to localize vacancies in solids
- Quantum cases: let the observable be the quantum average then $\mathcal{O}(x) = \langle \psi(r;x) | \hat{\mathcal{O}}(r,x) | \psi(r;x) \rangle$

$$\nabla \mathcal{O}(x) = <\nabla \psi(r;x)|\hat{\mathcal{O}}(r,x)|\psi(r;x)>$$

$$+ <\psi(r;x)|\hat{\mathcal{O}}(r,x)|\nabla \psi(r;x)>$$

$$+ <\psi(r;x)|\nabla \hat{\mathcal{O}}(r,x)|\psi(r;x)>$$

therefore for TAMD, and similar techniques, we need $\nabla \psi(r;x)$



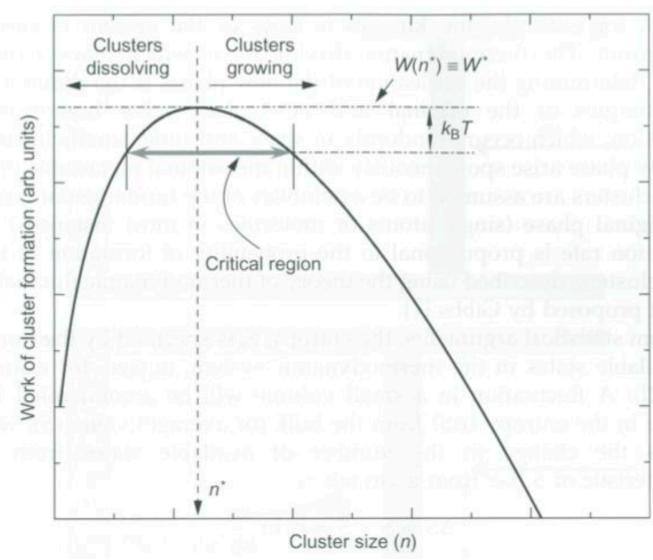




TAMC: application to the nucleation of a moderately undercooled L-J liquid

Targets

- Get the free energy as a function of the number of atoms of a given crystalline nucleus
- Critical size of the nucleus
- Mechanism of growth of the nucleus (hopefully)



Typical free energy as function of the number of atoms in the crystalline nucleus







Collective variable for nucleation

- Nucleus Size (NS): $\bar{\mathcal{N}}(x)$
 - Number of atoms in the largest cluster of (i) connected, (ii) crystal-like atoms
 - (i) Two atoms with $x_{ij} \leq 1.4 \ \sigma$ are connected when their q_{6m} are almost parallel¹

$$\frac{\left|\sum_{m=-6}^{6} q(i)_{lm} q(j)_{lm}\right|}{|q(i)_{lm}| |q(j)_{lm}|} \ge 0.5$$

- (ii) Crystal-like atoms: atoms with 7 or more connected atoms¹
- To identify the largest cluster one has to use methods of graph theory (e.g. the "Deep First search" which we used)

The NS is mathematically well defined but non analytical

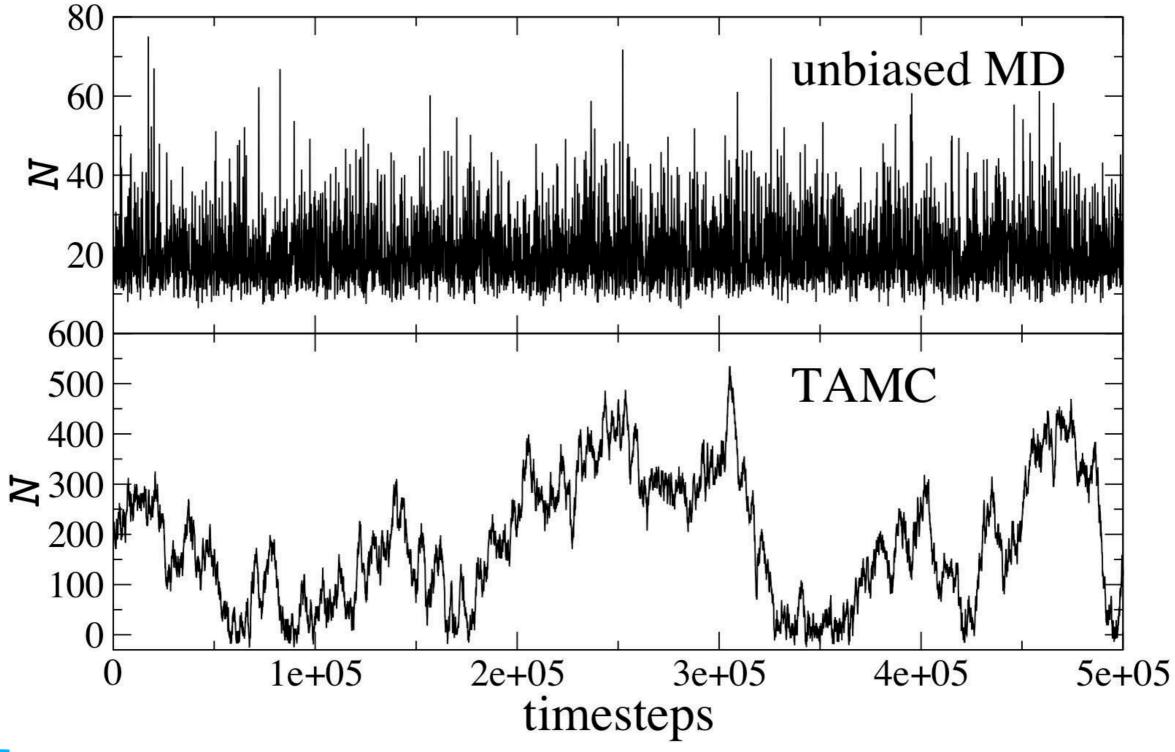
1) P. R. ten Wolde, M. J. Ruiz-Montero and D. Frenkel, J. Chem. Phys. 104 (1996) 9932







Results: timeline MD vs TAMC

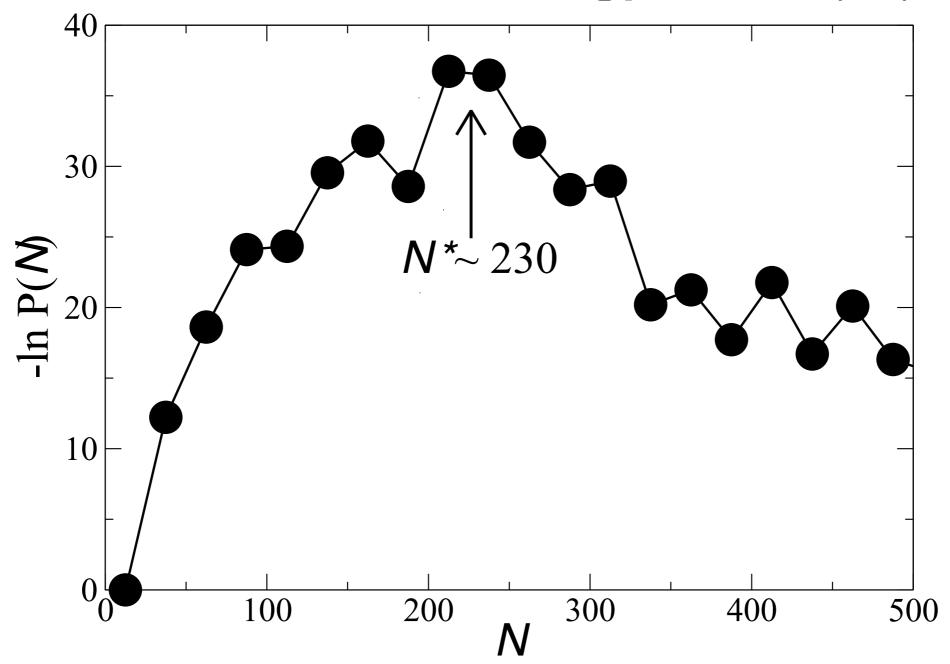








Results: free energy vs $\mathcal{N}(x)$



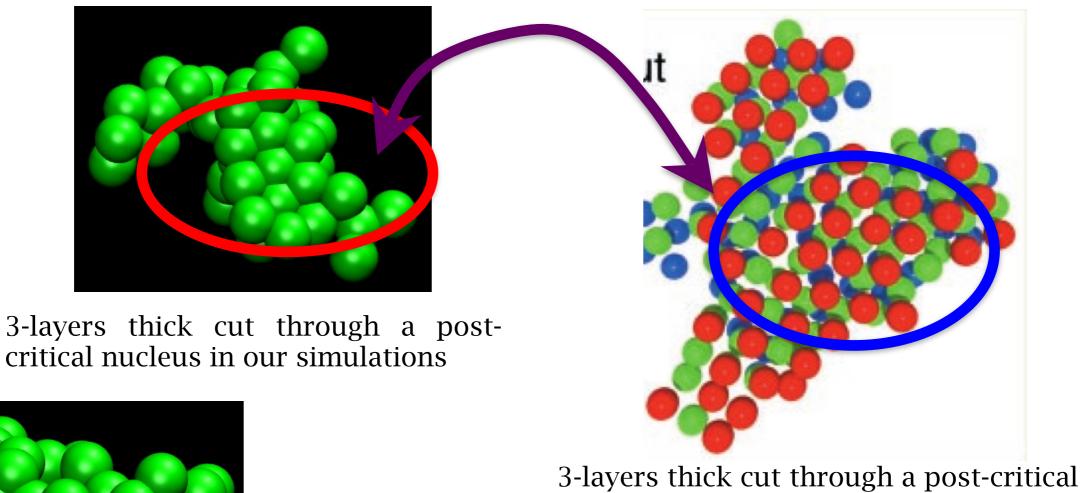
G. Ciccotti and S. Meloni, "Temperature Accelerated Monte Carlo (TAMC): a method for sampling the free energy surface of non-analytical collective coordinates" PCCP, 13, 5952 (2011)

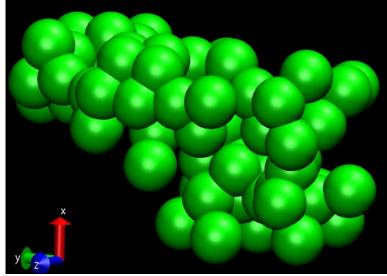






Results: nucleus configurations





an under-critical nucleus in our simulations



nucleus of colloids (by 3D imaging¹)







Non-Equilibrium MD (Dynamical-NEMD)

Hydrodynamic limit of $F_k(t)$

In books, after the derivation of the Navier-Stokes Eq.s (i.e. conservation laws (exact) + constitutive relations & local equilibrium hypothesis (phenomenological and approximate)) we are given

$$n_k(t) = f(k,t)$$
 - i.e an explicit form for $n_k(t)$

Then, is said, multiplying by $n_{-k}(0)$ and taking an "ensemble average", we get

$$F_k(t) = 1/N \ll n_k(t)n_{-k}(0) >>$$

BUT $n_k(t)$ is a macroscopic quantity, there is no ensemble over which to average. What kind of average is $\langle \langle \cdots \rangle \rangle$?







Hydrodynamic limit of $F_k(t)$

The question is: what is the meaning of the macroscopic $n_k(t)$ in Stat. Mech. and within the given approximations? Well,

$$n_k(t) = \langle \hat{n}_k \left(\Gamma(t) \right) | \hat{n}_{-k}(\Gamma) = n_{-k}(0) \rangle$$

i.e. is a standard conditional average over a suitable ensemble.

Multiplying by the condition and averaging over its probability distribution $P_{\hat{n}_{-k}}(n_{-k}(0))$ we get the $F_k(t)$: puzzle solved!

The question of this talk is: can we compute hydrodynamic fields, included $n_k(t)$, by some rigorous D-NEMD approach?







Relation between microscopic and macroscopic fields

How is defined the microscopic field $\hat{O}(\vec{x}, \Gamma)$ associated to the macroscopic field $O(\vec{x}, t)$?

$$\hat{O}(\vec{x}, \Gamma) = \sum_{i=1}^{N} \hat{\mathcal{O}}_i(\Gamma) \delta(\vec{x} - \vec{r}_i)$$

is the microscopic property associated to the field, $\mathcal{O}_i(\Gamma)$

e.g.:
$$n(\vec{x},t) \to \hat{\mathcal{O}}_i(\Gamma) = m_i, \ \vec{p}(\vec{x},t) \to \hat{\mathcal{O}}_i(\Gamma) = \vec{p}_i, \dots$$

The macroscopic field $O(\vec{x},t)$ is related to the corresponding microscopic field via a suitable average, conditional to the values of a set of scalar/field observables at t=0, $(\hat{C}(\vec{x},\Gamma) = C(\vec{x},t=0))$:

$$O(\vec{x}, t) = \langle \hat{O}(\vec{x}, \Gamma(t)) \rangle_{cond}$$







Dynamic NEMD [Ciccotti, Jacucci '75]

Compute the time-dependent NE average of a given observable

$$\frac{d\hat{O}}{dt} = -iL\hat{O}, \ \hat{O}(\vec{x}, t) = S(t)\hat{O}(\vec{x}, 0) \qquad iL = \{\mathcal{H}_0, \cdot\} + \{\mathcal{H}_p(t), \cdot\}$$

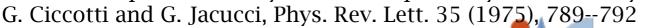
$$\frac{\partial \rho}{\partial t} = iL\rho, \quad \rho(\Gamma, t) = S^{\dagger}(t)\rho(\Gamma, 0)$$
Use the Onsager-Kubo equation:

$$O(\vec{x},t) = <\hat{O}(\vec{x},\Gamma)>_{t} = \int d\Gamma \hat{O}(\vec{x},\Gamma) \rho(\Gamma,t) \equiv \int d\Gamma \hat{O}(\vec{x},\Gamma) S^{\dagger}(t) \rho(\Gamma,0)$$

$$= \int d\Gamma S(t) \hat{O}(\vec{x},\Gamma) \rho(\Gamma,0) \equiv \langle S(t) \hat{O}(\vec{x},\Gamma) \rangle_{0}$$
Stationary, \mathcal{H}_{0} or otherwise driven, MD to sample $\rho(\Gamma,0)$

$$t \in \mathcal{H}_{0} + \mathcal{H}_{p} \text{ driven MD}$$
to compute $S(t) \hat{O}(\vec{x},\Gamma)$

"Direct Computation of Dynamical Response by Molecular Dynamics: The Mobility of a Charged Lennard-Jones Particle",





Conditional (initial) probability density function

$$\rho(\Gamma,0) = \rho\left(\Gamma|\hat{C}(\vec{x},\Gamma) = C(\vec{x},0)\right)$$

$$= \frac{\exp[-\beta H(\Gamma)] \prod_{\vec{x}} \delta\left(\hat{C}(\vec{x},\Gamma) - C(\vec{x},0)\right)}{\mathcal{Z}P_{\hat{C}(\vec{x})}(C(\vec{x},0))}$$

 Π indicates that the Dirac delta $\delta\left(\hat{C}(\vec{x},\Gamma) - C(\vec{x},0)\right)$ is valid over the entire ordinary space; \mathcal{Z} is the partition function and $P_{\hat{C}(\vec{x})}(C(\vec{x},0))$ denotes the probability to observe a value of the microscopic field $\hat{C}(\vec{x},\Gamma)$ equal to $C(\vec{x},0)$.

In practice, in simulations the fields are computed over a discretization of the ordinary space $\{\vec{x}_{\alpha}\}_{\alpha=1,M}$ and the $\delta(\vec{x}-\vec{r}_i)$ in their definition are mollified (e.g. replaced by gaussians) such that the field changes smoothly when particles move from one "cell" to another



Conditional averages by restrained MD

• The sampling of the conditional ensemble can be performed by restrained MD

$$<\hat{O}(\vec{x}_{\beta}, \Gamma(t))>_{cond} = \lim_{k \to \infty} \frac{\left\langle \hat{O}(\vec{x}_{\beta}, \Gamma(t)) \prod_{\vec{x}_{\alpha}} \exp\left[-\beta k/2 \left(\hat{C}(\vec{x}_{\alpha}, \Gamma) - C(\vec{x}_{\alpha})\right)^{2}\right] \right\rangle_{H}}{\left\langle \prod_{\vec{x}_{\alpha}} \exp\left[-\beta k/2 \left(\hat{C}(\vec{x}_{\alpha}, \Gamma) - C(\vec{x}_{\alpha})\right)^{2}\right] \right\rangle_{H}}$$

where: $=\lim_{k\to\infty} <\hat{O}(\vec{x}_{\beta},\Gamma(t))>_{H'}$

$$H'(\Gamma) = H(\Gamma) + \sum_{\vec{x}_{\alpha}} k/2 \left(\hat{C}(\vec{x}_{\alpha}, \Gamma) - C(\vec{x}_{\alpha}) \right)^{2}$$







Simulation Details

- 171500 particles: 88889 particles A, 82611 particles B
- Pair potential: $u^{AA}(r) = u^{BB}(r) = 4\varepsilon \left(\left(\frac{\sigma}{r} \right)^{12} \left(\frac{\sigma}{r} \right)^{6} \right)$ $u^{AB}(r) = 4\varepsilon \left(\frac{\sigma}{r} \right)^{12}$
- Simulation box: $\sim (90\sigma \times 45\sigma \times 45\sigma)$
- Average density: 1.024 particles $x \sigma^3$
- Temperature: 1.5 ε/k_b
- Simulation time:
 - Restrained MD: 75000 steps
 - Unrestrained MD: 600000 steps
- fields are averaged over (only) 40 unrestrained trajectories



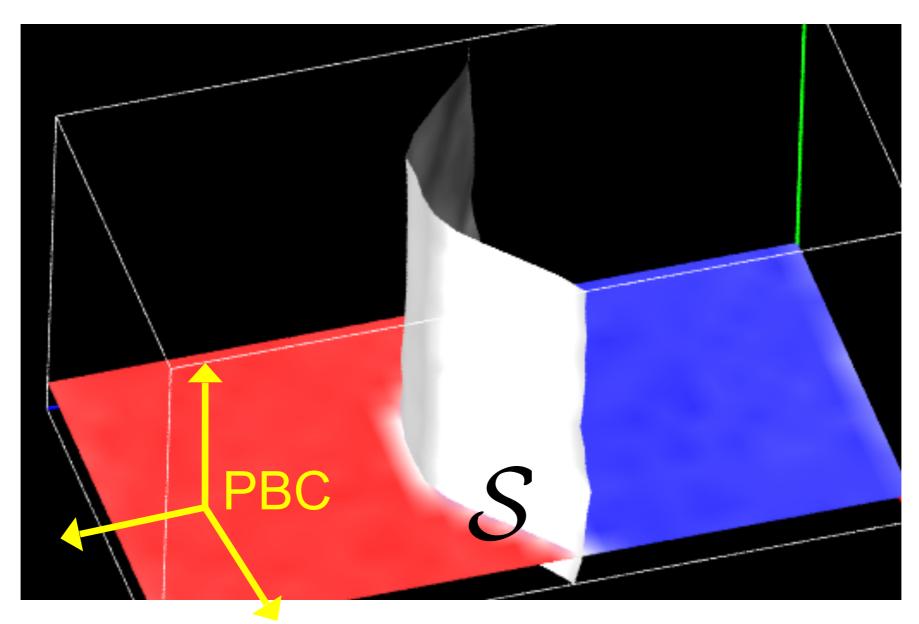






Hydrodynamic evolution of an interface by restrained- and NE-MD

$$\Delta n(\vec{x}_{\alpha}, \Gamma) = n^{A}(\vec{x}_{\alpha}, \Gamma) - n^{B}(\vec{x}_{\alpha}, \Gamma) = 0, \quad \vec{x}_{\alpha} \in S$$

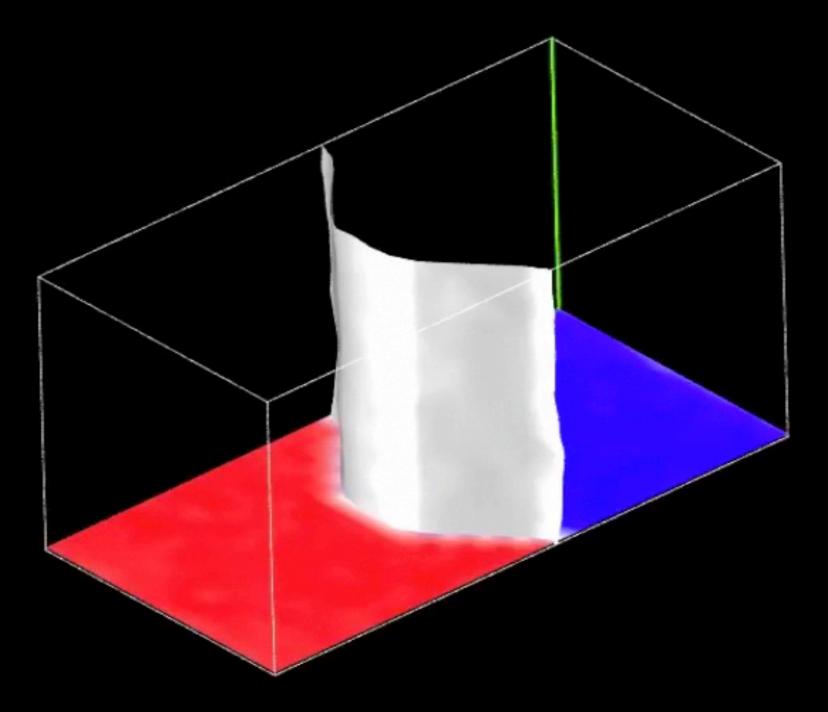


"Hydrodynamics from statistical mechanics: combined dynamical-NEMD and conditional sampling to relax an interface between two immiscible liquids.", S. Orlandini, S. Meloni, G. Ciccotti, Phys. Chem. Chem. Phys. 13, 13177 (2011)



Hydrodynamics from Dynamical Non-Equilibrium Molecular Dynamics

$$\Delta n(\vec{x}_{\alpha}, t) = 0$$



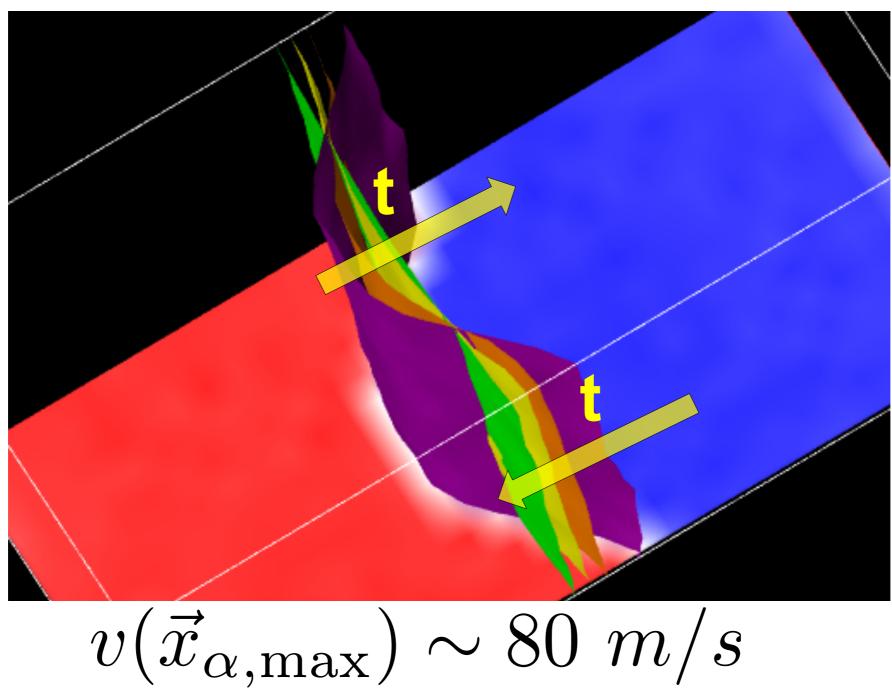


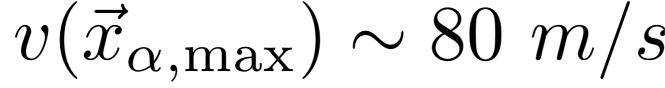




Hydrodynamics from Dynamical Non-Equilibrium Molecular Dynamics

$$\Delta n(\vec{x}_{\alpha}, t) = 0$$









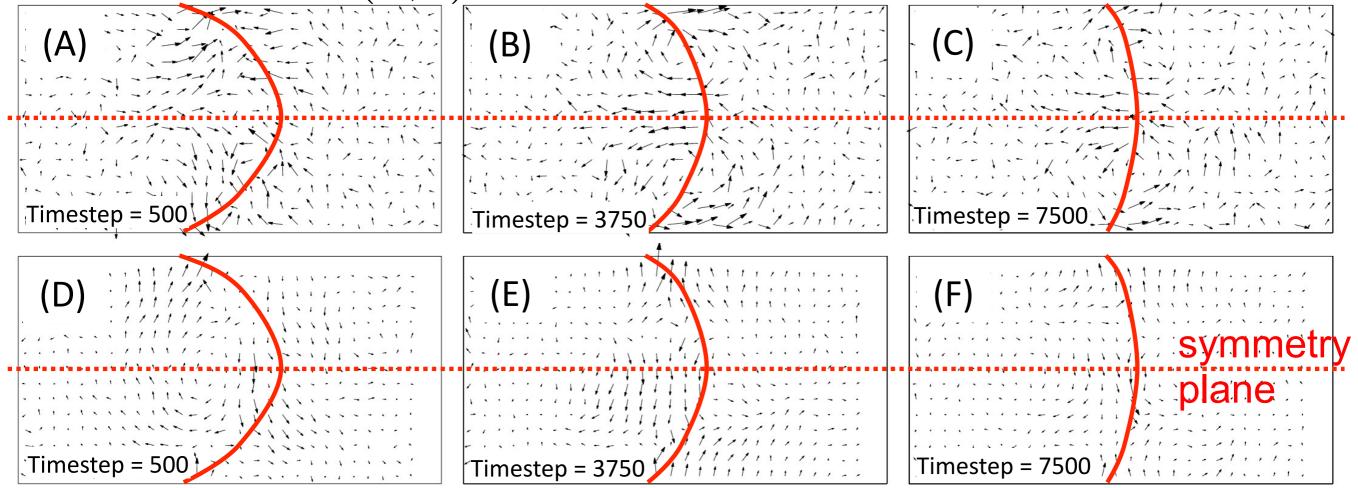


Hydrodynamics from Dynamical Non-Equilibrium Molecular Dynamics

Rigorous non-equilibrium ensemble averages vs local time averages (1)

$$\vec{v}(\vec{x},t) = \frac{\langle \hat{\vec{p}}(\vec{x},t) \rangle_{H'}}{n(\vec{x},t)}$$

The surface relaxes to the equilibrium by forming initially a two-tail profile of the velocity field that then stabilizes into a double-roll profile



$$\vec{v}(\vec{x},t) = \frac{1}{2\tau} \int_{t-\tau}^{t+\tau} ds \frac{\hat{\vec{p}}(\vec{x},s)}{n(\vec{x},t)}$$

The velocity field obtained via the local time average technique violates the symmetry of the problem





Conclusions

- ① Molecular Dynamics Simulations (MS) are a regular chapter of Theoretical Physics BUT they are amenable to real applications both for the analysis of experiments and for engineering
- 2 New algorithms have been, and continue to be, introduced to circumvent practical limitations of the MS, i.e. problems which cannot be solved by brute force. A good example has been TAMD/TAMC
- 3 Also NonEquilibrium situations can be confronted by a proper use of Onsager-Kubo relationship. This opens the way to represent directly transport but also to atomistic simulations (without constitutive hypotheses) of hydrodynamical phenomena.

Dynamical systems with (holonomic) constraints

$$\mathcal{L}\left(\{\mathbf{r},\dot{\mathbf{r}}\}\right) = \frac{1}{2} \sum_{i,\alpha} m_{i\alpha} \dot{\mathbf{r}}_{i\alpha}^2 - \mathcal{U}\left(\{\mathbf{r}\}\right) \qquad \left\{ \begin{aligned} i &= 1, N \text{ # molecules} \\ \alpha &= 1, n_i \end{aligned} \right. \\ \text{plus} \qquad \sigma_k\left(\{\mathbf{r}\}\right) = 0 \,, \qquad k = 1, f \text{ # constraints} \\ \text{e.g.} \qquad \sigma\left(\{\mathbf{r}\}\right) = \left(\mathbf{r}_{i\alpha} - \mathbf{r}_{i\beta}\right)^2 - d_{i,\alpha\beta}^2 = 0 \end{array}$$

To keep the trajectories satisfying the constraints $(\sigma(\{\mathbf{r}(t)\}) = 0, \forall t)$ we must have also $(\dot{\sigma} = \dot{\mathbf{r}} \cdot \nabla_r \sigma(\{\mathbf{r}(t)\}) = 0, \forall t)$

Moreover, if the constraint forces do not do work (i.e. conserve energy)

$$\dot{\mathbf{r}} \cdot \mathcal{G} \left(\{ \mathbf{r} \} \right) = 0, \quad \mathcal{G}_{i\alpha} = -\sum_{k=1}^{3} \lambda_k \nabla_{i\alpha} \sigma_k \equiv -\lambda \cdot \nabla_{i\alpha} \sigma$$

where the intensity λ of the constraint force has to be determined







SHAKE (1)

By differentiating two times the constraint relations σ 's we find

$$\ddot{\sigma}_k = \ddot{\mathbf{r}} \cdot \nabla_r \sigma_k + (\dot{\mathbf{r}}\dot{\mathbf{r}}) \cdot (\nabla_r \nabla_r) \, \sigma_k = 0$$
 (*)

and from
$$\frac{d}{dt}\frac{\partial \mathcal{L}}{\partial \dot{\mathbf{r}}} - \frac{\partial \mathcal{L}}{\partial \mathbf{r}} = \mathcal{G}$$
 i.e. $m\ddot{\mathbf{r}} = \mathbf{F} - \lambda \cdot (\nabla_r \sigma)$ (**)

$$\ddot{\sigma} = \frac{1}{m} \left(\mathbf{F} - (\lambda \nabla_r) \cdot \sigma \right) \cdot \nabla_r \sigma + (\dot{\mathbf{r}} \dot{\mathbf{r}} \cdot \nabla_r \nabla_r) \sigma = 0 \quad (***)$$

from which the <u>useless</u> solution of the linear system (***)

$$\lambda = \mathcal{Z}^{-1} \left[\left(\mathbf{F} \cdot \nabla_r \right) \sigma + \left(\dot{\mathbf{r}} \dot{\mathbf{r}} \cdot \nabla_r \nabla_r \right) \sigma \right], \qquad \mathcal{Z}_{k\ell} = \nabla_r \sigma_k \cdot \nabla_r \sigma_\ell$$

due to the problem of the algorithmic error, destroying the model.







SHAKE (2)

E.g.

$$\mathbf{r}(t+h) = \mathbf{r}(t) + h\dot{\mathbf{r}} + \frac{h^2}{2m} \left[\mathbf{F}(t) + \mathcal{G}(t) \right] + \mathcal{O}(h^3) \tag{+}$$

 $\mathcal{G}(t)$ exact at all timesteps will destroy in time the conservation of constraints

Instead, exploiting the "freedom" to choose the λ to satisfy the constraints one can write the set of equations

$$\sigma_k\left(\left\{\mathbf{r}(t+h)\right\}\right) = 0, \qquad k = 1, \dots, f \tag{++}$$

where, plugging e.g. (+) in (++) one gets λ values which satisfy exactly the constraints at timestep (t+h) without increasing the algorithmic error

(++) can be solved by any efficient algorithm. E.g. the Newton-Raphson method







SMwC (1)

Statistical Mechanics of systems with constraints in cartesian coordinates

In analytical mechanics, in presence of constraints

$$\mathbf{r} \longrightarrow \mathbf{r}(\mathbf{q})$$
 $\mathbf{r} : 3N ; \mathbf{q} : 3N - f$
 $\dot{\mathbf{r}} = \dot{\mathbf{q}} \cdot \nabla_{a} \mathbf{r}$

$$\mathcal{L}'\left(\mathbf{r},\dot{\mathbf{r}}\right)\longrightarrow\mathcal{L}'\left(\mathbf{q},\dot{\mathbf{q}}\right)\quad\Longrightarrow\quad\mathcal{H}\left(\mathbf{q},\mathbf{p}^{q}\right)\quad\left(\mathbf{p}^{q}=\frac{\partial\mathcal{L}}{\partial\dot{\mathbf{q}}}\right)$$

and, for Statistical Mechanics

$$Q = \int d\mathbf{q} d\mathbf{p}^q e^{-\beta \mathcal{H}(\mathbf{q}, \mathbf{p}^q)}$$

Now: instead of assuming the constraints satisfied, let us take the constraint functions as f generalized coordinates. Then

$$\mathbf{r} \longleftrightarrow ((\mathbf{q}, \, \sigma) \equiv \mathbf{u}) \quad \mathbf{p}^r \longleftrightarrow (\mathbf{p}^u \equiv (\mathbf{p}^q, \, \mathbf{p}^\sigma)) \text{ with } \mathbf{p}^u = \mathbf{M}\dot{\mathbf{u}}$$

The constrained system will be obtained putting back

$$\sigma(\mathbf{r}) = 0, \quad \dot{\sigma}(\mathbf{r}, \dot{\mathbf{r}}) = 0$$







SMwC (2)

Considering that

$$\mathcal{L}(\mathbf{r}, \dot{\mathbf{r}}) = \mathcal{K}(\dot{\mathbf{r}}) - \mathcal{V}(\mathbf{r}) = \mathcal{L}(\mathbf{u}, \dot{\mathbf{u}}) = \frac{1}{2}\dot{\mathbf{u}}^T\mathbf{M}\dot{\mathbf{u}} - \mathcal{V}'(\mathbf{u})$$

$$\mathbf{p}^{u} = \begin{pmatrix} \mathbf{p}^{q} \\ \mathbf{p}^{\sigma} \end{pmatrix} = \frac{\partial \mathcal{L}'}{\partial \dot{\mathbf{u}}} = \mathbf{M}\dot{\mathbf{u}} = \begin{pmatrix} \mathbf{A} & \mathbf{B} \\ \mathbf{B}^{T} & \Gamma \end{pmatrix} \begin{pmatrix} \dot{\mathbf{q}} \\ \dot{\sigma} \end{pmatrix}$$

i.e.
$$\dot{\mathbf{u}} = \begin{pmatrix} \dot{\mathbf{q}} \\ \dot{\sigma} \end{pmatrix} = \mathbf{M}^{-1} \mathbf{p}^u = \begin{pmatrix} \Delta & \mathbf{E} \\ \mathbf{E}^T & \mathbf{Z} \end{pmatrix} \begin{pmatrix} \mathbf{p}^q \\ \mathbf{p}^\sigma \end{pmatrix}$$

we have that

$$\dot{\sigma} = 0$$

corresponds to

$$\dot{\sigma} = \mathbf{E}^T \mathbf{p}^q + \mathbf{Z} \mathbf{p}^\sigma = 0$$

i.e. to \mathbf{p}^{σ} not zero, for $\dot{\sigma}=0$, but equal to $\tilde{\mathbf{p}}^{\sigma}=\tilde{\mathbf{Z}}^{-1}\tilde{\mathbf{E}}^T\mathbf{p}^q$ and $\mathbf{p}^{\sigma}+\tilde{\mathbf{p}}^{\sigma}=\tilde{\mathbf{Z}}^{-1}\dot{\sigma}$







SMwC(3)

Now our result is at hand, since

$$\mathcal{L}'(\mathbf{u}, \dot{\mathbf{u}}) \longleftrightarrow \mathcal{H}'(\mathbf{p}^u, \mathbf{u})$$

$$\mathcal{H}_c(\mathbf{p}^q, \mathbf{q}) = \mathcal{H}'(\mathbf{p}^q, \mathbf{p}^\sigma = \mathbf{Z}^{-1} \mathbf{E}^T \mathbf{p}^q, \mathbf{q}, \mathbf{q}, \sigma = 0)$$

SO

$$Q = \int d\mathbf{q} d\mathbf{p}^{q} e^{-\beta \mathcal{H}_{c}} = \int d\mathbf{u} d\mathbf{p}^{u} e^{-\beta \mathcal{H}'(\mathbf{p}^{u}, \mathbf{u})} \delta\left(\sigma\right) \delta\left(\mathbf{p}^{\sigma} - \tilde{\mathbf{p}}^{\sigma}\right)$$
$$= \int d\mathbf{r} d\mathbf{p}^{r} e^{-\beta \mathcal{H}(\mathbf{p}^{r}, \mathbf{r})} \delta\left(\sigma(\mathbf{r})\right) \delta\left(\mathbf{Z}^{-1} \dot{\sigma}(\mathbf{r}, \mathbf{p}^{r})\right)$$

where $d\mathbf{u}d\mathbf{p}^u = d\mathbf{r}d\mathbf{p}^r$ since the transformation is canonical and the Jacobian is = 1

Moreover
$$\varrho(\mathbf{r}, \mathbf{p}^r) = \frac{1}{\mathcal{Q}} e^{-\beta \mathcal{H}(\mathbf{p}^r, \mathbf{r})} \delta\left(\sigma(\mathbf{r})\right) \delta\left(\mathbf{Z}^{-1} \dot{\sigma}(\mathbf{r}, \mathbf{p}^r)\right)$$

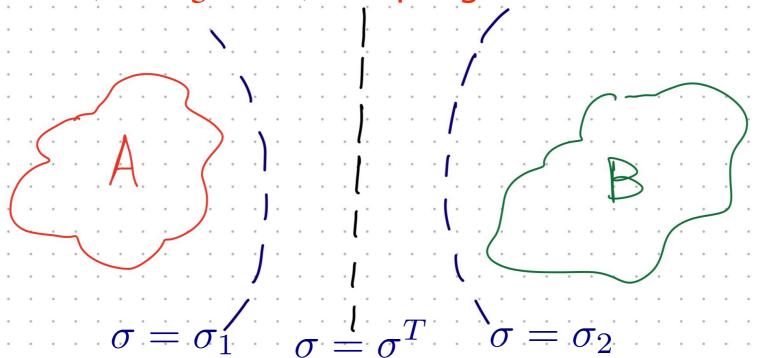






ASBM(1)

Accelerated (but rigorous) Sampling: Blue Moon Ensemble



We assume a metastability in configurational space and the existence of at least one configurational function (also often improperly called a reaction coordinate) such that its values going from A to B grow monotonically from σ_A to $\sigma_B > \sigma_A$ and can then be used to characterise the STATE of the system at and between the two metastable states. The system spends the majority of its time in A or B and very little in between. The transitional region has almost zero probability to be visited and MD is unable to sample the full space







ASBM(2)

We would like to know $\mathcal{P}_{\sigma}\left(\sigma'\right)$ for all possible values (probable and improbable) of σ' .

1. using $\sigma(\{\mathbf{r}\})$ as a constraint we can sample by MD any chosen region

2. By knowing the probability in presence of a constraint (as shown before) we can demonstrate that

(X)
$$\frac{d \ln \mathcal{P}_{\Xi}(\xi')}{d\xi'} \propto \langle \mathbf{F}_{\xi'} \rangle_{cond} = \frac{\langle corr \, \mathbf{F}_{\xi'} \rangle_{constrained}}{\langle corr \rangle_{constrained}}$$

where the second term is virtually impossible to sample while the third is obtained with full statistics.

3. the integral of (X) [Thermodynamical Integration] provides a direct measure of the strength of the metastability





