

Molecular equilibrium sampling and free energy computations: some open topics

Mathias Rousset⁽¹⁾

(1) INRIA Lille-Nord Europe and Laboratoire Paul Painlevé,
SIMPAF project-team, Villeneuve d'Ascq, France.

- Joint work with: [T. Lelièvre](#) and [G. Stoltz](#) (MicMac INRIA project team, ENPC, Marne-la-Vallée, France). Talk mainly based on ideas out of:
 - 11' T. Lelièvre, MR and G. Stoltz: Langevin dynamics with constraints and computation of free energy differences.
 - 10' T. Lelièvre, MR and G. Stoltz: Free energy computation: a mathematical perspective.
- In this talk, I will focus on:
[Errors in equilibrium sampling](#) and free energy computation of [molecular systems](#) coupled to a thermostat (NVT = fixed number of particles, volume and temperature).

- Errors is a **puzzling** aspect of the field: everyone computes but no one knows **how far** we are from **"reality"**.
- In this talk I will **NOT** focus on:
 - Extraction of **dynamical quantities** (exit times, Transition State Theory, etc..)
 - Computation of **non-equilibrium steady states**.
 - Errors in computation of (effective) potentials **quantum chemistry**.

- There are **still open questions** from the mathematical/theoretical side.
- In particular some source of **errors remained uncontrolled** , e.g.:
 - **Time-step** errors.
 - **Quantum** corrections.
 - Model **reduction**.
- Important in **practice**:
 - To track the origin of **errors/aberrations** in simulations.
 - To increase **efficiency** by focusing on the **bottlenecks errors**.

OVERVIEW ON MOLECULAR SAMPLING AND FREE ENERGY

REALITY HAS AN EQUATION !!. N = number of atoms. Globally neutral.

- Building block: **Full Schrödinger** operator (q position of nuclei, x position of electrons, Z_i = number of protons in nucleus i , N_i = number of neutrons in nucleus i) in atomic units:

$$\left\{ \begin{array}{l} \epsilon^2 = \frac{m_e}{m_p} \sim 10^{-3}, \\ H_n = - \sum_{i \in \text{nuc}} \frac{\epsilon^2}{2(Z_i + N_i)} \Delta_{q_i} + \sum_{i \in \text{nuc}, j \in \text{nuc}} \frac{Z_i Z_j}{|q_j - q_i|}, \\ H_e = - \frac{1}{2} \Delta_x + \sum_{i \in e, j \in e} \frac{1}{|x_j - x_i|}, \\ V_c(x, q) = - \sum_{i \in \text{nuc}, j \in e} \frac{Z_i}{|x_j - q_i|}, \\ H_{tot} = H_p + H_e + V_c \times, \end{array} \right.$$

- You need to add **spin variable** to electrons and **skew-symmetry** constraint.
- Schrödinger equation: $i\varepsilon\partial_t\psi = H_{tot}\psi$.
- Usual temperature, in atomic unit, is small: $\beta \sim 10^3$.

- Standard route: the **Born-Oppenheimer approximation**:
 - Electrons are **fast (in fact stationary)** and nuclei are **slow**.
 - So that you **neglect coupling between eigenstates** of the electrons (adiabatic assumption).
 - You get the decoupled equations for $k = 0 \dots \infty$:

$$\begin{cases} i\epsilon\partial_t\Psi_n = H_n\Psi_n + E_e^k(q) \times \Psi_n, \\ (H_e + V_c(\cdot, q))\Psi_{e,q} = E_e^k(q)\Psi_{e,q} \quad (k\text{-th eigenvalue problem}). \end{cases}$$

- Taking $\varepsilon \rightarrow 0$ (semi-classical limit), and $k = 0$ (groundstate), we get ab-initio classical dynamics:

$$\begin{cases} Z_i \ddot{q}_i = -\nabla_{q_i} E_e^0(q), & i = 1 \dots N. \\ (H_e + V_c(\cdot, q)) \Psi_{e,q} = E_e^0(q) \Psi_{e,q} & \text{(eigenvalue/minimization problem)}. \end{cases}$$

- It is the standard **ab-initio** model for MD.

- In many cases effective potentials are used with 2,3,4- body effective potentials.
- Even if untrue in practice, we assume that we have a standard matching procedure (MP) to compute effective potentials n -body potentials from any N -body potential:

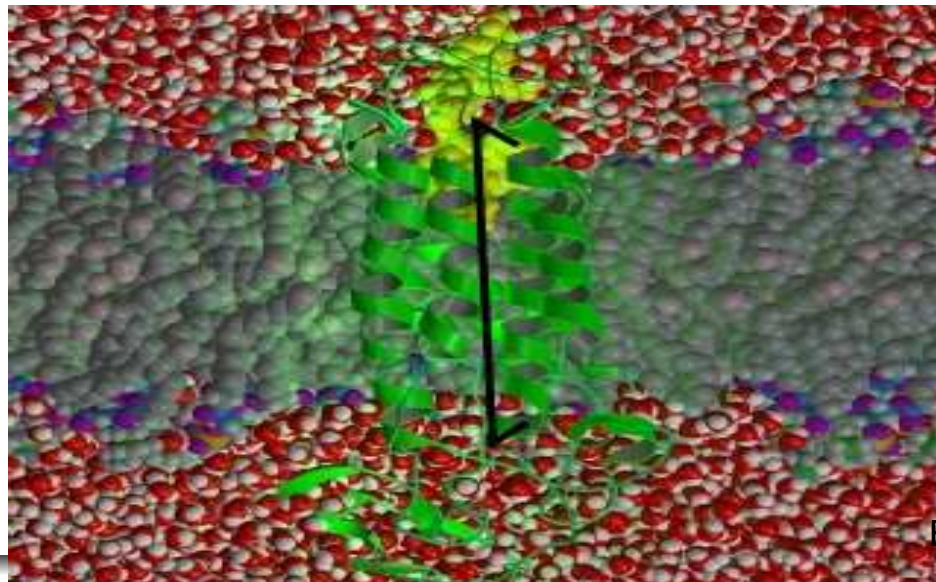
$$\text{molecular system} \xrightarrow{MP} (V_n(q_{i_1}, \dots, q_{i_n}), i_1 \dots i_n \in C_N^n, n = 2 \dots N)$$

- In principle: independant of $\varepsilon, \beta, \Delta t$. Computed from idealized quantum calculation, not by fitting parameters to reality !!
- This step requires a lot of quantum chemistry, and sometimes kitchen/crystal ball science (CHARMM force field).
- Other topic: Coarse-graining = model reduction in ${}^6\tilde{N}$ models ($\tilde{N} < N$ meta-atoms).

- Short range interaction (covalent): Rapidly oscillating quantity ($\sim 10^{-15}$ s).
- Large range interaction (electrostatic): ($\sim 10^{-12}$ s).
- **Reaction coordinates** (slow macroscopic variables $\sim 10^{-9}$ s, limit of computational range):

$$\xi_{\text{slow}} : \mathbb{R}^{3N} \rightarrow \mathbb{R}$$

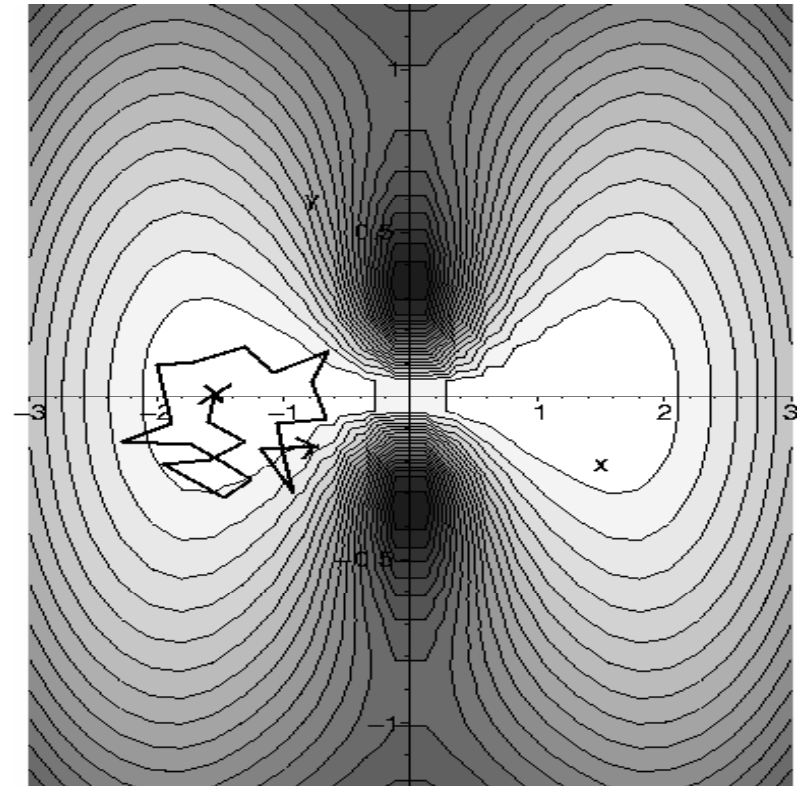
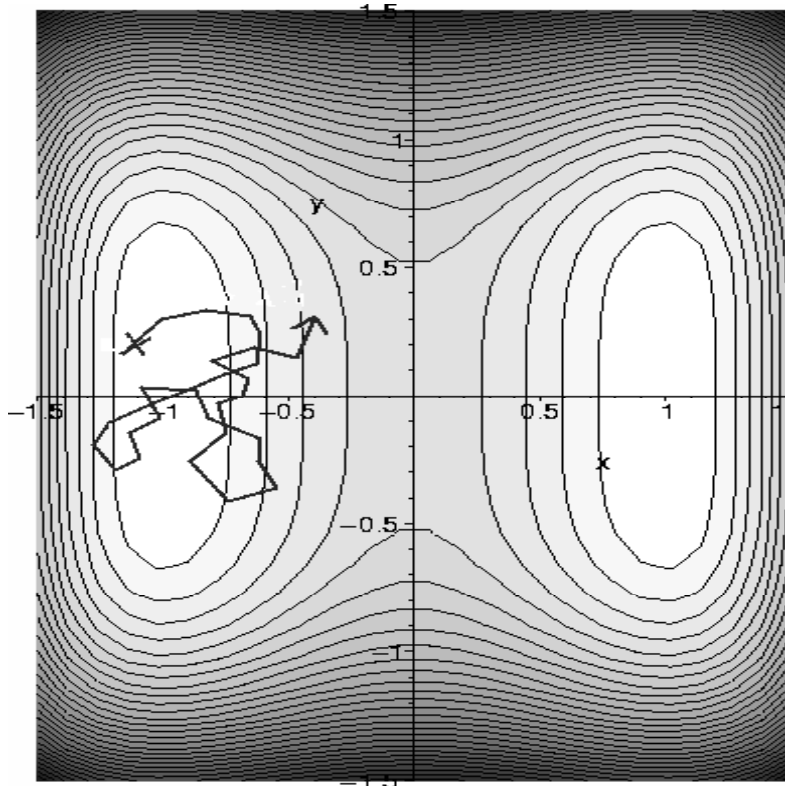
Example:



Slow degrees of freedom: Meta-stability

Dynamical **barriers** = **metastability** = exponential slowdown of equilibrium convergence:

- **Energetic** (low temperature) or **entropic** (high temperature) barriers:



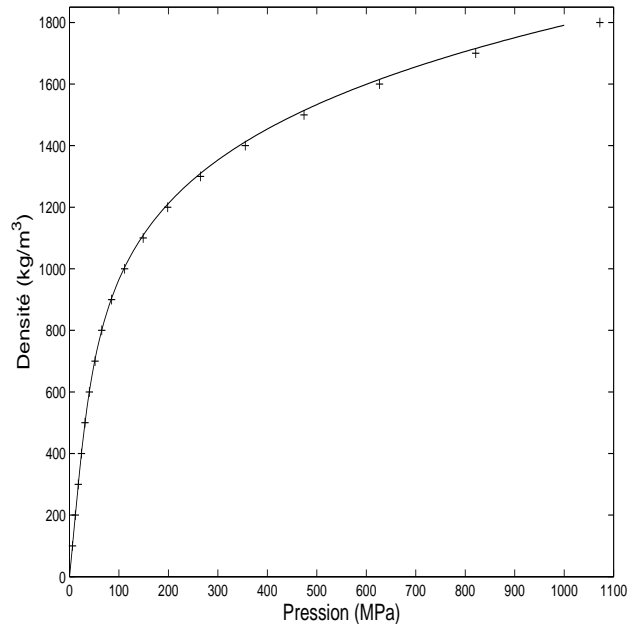
- Introduction of a coupling with a thermostat of **temperature**, $\beta^{-1} = k_b T$.
- Ergodic assumption:

$$\begin{array}{ccc} \text{Time average} & = & \text{Spatial average} \\ \frac{1}{T} \int_0^T \varphi(q(t)) dt & \xrightarrow{T \rightarrow \infty} & \langle \varphi \rangle \end{array}$$

- Typical configurations given by **Boltzmann Law** (NVT case):

$$\langle \varphi \rangle = \frac{\int \varphi(q) e^{-\beta V(q)} dq}{\int e^{-\beta V(q)} dq}$$

- **Statistical** nature of quantities to be computed:
- Boltzmann gives all **equilibrium physical quantities**. Ex: pressure
 $\propto E_c - 2\langle q \cdot \nabla V(q) \rangle$.
- **Non-ideal equation of state** of Argon at 300 K:



What is free energy F ?

- Free energy is defined by the **equilibrium (marginal)** distribution of $\xi_{\text{slow}} \in \mathbb{R}^d$, a "slow" variable, or reaction coordinates. For any observable φ of ξ_{slow}

$$\langle \varphi(\xi_{\text{slow}}(q)) \rangle = \frac{\int_{\mathbb{R}^d} \varphi(z) e^{-\beta F(z)} dz}{Z}$$

- Define $\mathcal{M}_z = \{q \in \mathbb{R}^d \mid \xi_{\text{slow}}(q) = z\}$ the sub-manifold associated with the value z of the reaction coordinate ξ_{slow} .
- Define $\delta_{\xi_{\text{slow}}(q)=z}(dq)$ the **conditional surface measure on \mathcal{M}_z** verifying the slice integration $dq = \delta_{\xi_{\text{slow}}(q)=z}(dq) dz$

What is free energy F ?

- We get, up to an additive constant:

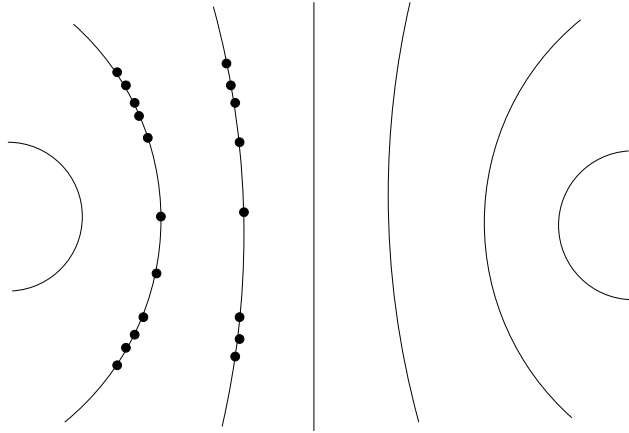
$$F(z) = -\beta^{-1} \ln \int_{\mathcal{M}_z} e^{-\beta V(q)} \delta_{\xi_{\text{slow}}(q)=z}(dq)$$

- Goal: numerical computation of $z \mapsto F(z)$

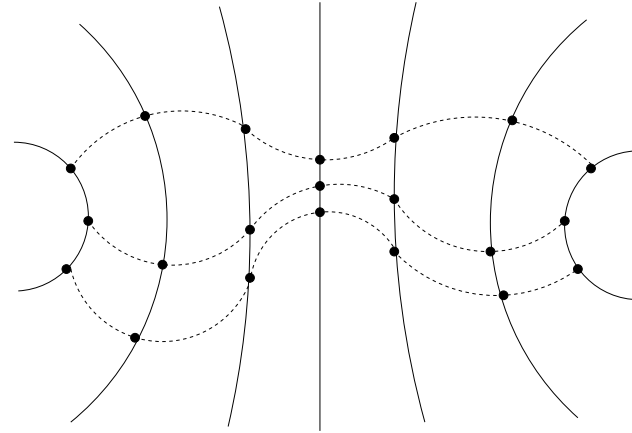
- In fact in the context of the present talk ([error analysis](#)), free energy computation is arguably [reducible to the sampling problem](#).
- We need to add [non-equilibrium](#) simulation of [Jarzynski](#) type. But, I arguably the discussion on errors is strictly [similar](#) to the case of sampling.
- Two main class of methods:
 - [Adaptive biasing](#) methods.
 - [Non-equilibrium simulation of Jarzynski](#) type (Thermodynamic Integration as a limit).

The enemy is **metastability**.

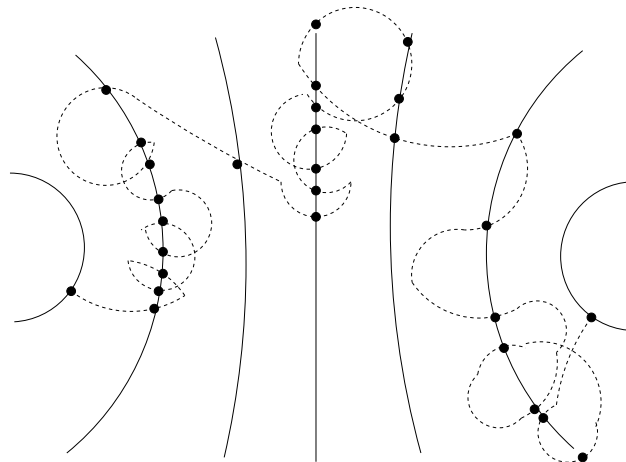
Fixed constraint



Switching constraint



Biasing (with or without adaptivity)



- Compute adaptively a biasing potential of the form:

$$F_{\text{approx}}(\xi_{\text{slow}}(q))$$

- Perform sampling with the biasing potential F_{approx} .
- The bias is removable since F_{approx} and the equilibrium distribution is known.

- Add a constraining force (switching) to a molecular system of the form $-\nabla_q \xi_{\text{slow}} \lambda(t)$, where $\lambda(t)$ is a Lagrange multiplier with constraints $\xi_{\text{slow}}(q_t) = z(t)$.
- Take care with the thermostat.
- Starts at equilibrium .
- Compute the energy variation of the system which is only due to the switching (denoted $\mathcal{W}_T =$ the work).
- Use the identity: $F(z(T)) - F(z(0)) = -\beta \ln \mathbb{E}(e^{-\beta(\mathcal{W}_{0,T} + \text{Corr}_T - \text{Corr}_0)})$.
- Requires several replicas to average \mathbb{E} .

ERRORS

- **Quantum** errors:
 - Error in adiabatic decoupling. ("conical crossings"+ ϵ, β , See later)
 - Non-classical behavior of nuclei. (ϵ, β , See later)
- **Model reduction** errors:
 - 1, 2, 3-body effective potentials. (Really a mess).
 - "Constraints" errors (due to "freezing" fast degrees of freedom). (See later).
 - Coarse-graining. (Really model dependent).
- Errors in **time-stepping**
 - Time-step errors. (Δt , See later)
 - Bonus: efficiency of the sampling method. (See later)

ERRORS FROM TIME-STEP AND SAMPLING EFFICIENCY

- Here: **Newton equation** with **stochastic thermostat**. V is assumed to be a known classical potential. N -body system.
- Equilibrium distribution: $NVT \propto e^{-\beta V(q)} dq$ for positions.

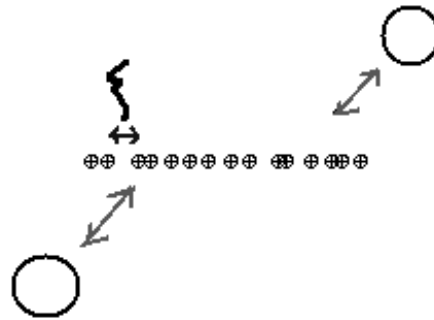
- Model: a **stiff** Hamiltonian system $H : \mathbb{R}^{6N} \rightarrow \mathbb{R}$:

$$H(p, q) = \frac{1}{2} p^T M^{-1} p + V(q)$$

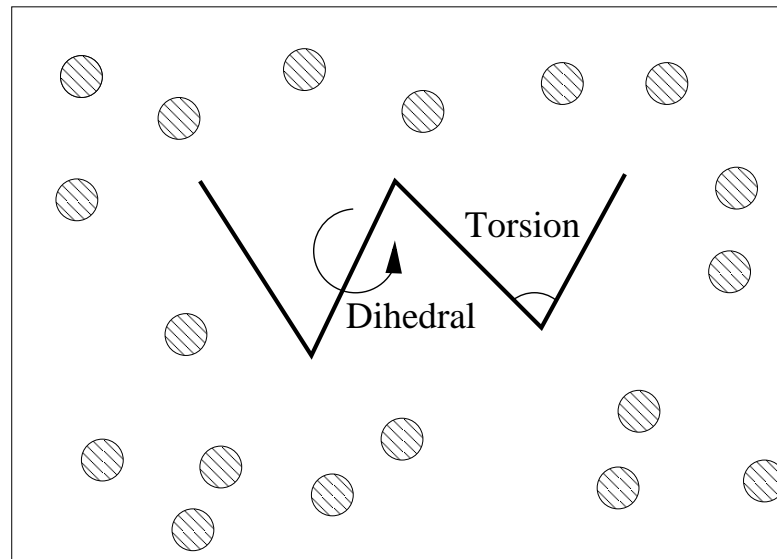
- $M = \text{diag}(m_1, \dots, m_N)$.
- Assume: **fastest** degrees of freedom (fDOFs) are known

$$\begin{cases} \xi_{\text{fast}} : \mathbb{R}^{3N} \rightarrow \mathbb{R}^n, \\ \{q \in \mathbb{R}^{3N} \mid \xi_{\text{fast}}(q) = 0\} = \text{"slow manifold"}. \end{cases}$$

- Ex.: Large molecule + slow environment:



- ξ_{fast} can be: bond lengths, bond angles, dihedral angles .



- Coupling with a **thermostat**: Newton/Hamiltonian motion + Ornstein-Uhlenbeck process.

$$\left\{ \begin{array}{l} dQ_t = M^{-1} P_t dt \\ dP_t = \underbrace{-\nabla V(Q_t)}_{\text{Hamilton}} - \underbrace{\gamma M^{-1} P_t dt}_{\text{Dissipation}} + \underbrace{\sigma dW_t}_{\text{Fluctuation}} \end{array} \right. \quad \sigma \sigma^T = \frac{2\gamma}{\beta} = 2\gamma k_b T$$

- Equilibrium distribution: $\frac{1}{Z} e^{-\beta H(p,q)}$.

- Newton/Hamilton $H(q, p) = \frac{1}{2}p^T M^{-1}p + V(q) = \text{kin.} + \text{pot. energy} :$

$$\begin{cases} \frac{d}{dt} \begin{pmatrix} q_t \\ p_t \end{pmatrix} = J \nabla_{q,p} H(q_t, p_t) & (1) \\ J := \begin{pmatrix} 0 & Id \\ -Id & 0 \end{pmatrix} \end{cases}$$

- $J =$ symplectic matrix:
 - Conservation of phase-space measure $dqdp$ (Liouville theorem = int. by parts).
 - Conservation of energy H .
- Reversibility $t \mapsto (q_{T-t}, -p_{T-t})$ is still solution of (1).

- Ornstein-Uhlenbeck process on momenta (stochastic thermostat):

$$dP_t = -\gamma M^{-1} P_t + \sigma dW_t, \quad \sigma \sigma^T = \frac{2\gamma}{\beta}, \quad (1)$$

- Symmetry: $t \mapsto -P_t$ solution of (1).
- Reversibility wrt to equilibrium distribution: If

$$\text{Law}(P_0) = \frac{1}{Z} e^{-\beta \frac{1}{2} p^T M^{-1} p} dp$$

then the probability distribution of paths is invariant under time reversal:

$$\text{Law}(t \mapsto P_t, t \in [0, T]) = \text{Law}(t \mapsto \pm P_{T-t}, t \in [0, T])$$

- Ergodicity, whatever P_0 :

$$\lim_{T \rightarrow +\infty} \frac{1}{T} \int_0^T \varphi(P_t) dt = \int_{\mathbb{R}^{3N}} \varphi(p) \frac{1}{Z_k} e^{-\beta \frac{1}{2} p^T M^{-1} p} dp \quad a.s.$$

- Mixing, whatever P_0 :

$$\lim_{t \rightarrow +\infty} Law(P_t) \xrightarrow{\text{exp}} \frac{1}{Z_k} e^{-\beta \frac{1}{2} p^T M^{-1} p} dp.$$

- Reversibility up to **momenta reversal** wrt to invariant distribution: If

$$\text{Law}(Q_0, P_0) = \frac{1}{Z} e^{-\beta H} dq dp$$

then the probability distribution of paths is invariant under time reversal:

$$\text{Law}(t \mapsto (Q_t, P_t), t \in [0, T]) = \text{Law}(t \mapsto (Q_{T-t}, -P_{T-t}), t \in [0, T])$$

THIS IS DETAILED BALANCE

- Ergodicity + mixing from the hypoellipticity of associated operator:

$$\lim_{T \rightarrow +\infty} \frac{1}{T} \int_0^T \varphi(Q_t, P_t) dt = \int_{\mathbb{R}^{6N}} \varphi(q, p) \frac{1}{Z} e^{-\beta H(q, p)} dq dp \quad a.s.$$

- Hamiltonian dynamics: Verlet **explicit** integration.

$$\left\{ \begin{array}{l} p^{n+1/2} = p^n - \frac{\Delta t}{2} \nabla V(q^n), \\ q^{n+1} = q^n + \Delta t M^{-1} p^{n+1/2}, \\ p^{n+1} = p^{n+1/2} - \frac{\Delta t}{2} \nabla V(q^{n+1}) \end{array} \right.$$

- **Conserve qualitative properties of continuous Hamiltonian dynamics:**
 - **Time-reversible** under $p \rightarrow -p$.
 - Conserve **symplectic form** ($2d$ - area in all planes) and thus volume $d p d q$.
 - Quasi-conservation of energy $H_{\Delta t, n} = H + O(\Delta t^2 \times \log(\Delta t n))$
 - Does **not work** if H is not separable $H = T(p) + V(q)$.

- One can thus get **detailed balance** up to an energy drift, that is to say:

$$\begin{aligned}
 & e^{-\beta H(q_n, p_n)} \underbrace{\delta \Phi(q_n, p_n)}_{\text{numerical flow}}(dq_{n+1} dp_{n+1}) dq_n dp_n = \\
 & \underbrace{e^{-\beta(H(q_n, p_n) - H(q_{n+1}, p_{n+1}))}}_{\text{energy variation}} \times \\
 & e^{-\beta(H(q_{n+1}, p_{n+1}))} \delta \Phi(q_{n+1}, p_{n+1})(dq_n dp_n) dq_{n+1} dp_{n+1},
 \end{aligned}$$

- Versions when adding a Orstein-Uhlenbeck thermostat process.

- GHMC = Metropolis for **time-step errors of Langevin equations**.
 1. Evolve **momenta** according to the **Ornstein-Uhlenbeck process** during Δt compute the energy $H(q^n, p^{n+1/4})$.
 2. Integrate the **Hamiltonian equations** of motion according to the Verlet scheme. Get $(\tilde{q}^{n+1}, \tilde{p}^{n+1})$ and compute the energy $H(\tilde{q}^{n+1}, \tilde{p}^{n+1})$.
 3. Accept the proposal with probability

$$\min(1, \exp(-\beta(H(\tilde{q}^{n+1}, \tilde{p}^{n+1}) - H(q^n, p^{n+1/4}))))$$

otherwise, reject and **reverse momenta** by setting

$$(q^{n+1}, p^{n+1}) = (q^n, -p^{n+1/4});$$

- Ergodic Markov chain **reversible wrt exact stationary distribution**
 $e^{-\beta H(q,p)} dq dp$ up to $p \rightarrow -p$ (**detailed balance**).

Overdamped dynamics:

- Replace step (1) by immediate equilibrium (**high friction limit**).
- Verlet yields **Euler discretized overdamped stochastic differential equation**:

$$Q^{n+1} = Q^n - \Delta t^2 / 2M^{-1} \underbrace{\nabla V(Q^n)}_{drift} + \Delta t \sqrt{\beta^{-1} M^{-1}} \underbrace{\mathcal{N}(0, 1)}_{diffusion}$$

- Yields Metropolis algorithm when add the rejection rule.
- Yields a Markov chain reversible wrt $e^{-\beta V(q)} dq$.

- The **average rejection rate** in Metropolis/ Hybrid Monte Carlo (both Langevin and overdamped) ONLY depend the time-step and NOT on the friction. This enables to define a **critical time step** with **rejection prescribed at 1/50** valid independantly on the friction parameter.
- However, if we denote the **mixing time of Langevin dynamics** T_{mix} , the computational cost of discretized Langevin without Metropolis step is:

$$N_{mix} = T_{mix} / \Delta t_{crit}.$$

- Now denote the **mixing "time"** $T_{mix,ov}^2$ of the overdamped dynamics:

$$dQ_s = -M^{-1}\nabla_q V ds + \sqrt{2\beta^{-1}M^{-1}}dW_s.$$

$T_{mix,ov}^2$ is homogenous to $[T^2]$ but **independent of Δt_{crit}** . Then according to the effective time step of overdamped above:

$$N_{mix,ov} = 2T_{mix,ov}^2 / \Delta t_{crit}^2.$$

- Thus if $\Delta t_{crit} \ll T_{mix,ov}$ then:

$$\frac{N_{mix}}{N_{mix,ov}} = \frac{\Delta t_{crit}}{T_{mix,ov}} \frac{T_{mix}}{T_{mix,ov}} \ll 1$$

- Langevin is much better than overdamped (**OPEN**: to precise)..
- **BUT Momenta flip in GHMC** make sampling efficiency similar to overdamped (**OPEN**: to precise).
- **Another problem**: Rejections in Metropolis depends badly on dimensionality of fast-degrees of freedom.
- **Open practical/theoretical question**: how can we obtain a **compromise between time-step bias and sampling efficiency** ??
If your favorite chemists gives you a code, can you construct a black box algorithm which computes the "good" time-step, computes weights and rejections, in order to achieve this compromise ?

- d evolving constraints associated with the reaction coordinate;

Hamiltonian:

$$\begin{cases} H(p, q) = \frac{1}{2}p^T M^{-1}p + V(q) \\ \xi_{\text{slow}}(q) = z(t) \end{cases} \quad (C)$$

- Verlet explicit integration + constraints solver with Newton/Gauss solver .

$$\left\{ \begin{array}{l} p^{n+1/2} = p^n - \frac{\Delta t}{2} \nabla V(q^n) - \nabla \xi_{\text{slow}}(q^n) \lambda^{n+1/2}, \\ q^{n+1} = q^n + \Delta t M^{-1} p^{n+1/2}, \\ \xi_{\text{slow}}(q^{n+1}) = z(t_{n+1}) \quad (C_q), \\ p^{n+1} = p^{n+1/2} - \frac{\Delta t}{2} \nabla V(q^{n+1}) - \nabla \xi_{\text{slow}}(q^{n+1}) \lambda^{n+1} \\ \nabla \xi_{\text{slow}}(q^{n+1})^T M^{-1} p^{n+1} = \frac{z(t_{n+1}) - z(t_n)}{\Delta t} \quad (C_p), \end{array} \right.$$

- Time-reversible under $p \rightarrow -p$ and $z(t_{n+1}) \leftrightarrow z(t_n)$. Conservation of the symplectic structure.

Conservation of the symplectic structure

- It may seem surprising to **obtain symplecticity** for a mapping between two different phase-space.
- In fact, the numerical flow is symplectic in the **full phase-space** \mathbb{R}^{6N} , and symplecticity is inherited on constraints manifolds.
- One can thus get detailed balance up to an energy drift, that is to say:

$$\begin{aligned}
 & e^{-\beta H(q_n, p_n)} \underbrace{\delta_{\Phi_{z_n, z_{n+1}}(q_n, p_n)}(dq_{n+1} dp_{n+1})}_{\text{numerical flow at } z_n} \underbrace{\sigma_{z_n}(dq_n dp_n)}_{\text{phase-space measure}} = \\
 & \underbrace{e^{-\beta(H(q_n, p_n) - H(q_{n+1}, p_{n+1}))}}_{\text{work = energy variation}} \times \\
 & e^{-\beta(H(q_{n+1}, p_{n+1}))} \delta_{\Phi_{z_{n+1}, z_n}(q_{n+1}, p_{n+1})}(dq_n dp_n) \sigma_{z_{n+1}}(dq_{n+1} dp_{n+1}),
 \end{aligned}$$

- Add carefully a thermostat.
- Define the work as

$$\mathcal{W}_{0,T} = \sum_n \underbrace{\Delta H_n}_{\text{energy variation of hamiltonian int}}$$

energy variation of hamiltonian int

- This yields a discrete Jarzynski identity:

$$F(z(T)) - F(z(0)) = -\beta \ln \mathbb{E}(e^{-\beta(\mathcal{W}_{0,T} + Corr_T - Corr_0)})$$

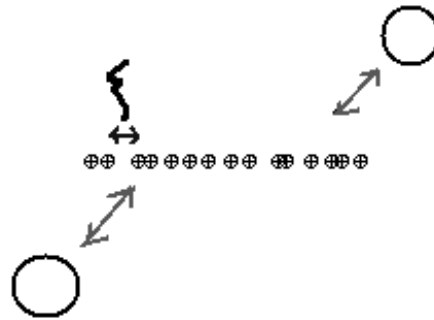
- Thus **time-reversibility + symplecticity** yields:
 - (i) Detailed balance up to a factor.
 - (ii) Metropolis correction in dynamics with rejection.
 - (iii) Time-step correcting weights in non-equilibrium Jarzynski.
- OPEN: Show theoretically following idea: there is an **incompressible source of error** : **bias from time-step** in Langevin \sim **variance from mixing slow-down** in Metropolis \sim **variance** in Jarzynski work **exponential weights**. **I DO NOT KNOW HOW TO DO BUT I SUSPECT IT IS TRUE IN SOME REGIME** .

- OPEN 2: IS THERE A WAY TO ACHIEVE A BIAS/VARIANCE COMPROMISE REGIME ?
- OPEN 3: CAN YOU PROVE THAT NOTHING MORE CLEVER CAN BE DONE (no trick theorem) ?

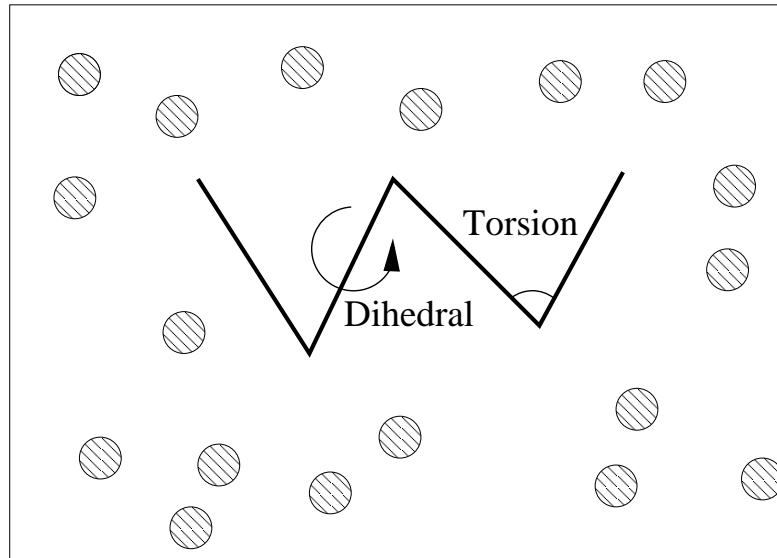
ERRORS FROM CONSTRAINTS

- Many MD codes enforce **holonomic constraints on ξ_{fast}** to enlarge time-step.
- This is called **molecular constraints** and **introduce errors** (assuming you know that the free dynamics is true !!!).
- Assuming that the free dynamics is true, I will show how to **recover exact sampling for free** (IMMP, MR, P. Plechac 2010).

- Ex.: Large molecule + slow environment:



- ξ_{fast} can be: bond lengths, bond angles, dihedral angles .



- Verlet explicit integration + constraints solver with Newton/Gauss solver .

$$\left\{ \begin{array}{l} p^{n+1/2} = p^n - \frac{\Delta t}{2} \nabla V(q^n) - \nabla \xi_{\text{fast}}(q^n) \lambda^{n+1/2}, \\ q^{n+1} = q^n + \Delta t M^{-1} p^{n+1/2}, \\ \xi_{\text{fast}}(q^{n+1}) = 0 \quad (C_q), \\ p^{n+1} = p^{n+1/2} - \frac{\Delta t}{2} \nabla V(q^{n+1}) - \nabla \xi_{\text{fast}}(q^{n+1}) \lambda^{n+1} \\ \nabla \xi_{\text{fast}}(q^{n+1})^T M^{-1} p^{n+1} = 0 \quad (C_p), \end{array} \right.$$

- Same properties as non-constrained case: Time-reversible under $p \rightarrow -p$, conserve $\sigma_{T^*\Sigma}(dpdq)$. Quasi-conserve energy $H_{\Delta t, n} = H + O(\Delta t^2 \times \log(\Delta t n))$. Can be Metropolized to GHMC.

- Idea from *Bennett 70's*. Penalize fDOFs with **mass-matrix penalty**:

$$M = \text{Diag}(m_1, ..m_d) \rightarrow M_\nu(q) = M + \nu^2 \nabla_q \xi_{\text{fast}} (\nabla_q \xi_{\text{fast}})^T$$

- ν is the **penalty parameter** ($\nu = 0$ gives back original dynamics).
- The dependance $q \mapsto M_\nu(q)$ needs to be corrected to conserve equilibrium distribution:

$$\begin{aligned} V(q) \rightarrow V_{\text{eff},\nu}(q) &= V(q) + \frac{1}{2\beta} \ln \det(M_\nu(q)) \\ &= V(q) + \frac{1}{2\beta} \ln \det(\nu^{-2} Id + (\nabla_q \xi_{\text{fast}})^T M^{-1} \nabla_q \xi_{\text{fast}}) \end{aligned}$$

- Pb: Hamiltonian **non longer separable** and $q \mapsto M_\nu(q)$ can be **stiff**.

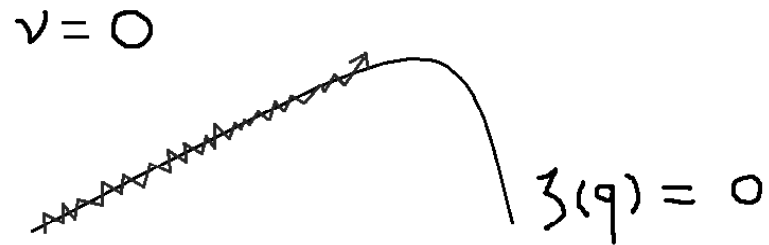
- Extend state-space with n new DOFs with mass 1 (z, p_z) associated with each fDOFs; and n constraints. Extended Hamiltonian:

$$\left\{ \begin{array}{l} H_{\text{IMMP}}(p, p_z, q, z) = \frac{1}{2} p^T M^{-1} p + \underbrace{\frac{1}{2} p_z^T p_z}_{\text{penalty}} + V_{\text{eff}, \nu}(q) \\ \xi_{\text{fast}}(q) = \frac{z}{\nu} \end{array} \right. \quad (C)$$

- Formulation equivalent to explicit mass-matrix penalisation M_ν .
- When $\nu \rightarrow 0$, exact Hamiltonian is recovered.
- When $\nu \rightarrow +\infty$, Hamiltonian system with constraints $\xi_{\text{fast}}(q) = 0$.

Picture

- Velocities are penalized away from the tangent slow bundle.
- Boltzmann distribution in position is conserved.



- Since the model is Hamiltonian, you can recover all the weighting/Metropolis tools and perform (**exact sampling**) in position:

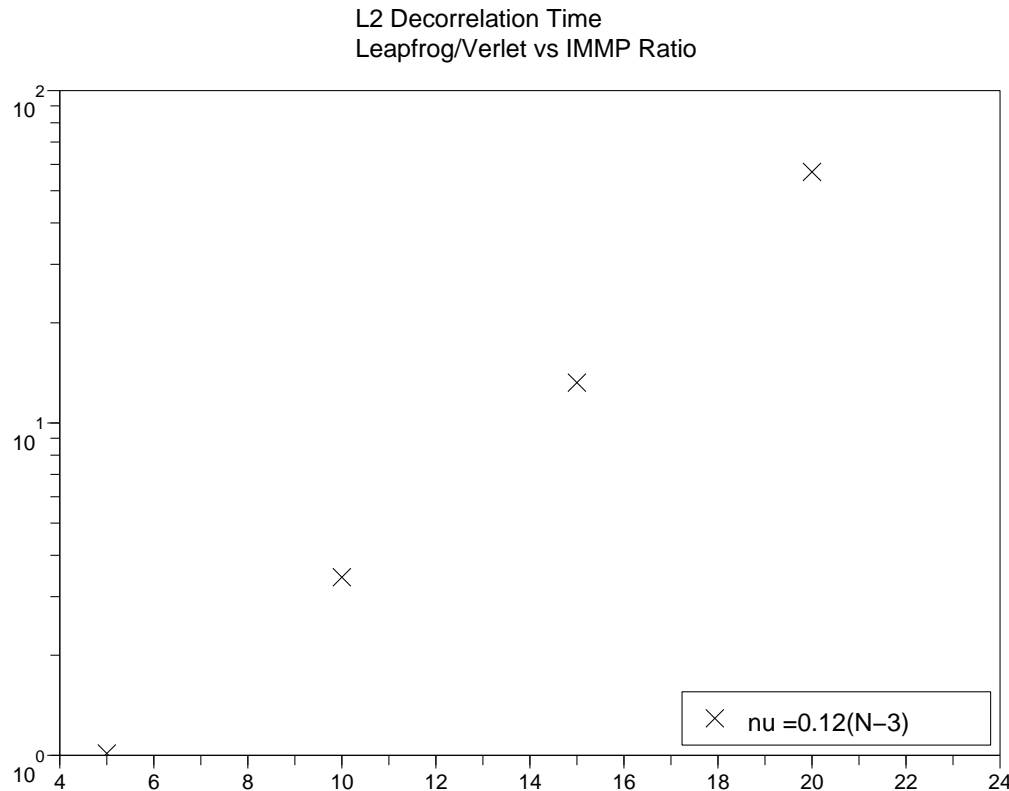
$$\propto e^{-\beta V(q)} dq.$$

- The fastest degrees of freedom are **slowed down** as desired !!

Numerical results: Alkane+sampling

ξ_{fast} = torsion angles, the rest is constrained. SIMU !!

l^2 decorrelation time of the end-to-end alkane length in terms of Monte-Carlo iteration steps. The ratio between the Verlet integration and the IMMP integration is given. Note that the y -axis is in logarithmic scale, and an exponential gain occurs.



ERRORS FROM QUANTUM

- The subject IS **NOT cleaned** up at all.
- (i) Errors from failure of **semi-classical** limit for nuclei.
- (ii) Errors from failure of **adiabatic decoupling** .

- A simple computation shows that

$$\hbar\omega \gg \beta^{-1}$$

where ω is the frequency of most bond oscillations !! (low temperature case). This is in apparent contradiction with the classicality of the nuclei dynamics.

- (i) OPEN: study the **commutation of the limit $\varepsilon \rightarrow 0$ and $\beta \rightarrow +\infty$** for a given potential. It is not clear to me what is happening. A possible route: starts directly from the Gibbs quantum state:

$$\frac{\text{Tr}(e^{-\beta H_\varepsilon}, \cdot)}{\text{Tr}(e^{-\beta H_\varepsilon})}.$$

- (ii) OPEN: study the **commutation** between the (i) process and the **Matching Procedure** to compute effective potential.
- (iii) OPEN: Find a non-controversial **effective potential for bonded interactions**.

- The cause is **conical crossing** (or "diabolic points"), and is defined as the submanifold where $E_e^0(q) = E_e^1(q)$.
- Many real life examples exhibit case where the chemistry **DO USE canonical crossing** (coupling with light).
- Here again, the dynamics is quite well-known (**surface hopping, Landau-Zener formula ...**), but :
- (i) OPEN: expand the semi-classical limit of the **quantum Gibbs distribution**.
- (ii) OPEN: find **all Langevin** versions of **surface hopping** that samples the expanded Gibbs distribution.
- (ii) OPEN: find the most **efficient** one...

- Even for simple cases there is a lack of **benchmarking**.
- It would be useful to make a systematic study of **ALL** sources of errors on simple molecular examples, and a review on the most reasonable methods.