# Understanding Molecular Simulation

Advanced MC

# **Advanced MC Sampling**

- Exotic ensembles
- Improved sampling:
  - Rejectionless cluster moves
  - include rejections: waste recycling
- Computing the chemical potential
  - Widom insertion
  - overlapping distributions
- Biased sampling: umbrella sampling
- The problem of computer time

## **Summary ensembles**

Ensemble	Constant (Imposed)	Fluctuating (Measured)	Function
NVT	N,V,T	Р	βF=-InQ(N,V,T)
NPT	N,P,T	V	βG=-lnQ(N,P,T)=βF+βPV
μVT	μ,V,T	N	$\beta \Omega = -\ln Q(\mu, V, T) = -\beta PV$

# Semi grand ensemble

- For mixtures an additional ensemble exists: semigrand
- Constant N $\Delta\mu$ VT or N $\Delta\mu$ PT :
  - The difference between the chemical potentials of the components is fixed
  - total number of particles is fixed
  - Pressure or volume fixed
- For a binary mixture
  - $N_1$ ,  $N_2$  are allowed to change, but  $N_1+N_2=N$
  - $\mu_2$ ,  $\mu_1$  are allowed to change, but  $\Delta \mu = \mu_2$   $\mu_1$

$$\Xi = \frac{\beta P}{N! \Lambda^{3N}} \int dV V^N \exp[-\beta PV] \times \sum_{identities} \exp[\beta N_2 \Delta \mu] \times \int ds^N \exp[-\beta U(s^N)]$$
  
In binary mixture just 1,2

#### MC move changes identity



 $P_{acc}(1 \rightarrow 2) = \min[1, \exp[-\beta U + \beta \Delta \mu]]$ 

Advantage: at high densities still good acceptance

## **Summary ensembles**

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NPT	N,P,T	V	βG=-InQ(N,P,T)=βF+βPV
μVT	μ,V,T	N	βΩ=-lnQ(μ,V,T)=-βPV
Δμ <b>V</b> Τ	ΔμVΤ	μ <sub>i</sub> ,N <sub>i</sub>	$\beta Y = -\ln \Xi(\Delta \mu, V, T)$

#### **Exotic ensembles**

What to do with a biological membrane?



## Model membrane: Lipid bilayer



hydrophilic head group

two hydrophobic tails



# Surface Tension y controls area per lipid

γ<0	compressed bilayer
γ=0	tensionless bilayer
γ>0	stretched bilayer

# Simulations at imposed surface tension

- Simulation to a constant surface tension
  - Simulation box: allow the area of the bilayer to change in such a way that the volume is constant.





#### **Constant surface tension simulation**

$$N_{N\gamma T}(A,\mathbf{r}^N) \propto \exp[-\beta(\mathbf{U}(\mathbf{r}^N)-\gamma A)]$$

$$A \implies A'$$
$$L \implies L'$$
$$A \perp = A' \perp' = V$$

$$P_{acc} = \min(1, \exp\{-\beta[U(\mathbf{s}^{N}; \mathbf{A}') - U(\mathbf{s}^{N}; \mathbf{A}) - \gamma(\mathbf{A}' - \mathbf{A})]\})$$

## Tensionless state: γ= 0



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Conventional MC performs a random walk in configuration space, such that the number of times that each point is visited, is proportional to its Boltzmann weight.

$$n(\mathbf{r}^N) = c \exp[-\beta \mathcal{U}(\mathbf{r}^N)]$$

$$\frac{\operatorname{acc}(o \to n)}{\operatorname{acc}(n \to o)} = \frac{\mathcal{N}(n)}{\mathcal{N}(o)} = \exp\{-\beta[\mathcal{U}(n) - \mathcal{U}(o)]\}$$

Metropolis, Rosenbluth,Rosenbluth,

Teller and Teller choice:



$$\operatorname{acc}(o \to n) = \min\left(1, \exp\{-\beta[\mathcal{U}(\mathbf{r'}^N) - \mathcal{U}(\mathbf{r}^N)]\}\right)$$

# Satisfactory?



Solution of conflict: play with the a-priori probabilities of trial moves:

$$\alpha(o \to n) \neq \alpha(n \to o)$$

$$\frac{\operatorname{acc}(o \to n)}{\operatorname{acc}(n \to o)} = \frac{\alpha(n \to o)}{\alpha(o \to n)} \exp\{-\beta[\mathcal{U}(n) - \mathcal{U}(o)]\}.$$

In particular, if:

$$\frac{\alpha(n \to o)}{\alpha(o \to n)} = \exp\{-\beta[\mathcal{U}(o) - \mathcal{U}(n)]\}.$$

Then

$$\frac{\operatorname{acc}(o \to n)}{\operatorname{acc}(n \to o)} = 1 \qquad (100\% \operatorname{acceptance})$$

100% acceptance can be achieved in special cases: e.g. Swendsen-Wang, Wolff, Luyten, Whitelam-Geissler, Bortz-Kalos-Lebowitz...

General idea: construct "cluster moves"

Simplest example: Swendsen-Wang



Illustration: 2D Ising model.

Snapshot: some neighbors are parallel, others antiparallel



Number of parallel nearest-neighbor pairs:  $N_p$ Number of anti-parallel nearest neighbor pairs is:  $N_a$ Total energy:  $U = (N_a - N_p) J$ 



Make "bonds" between parallel neighbors. The probability to have a bond (red line) between parallel neighbors is **p** (as yet undetermined). With a probability **1-p**, parallel neighbors are not connected (blue dashed line).



Form clusters of all spins that are connected by bonds. Some clusters are all "spin up" others are all "spin down".

Denote the number of clusters by **M**.



Now randomly flip clusters. This yields a new cluster configuration with probability  $P_{(flip)} = (1/2)^{M}$ .

Then reconnect parallel spins



Next: forget about the "bonds"...



New spin configuration!

$$P_o P_{clus}(o) P_{flip}(M) P_{acc}(o \to n)$$

\_

$$P_n P_{clus}(n) P_{flip}(M) P_{acc}(n \to o)$$

$$\exp(-\beta U_o)p^{n_c}(1-p)^{N_p(o)-n_c}(1/2)^M P_{acc}(o \to n)$$

$$\exp(-\beta U_n)p^{n_c}(1-p)^{N_p(n)-n_c}(1/2)^M P_{acc}(n \to o)$$



Moreover, we want 100% acceptance, i.e.:

$$P_{acc}(o \rightarrow n) = P_{acc}(n \rightarrow o) = 1$$

$$\exp(-\beta U_o)p^{n_c}(1-p)^{N_p(o)-n_c}(1/2)^M P_{acc}(o \to n)$$

$$\exp(-\beta U_n)p^{n_c}(1-p)^{N_p(n)-\eta_c}(1/2)^M P_{acc}(n \to o)$$

#### Hence:

$$\exp(-\beta U_o)(1-p)^{N_p(o)} = \exp(-\beta U_n)(1-p)^{N_p(n)}$$

$$\exp(\beta(U_n - U_o)) = (1 - p)^{N_p(n) - N_p(o)}$$

#### But remember:

$$U_n - U_o = J(N_a(n) - N_p(n)) - J(N_a(o) - N_p(o))$$

or

$$\Delta U = J(\Delta N_a - \Delta N_p)$$

# But: $\Delta N_a = -\Delta N_p$ and therefore $\Delta U = -2J\Delta N_p$

$$\exp(\beta(U_n - U_o)) = \exp(-2\beta J(N_p(n) - N_p(o)))$$

Combining this with:  $\exp(\beta(U_n - U_o)) = (1 - p)^{N_p(n) - N_p(o)}$ 

we obtain:

$$p = 1 - \exp(-2\beta J)$$

100% acceptance!!!





#### Why not?



For Swendsen-Wang, we generate very many trial states : with **n** clusters, **2**<sup>**n**</sup> possible states...)

...and yet we accept only **one**!

## HERETICAL MC SAMPLING

Include "rejected" moves in the sampling

Why is this heretical?:

Metropolis "importance" sampling is **based** on the earlier (Ulam/von Neumann) **rejection** method applied to **random** MC sampling
This is the key:

$$\sum_{m} \rho(m) \pi_{mn} = \rho(n)$$

The transition matrix  $\pi$  leaves the equilibrium distribution  $\rho$  unchanged.

$$\langle A \rangle_{\rho} = \sum_{n} A_{n} \rho_{n}$$

we can rewrite this using 
$$\sum \rho_m \pi_{mn} = \rho_n$$

$$\sum_{n} A_n \rho_n = \sum_{n} \sum_{m} A_n \rho_m \pi_{mn} = \sum_{m} \rho_m \sum_{n} A_n \pi_{mn}$$

$$=\sum_{m}\rho_{m}\sum_{n}A_{n}\pi_{mn} \Leftrightarrow \langle A \rangle_{\rho} = \left\langle \sum_{n}\pi_{mn}A_{n} \right\rangle_{\rho_{m}}$$

$$\langle A \rangle_{\rho} = \left\langle \sum_{n} \pi_{mn} A_{n} \right\rangle_{\rho_{m}}$$

Note that <A> is no longer an average over "visited" states – we also include "rejected" moves in the sampling.

$$\langle A \rangle_{\rho} = \left\langle \sum_{n} \pi_{mn} A_{n} \right\rangle_{\rho_{m}}$$

This relation also holds for any set of "connected" trial states: i.e. the possible final states of a single (decent) MC trial move.

$$\langle A \rangle_{\rho} = \left\langle \sum_{n} \pi_{mn} A_{n} \right\rangle_{\rho_{m}}$$

For instance: in conventional MC, there would be only two states (the "old" state and the "new" state)

$$\langle A \rangle_{\rho} = \left\langle \sum_{n} \pi_{mn} A_{n} \right\rangle_{\rho_{m}}$$

But in other algorithms, there are many. e.g. in the Swendsen-Wang algorithm: **n** clusters that can be flipped  $\Rightarrow 2^n$  connected states.

The more parallel the algorithm, the better...

$$\langle A \rangle_{\rho} = \left\langle \sum_{n} \pi_{mn} A_{n} \right\rangle_{\rho_{m}}$$

Note that the transition matrix that is used in the averaging need not be the same as the one used in sampling the  $\rho_{m}$ .

(e.g. one could be "Barker" and the other "Metropolis")

$$\langle A \rangle_{\rho} = \left\langle \sum_{n} \pi_{mn} A_{n} \right\rangle_{\rho_{m}}$$

## How to sample this?

$$\pi_{nm} = \alpha_{nm} P_{acc}(nm)$$
$$\langle A \rangle_{\rho} = \sum_{m} \rho_{m} \left\langle \sum_{n} P_{acc}(mn) A_{n} \right\rangle_{\alpha}$$

### Slightly dishonest and slightly trivial example:

# Sampling the magnetization of a 2D Ising system

## Compare:

- Normal (Swendsen-Wang) MC (sample one out of 2<sup>n</sup> states)
- 2. Idem + "waste recycling" (sample all 2<sup>n</sup> states)



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A central quantity in the study of (phase) equilibria is the chemical potential  $\mu$ .

We have an "intuitive" picture of the temperature T and the pressure P.

Can we gain a similar "intuitive" understanding of the chemical potential?

First, look at the formal definition:

$$dE = TdS - PdV + \mu dN$$

$$dF = -SdT - PdV + \mu dN$$

#### Hence:

$$\mu = \left(\frac{\partial E}{\partial N}\right)_{S,V} = \left(\frac{\partial F}{\partial N}\right)_{T,V}$$

Using  $F=-k_BT \ln Q(N,V,T)$  we can write:

$$\mu = \left(\frac{\partial F}{\partial N}\right)_{V,T}$$

$$= -k_B T \lim_{\Delta N \to 0} \frac{\ln Q(N + \Delta N, V, T) - \ln Q(N, V, T)}{\Delta N}$$

$$= -k_B T \ln \left(\frac{Q(N + 1, V, T)}{Q(N, V, T)}\right)_{V,T}$$

For N >> 1



with  $\rho \equiv (N/V)$ 

The ideal-gas result, we had obtained before. Now, with interacting particles:

$$\mu = \lim_{N \to \infty} -k_B T \ln \left[ \frac{Q(N+1, V, T)}{Q(N, V, T)} \right]$$
$$\equiv \mu^{\text{id}} + \mu^{\text{excess}}$$

We can there write the "excess" part of the chemical potential as:

$$\mu^{\text{ex}} = k_B T \left[ \ln \frac{Q(N+1,V,T)}{Q(N,V,T)} - \ln \left( \frac{V^3}{\Lambda^3(N+1)} \right) \right]$$
$$= k_B T \ln \left[ \frac{Q(N+1,V,T)\Lambda^3(N+1)}{VQ(N,V,T)} \right]$$

Now we use the explicit expression for Q(N,V,T):

$$Q(N, V, T) = \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N e^{-\beta U(\mathbf{r}^N)}$$

$$\mu^{\mathbf{ex}} = -k_B T \ln \left( \frac{\int d\mathbf{r}^{N+1} e^{-\beta U(\mathbf{r}^{N+1})}}{V \int d\mathbf{r}^N e^{-\beta U(\mathbf{r}^N)}} \right)$$

We define:

$$\Delta U(\mathbf{r}^N;\mathbf{r}_{N+1}) \equiv U(\mathbf{r}^{N+1}) - U(\mathbf{r}^N)$$

Then

$$\begin{split} \mu^{\mathbf{e}\mathbf{x}} &= -k_B T \ln\left(\frac{\int d\mathbf{r}^{N+1} e^{-\beta \left[U(\mathbf{r}^N) + \Delta U(\mathbf{r}^N; \mathbf{r}_{N+1})\right]}}{V \int d\mathbf{r}^N e^{-\beta U(\mathbf{r}^N)}}\right) \\ &= -k_B T \ln\left(\frac{1}{V} \int d\mathbf{r}_{N+1} \frac{\int d\mathbf{r}^N e^{-\beta \left[U(\mathbf{r}^N) + \Delta U(\mathbf{r}^N; \mathbf{r}_{N+1})\right]}}{\int d\mathbf{r}^N e^{-\beta U(\mathbf{r}^N)}}\right) \end{split}$$

$$\mu^{\mathbf{e}\mathbf{x}} = -k_B T \ln\left(\frac{1}{V} \int d\mathbf{r}_{N+1} \frac{\int d\mathbf{r}^N e^{-\beta U(\mathbf{r}^N)} e^{-\beta \Delta U(\mathbf{r}^N;\mathbf{r}_{N+1})}}{\int d\mathbf{r}^N e^{-\beta U(\mathbf{r}^N)}}\right)$$
$$= -k_B T \ln\frac{1}{V} \int d\mathbf{r}_{N+1} \left\langle e^{-\beta \Delta U(\mathbf{r}^N;\mathbf{r}_{N+1})} \right\rangle_N$$

For a homogeneous system, the average does not depend on the position  $\mathbf{r}_{N+1}$ . Hence

$$\beta \mu^{\rm ex} = -\ln \left\langle e^{-\beta \Delta U} \right\rangle_N$$

In words: the excess chemical potential is equal to minus the logarithm of the average Boltzmann factor associated with the **random insertion** of an additional particle N+1 in an N-particle system.

So, finally, we get:

$$\mu = \mu_{id.gas} - kT \ln \left( \int ds_{N+1} \langle \exp(-\beta \Delta U(s_{N+1}, s^N)) \rangle \right)$$

Interpretation:

- 1. Evaluate  $\Delta U$  for a random insertion of a molecule in a system containing N molecule.
- 2. Compute  $\exp(-\beta \Delta U)$
- 3. Repeat M times and compute the average "Boltzmann factor"  $< \exp(-\beta \Delta U) >$
- 4. Then  $\mu_{excess} = -kT \ln \langle \exp(-\beta \Delta U) \rangle$

#### Algorithm 16 (Widom Test Particle Insertion)

```
subroutine Widom
xtest=box*ranf()
call ener(xtest,entest)
wtest=wtest
+ +exp(-beta*entest)
return
end
```

excess chemical potential via the addition of test particles generate a random position determine energy update Boltzmann factor in (7.2.5) This method to evaluate  $\mu$  is commonly known as **Widom**'s "particle-insertion method".



Ben Widom

Example: excess chemical potential of "hard" particles



$$\langle e^{-\beta\Delta U} \rangle = P_{\text{overlap}} \times 0 + P_{\text{no overlap}} \times 1$$
  
=  $P_{\text{no overlap}}$ 

In the case of hard particles, the excess chemical potential is related to the probability of a successful random insertion:

$$\beta \mu^{\rm ex} = -k_B T \ln P_{\rm no \ overlap}$$

Example of an application:

We can use the particle-insertion method to derive an expression for the second virial coefficient.

# Lennard-Jones fluid



## **Other ensembles: NPT**



## Widom particle insertion

NVT:  

$$\beta \mu = \beta \ln(\rho) - \ln \left\langle \int ds_{N+1} \exp[-\beta \Delta U^+] \right\rangle_{NVT}$$

NPT:  

$$\beta \mu = \ln(\Lambda^{3}\beta P) - \ln\left\langle\frac{\beta PV}{N+1}\int ds_{N+1}\exp(-\beta\Delta U^{+})\right\rangle$$

#### ACCEPTANCE OF RANDOM INSERTION DEPENDS ON SIZE





#### ACCEPTANCE OF RANDOM INSERTION DEPENDS ON DENSITY



Insertion easy



Insertion difficult

Particle insertion continued....

$$\mu \approx \left(\frac{F(N+1,V,T) - F(N,V,T)}{(N+1) - N}\right)$$

therefore

$$\mu \approx -kT \ln \frac{Q(N+1,V,T)}{Q(N,V,T)}$$

But also

$$\mu \approx +kT \ln \frac{Q(N, V, T)}{Q(N+1, V, T)}$$

Interpreted as particle removal

As before:

$$Q(N, V, T) = \frac{1}{N! \Lambda^{3N}} \int dr^N \exp(-\beta U(r^N)) = \frac{1}{N! \Lambda^{3N}} V^N \int ds^N \exp(-\beta U(s^N; L))$$

With **s** a scaled coordinate:  $0 \le s \le 1$ 

 $\mathbf{r} = \mathbf{L} \mathbf{s}$  (is box size)

$$\frac{Q(N, V, T)}{Q(N+1, V, T)} =$$

$$\frac{(N+1)\Lambda^3}{V} \frac{\int ds^N \exp(-\beta U(s^N))}{\int ds^{N+1} \exp(-\beta U(s^{N+1}))}$$

$$= \frac{(N+1)\Lambda^3}{V} \frac{\int ds_{N+1} \int ds^N \exp(-\beta U(s^N))}{\int ds^{N+1} \exp(-\beta U(s^{N+1}))}$$

However, now write

$$U((s^N) \equiv U(s^{N+1}) - \Delta U(s_{N+1}, s^N)$$

$$\frac{Q(N,V,T)}{Q(N+1,V,T)} =$$

$$\frac{(N+1)\Lambda^3}{V} \langle \exp(+\beta \Delta U(s_{N+1}, s^N)) \rangle_{N+1}$$

And therefore

$$\mu = \mu_{id.gas} + kT \ln \left( \langle \exp(+\beta \Delta U(s_{N+1}, s^N)) \rangle_{N+1} \right)$$
Positive sign!

Interpretation:

- Evaluate ∆U for a random REMOVAL of a molecule in a system containing N+1 molecule.
- 2. Compute  $\exp(+\beta\Delta U)$
- 3. Repeat M times and compute the average "Boltzmann factor"
- 4. Then

 $\mu_{excess} = +kT \ln < \exp(+\beta \Delta U) >$ 

This approach DOES NOT WORK

What is wrong?

 $\exp(+\beta\Delta U)$ 

is not bounded. The average that we compute can be dominated by INFINITE contributions from points that are NEVER sampled.

What to do?

Consider the energy distribution upon insertion:

$$p_{0}(\Delta U) \equiv \frac{\int \exp(-\beta U_{N})\delta(\Delta U - U_{N+1} + U_{N})}{\int \exp(-\beta U_{N})}$$

And also consider the energy distribution for removal

$$p_1(\Delta U) \equiv \frac{\int \exp(-\beta U_{N+1})\delta(\Delta U - U_{N+1} + U_N)}{\int \exp(-\beta U_{N+1})}$$

 $p_0$  and  $p_1$  are related:

$$p_1(\Delta U) = \frac{\int \exp(-\beta(U_N + \Delta U))\delta(\Delta U - U_{N+1} + U_N)}{Q_{N+1}}$$
$$= \exp(-\beta\Delta U) \frac{\int \exp(-\beta U_N)\delta(\Delta U - U_{N+1} + U_N)}{Q_{N+1}}$$

$$\ln p_{1}(\Delta U) = \beta (\Delta F - \Delta U) + \ln p_{0} (\Delta U)$$

$$f_{0}(\Delta U) = \ln p_{0} (\Delta U) - 0.5\beta\Delta U$$

$$f_{1}(\Delta U) = \ln p_{1} (\Delta U) + 0.5\beta\Delta U$$

$$\beta\Delta I \quad \text{Fit } f_{0} \text{ and } f_{1} \text{ to two polynomials that only}$$

$$differ \text{ by a constant.}$$

$$f_{1}(\Delta U) = C_{1} \quad \text{simulate system 0: compute } f_{0}$$

$$f_{0}(\Delta U) = C_{0} + a\Delta U + b\Delta U^{2} + c\Delta U^{3}$$

$$\beta\Delta F = C_{1} - C_{0} \quad \text{This is known as the overlapping}$$

This is known as the overlapping distribution method (sec 7.2.3)
### µ from overlapping distributions



Does it work for hard spheres?

consider  $\Delta U=0$  $f_1(0) = f_0(0) + \beta \mu$  $f_1(0) = \ln(1) + (constant)$  $f_0(0) = \ln(P_{acc}) + (constant)$  $\beta \mu = -\ln(P_{acc})$ YFSI

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### From histograms to free-energy landscapes

The function  $p(\Delta U)$  is an example of a histogram where  $\Delta U(\mathbf{r}^N)$  plays the role of the coordinate. We can consider other coordinates Q(r<sup>N</sup>). Examples...

The probability that  $Q(r^N)$  has a value Q is given by

$$P(Q_0) = \frac{\int \exp(-\beta U(r^N))\delta(Q(r^N) - Q_0)}{\int \exp(-\beta U(r^N))}$$

From this probability we can derive (`define') the variation of the free energy with Q:

$$F(Q) = -kT \ln(P(Q)) + \text{constant}$$

### From histograms to free-energy landscapes

Free energy landscapes are interesting in many different contexts (phase transitions, protein folding, ...)

Sampling a free energy landscape is easy near the minima, and difficult near the maxima because P(Q) is small where F(Q) is large..

# $F(Q) = -kT \ln(P(Q)) + \text{constant}$

Naively, we could sample determine F(Q) by accumulating a histogram of the spontaneous fluctuations of Q.



Q

Problem: F(Q) is very noisy, except near its minimum.

## **Biased sampling**





## METROPOLIS SAMPLING

BIASED SAMPLING

#### Application of biased MC simulation to determine $P_w(Q)$

$$P_w(Q_0) = \frac{\int \exp[-\beta(U(r^N) + w(Q(r^N))]\delta(Q(r^N) - Q_0))}{\int \exp[-\beta(U(r^N) + w(Q(r^N))]}$$

But this we can rewrite as

$$P_w(Q_0) = \exp(-\beta w(Q_0)) \frac{\int \exp[-\beta U(r^N)] \delta(Q(r^N) - Q_0)}{\int \exp[-\beta (U(r^N) + w(Q(r^N))]}$$

Ideally, we should choose w(Q) = -F(Q), because then the biased histogram would be flat. But, of course, we don't know F(Q) a priori... Once we know the biased histogram

$$P_w(Q_0) = c \exp(-\beta w(Q_0)) P(Q_0)$$

we can reconstruct the unbiased histogram – up to a constant

$$-kT\ln(P(Q_0)) = -kT\ln(P_w(Q_0)) + w(Q_0) - kT\ln c$$

$$F(Q_0) = -kT \ln(P_w(Q_0)) + w(Q_0) - kT \ln c$$

In summary: We can choose the bias w(Q) such that any desired range of Q-values is sampled.

And we can correct for the bias.

For obscure reasons this is often called umbrella sampling



The only remaining problem is that the different parts of F(Q) are shifted with respect to each other.

#### Solutions:

- 1. Fit to a single curve (not very elegant, but effective)
- 2. Use `Ferrenberg-Swendsen' scheme to combine different parts of the histogram (more elegant, but more sensitive to noise)
- 3. Use the WHAM method of Kumar et al. or MBAR from Chodera et al.

**Result:** 



#### FINALLY...



#### Solutions:

#### **1.** Wait until there is enough computer power

Moore's law: computing power  $\Gamma$  increases a factor 10 every 5 years

$$\Gamma(\Delta t) = \Gamma_0 e^{\Delta t/\tau}$$

with  $\tau \approx 2.17$  years.

When to start a major calculation? Either start now, or wait a time  $\Delta t$  and use the computers that are available then:

For a computation involving X operations, the expected time to completion, using the current computing power  $\Gamma_0$ , is:  $t_0 = \frac{X}{\Gamma_0}$ .

### But if we first wait a time $\Delta t$ , it is:

$$t = \Delta t + \frac{t_0}{e^{\Delta t/\tau}}$$

#### Optimum? Compute extremum:

$$0 = 1 - \frac{t_0}{\tau} e^{-\Delta t/\tau}$$

Hence

$$1 = \frac{t_0}{\tau} e^{-\Delta t/\tau}$$

No solutions if  $t_0 < \tau$ , i.e. if the estimated computing time is less than 2.17 years.

In that case: start right away! Otherwise:

wait for a time interval

 $\Delta t = \tau \ln(t_0/\tau)$ 

then buy a computer, and start!!!

#### Solutions:

- **1. Wait until there is enough computer power**
- 2. Use cheaper models/ more efficient algorithms.

