Statistical thermodynamics for MD and MC simulations

knowing 2 atoms and wishing to know 10^{23} of them

Marcus Elstner and Tomáš Kubař

22 June 2016

Thermodynamic properties of molecular systems

statistical mechanics: the way from the properties of particles to the thermodynamic properties of ensembles via the partition function

- how is the thermodynamic equilibrium characterized?
- which quantities are of interest?
- how may these be derived from the partition function?
- how to derive the ensemble partition function from the partition function of a single molecule?
- how is partition function connected to phase-space density?
- MD simulation provides an alternative way to thermodynamic quantities
- it is difficult to obtain free energies
 from normal simulations

classical thermodynamics \rightarrow

Equilibrium and spontaneous processes

```
says which quantites are maximized/minimized in equilibrium
      and show a definite change during spontaneous processes
microcanonical ensemble.
   equilibrium reached if entropy S maximized
   process spontaneously if entropy increases: \Delta S > 0
canonical ensemble - more complex, as we need
   to consider system of interest together with surroundings
   (to identify equilibrium and spontaneity)
to calculate anything for the supersystem
  - impossible - alternative needed
```

thermodynamic equilibrium and spontaneous process

Free energy / enthalpy – fundamental property

How to keep our molecular system and drop the surroundings? introduce a new thermodynamic function:

- Helmholtz free energy F in canonical ensemble
- Gibbs free energy/enthalpy *G* in NPT ensemble

$$F = U - TS$$
 $G = H - TS = U + pV - TS$

- F = F(T, V) or G = G(T, P) depends on experimentally controllable variables T and V/P
- a 'stable' state of the system
 - minimum of F or G rather than of U
 - equivalent to maximization of entropy of universe
- F / G decreases in the course of spontaneous process
 - holy grail of MD simulation

Microscopic definition of entropy

```
W – number of ways ('microstates')in which a certain 'macrostate' may be realized(a certain occupation of energy levels of the system)
```

microscopic entropy:

$$S = k_{\mathsf{B}} \cdot \ln W$$

 $(k_{\rm B} - {\rm universal \ Boltzmann \ constant})$

Microscopic definition of entropy



Microscopic definition of entropy

- S tells us something about the travel of the system through the configuration (phase) space
- equivalently, S can be related to the order in the system
- low entropy few states are occupied
 - only a small part of configuration space accessible
 - ordered system
- high entropy many states are visited
 - extended regions of the configuration space covered
 - less ordered system

```
Example – pile of books on a desk
Jan Černý (Charles University in Prague, Dept Cellular Biology):
anthropy – "entropy of human origin"
```

Introduction

"... series of molecular structures generated by an MD simulation with the Berendsen thermostat does not represent the correct canonical ensemble" – what does this mean?

The phase space may be sampled (walked through) in various ways – just what is the correct way?

Closed system - canonical ensemble

system in thermal contact with the surroundings

- temperature rather than energy remains constant
- Boltzmann distribution of p_i applies:

$$p_i = \frac{\exp[-\beta \cdot \varepsilon_i]}{Q}$$

$$Q = \sum_{j} \exp[-\beta \cdot \varepsilon_j]$$

Q – canonical partition function (Zustandssumme)

derive the meaning of β – fall back to basic thermodynamics. . . :

$$\beta = \frac{1}{k_{\rm R}T}$$

Canonical partition function

```
Q – seems to be purely abstract...
```

but – to characterize the thermodynamics of a system Q is completely sufficient, all thermodynamic observables follow as functions of Q

how to obtain Q?

- there will be 2 ways, depending on the system studied

partition function connects the microscopic and macroscopic world

Using the partition function

we can get all thermodynamic functions from Q:

$$U = \langle E \rangle = k_{\rm B} T^2 \frac{\partial \ln Q}{\partial T}$$

$$S = k_{\rm B} T \cdot \frac{\partial \ln Q}{\partial T} + k_{\rm B} \cdot \ln Q$$

$$F = -k_{\rm B} T \cdot \ln Q$$

$$P = k_{\rm B} T \cdot \left(\frac{\partial \ln Q}{\partial V}\right)_T \qquad \text{(equation of state)}$$

$$H = U + pV$$

$$G = F + pV = H - TS$$

Michal Otyepka (Palacký University Olomouc, Dept Physical Chemistry):

"Just grab the partition function at the tail, and then you have everything!"

2 ways to thermodynamic properties

- simple molecules with 1 or few minima of energy
 - calculate the partition function (trans+vib+rot) (probably employ approximations IG+HO+RR)
 - derive properties from Q

Discrete systems

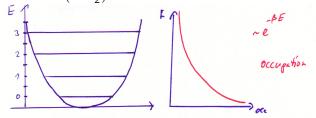
system with discrete energy levels E_i – partition function:

$$Q = \sum_{i} \exp[-\beta E_i]$$

Boltzmann distro function: (prob. of system in state E_i)

$$p_i = \frac{1}{Q} \exp[-\beta E_i]$$

example – HO: $E_i = (i + \frac{1}{2}) \cdot \hbar \omega$



Using the partition function of 1 molecule

partition function of a large system - simplifications possible:

 \blacksquare system of *n* identical and indistinguishable particles (gas):

$$Q=\frac{q^n}{n!}$$

- necessary effort reduced greatly!
 - get the molecular partition function q (calculate 1 molecule, or 2)
 - obtain the ensemble partition function Q and all thermodynamic quantities

Simple molecules

... with 1 or few well characterized minima for a certain minimum – consider contributions to energy:

$$E = E^{\text{el}} + E^{\text{trans}} + E^{\text{rot}} + E^{\text{vib}}$$

partition function follows as

$$Q = \exp\left[-\beta \left(E^{\text{el}} + E^{\text{trans}} + E^{\text{rot}} + E^{\text{vib}}\right)\right] =$$

$$= \exp\left[-\beta E^{\text{el}}\right] \cdot \exp\left[-\beta E^{\text{trans}}\right] \cdot \exp\left[-\beta E^{\text{rot}}\right] \cdot \exp\left[-\beta E^{\text{vib}}\right] =$$

$$= Q^{\text{el}} \cdot Q^{\text{trans}} \cdot Q^{\text{rot}} \cdot Q^{\text{vib}}$$

or

$$\ln Q = \ln Q^{\rm el} + \ln Q^{\rm trans} + \ln Q^{\rm rot} + \ln Q^{\rm vib}$$

Electronic partition function

usually: quite high excitation energy

ightarrow electronic ground state only populated:

$$E^{el}(0) = 0$$
 arbitrarily

electronic partition function:

$$Q^{\mathsf{el}} = \exp[-\beta E^{\mathsf{el}}(0)] + \exp[-\beta E^{\mathsf{el}}(1)] + \ldots \approx 1 + 0 + \ldots = 1$$

so this may be neglected ©

Translational partition function

calculated for quantum-mechanical particle (mass m) in a 3D box: energy levels:

$$E_{n_x,n_y,n_z} = \frac{h^2}{8m} \left(\frac{n_x^2}{L_x^2} + \frac{n_y^2}{L_y^2} + \frac{n_z^2}{L_z^2} \right)$$

quantum numbers $n_i = 1, 2, ...$ partition function:

$$Q^{\mathsf{trans}} = \left(\frac{2\pi m k_{\mathsf{B}} T}{h^2}\right)^{\frac{3}{2}} \cdot V$$

Rotational partition function

calculated for a rigid rotor (moments of inertia I_x): energy levels:

$$E_J = B \cdot J(J+1)$$

quantum number $J=0,1,2,\ldots$, degeneracy of levels 2J+1 rotational constant $B=\frac{\hbar^2}{8\pi^2I}$ (I – moment of inertia)

$$Q^{\mathsf{rot}} = \sum_{J=0}^{\infty} (2J+1) \exp\left[-J(J+1) \cdot \frac{B}{k_{\mathsf{B}}T}\right]$$

for asymmetric top with rotational constants B_x , B_y , B_z :

$$Q^{\text{rot}} = \sqrt{\frac{\pi \left(k_{\text{B}}T\right)^3}{B_x B_y B_z}}$$

Vibrational partition function

calculated with harmonic vibrational frequencies ω_k of the molecule (computation of Hessian in the minimum of potential energy)

each vibrational mode k is one harmonic oscillator

energy levels:

$$E_k^m = \left(m + \frac{1}{2}\right) \cdot \hbar \omega_k$$

where $E_k^0 = \frac{1}{2}\hbar\omega_k$ is zero point vibrational energy

partition function (using $\sum_{n=0}^{\infty} x^n = \frac{1}{1-x}$):

$$Q_k^{\mathsf{vib}} = \sum_{m=0}^{\infty} \exp\left[-\beta \left(m + \frac{1}{2}\right) \hbar \omega_k\right] = \frac{\exp\left[-\frac{1}{2}\beta \hbar \omega_k\right]}{1 - \exp\left[-\beta \hbar \omega_k\right]}$$

each molecule: N-6 vibrational modes = N-6 HOs example - H_2O : 3 modes (2 stretches, 1 bend)

Thermodynamic properties

for enthalpy -pV is needed - simple for IG:

$$pV = Nk_{\rm B}T$$

then, enthalpy and Gibbs free energy follow:

$$H = U + pV = U + Nk_BT$$

 $G = F + pV = F + Nk_BT$

thermal contributions – calculated by default
with many QCh and MD programs
whenever vibrational analysis is requested
reason – vibrational frequencies are computationally costly
while the thermodynamics is done 'for free'

Example – vibrational contributions

$$\ln Q_k = -\frac{1}{2}\beta\hbar\omega_k - \ln\left[1 - \exp[-\beta\hbar\omega_k]\right]$$

$$U_k = -\frac{\partial \ln Q_k}{\partial \beta} = \hbar\omega_k \left(\frac{1}{2} + \frac{1}{\exp[\beta\hbar\omega_k] - 1}\right)$$

consider this for all of N-6 vibrational DOFs, dropping ZPVE:

$$U^{\mathsf{vib}} = \sum_{k=1}^{N-6} \left(\frac{\hbar \omega_k}{\exp[\beta \hbar \omega_k] - 1} \right)$$

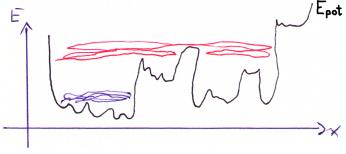
$$\begin{split} F^{\mathrm{vib}} &= -k_{\mathrm{B}} T \ln Q^{\mathrm{vib}} = \sum_{k=1}^{N} k_{\mathrm{B}} T \ln \left[1 - \exp[-\beta \hbar \omega_{k}]\right] \\ \frac{S^{\mathrm{vib}}}{k_{\mathrm{B}}} &= \frac{U^{\mathrm{vib}} - F^{\mathrm{vib}}}{k_{\mathrm{B}} T} = \sum_{k=1}^{N-6} \left(\frac{\beta \hbar \omega_{k}}{\exp[\beta \hbar \omega_{k}] - 1} - \ln \left[1 - \exp[-\beta \hbar \omega_{k}]\right]\right) \end{split}$$

2 ways to thermodynamic properties

- simple molecules with 1 or few minima of energy
 - calculate the partition function (trans+vib+rot) (probably employ approximations IG+HO+RR)
 - derive properties from Q
- flexible molecules, complex molecular systems
 - quantum mechanical energy levels cannot be calculated
 - a single minimum of energy not meaningful
 - do MD (MC) simulation instead, to sample phase space
 - evaluate time averages of thermodynamic quantities

Systems with continuous distribution of energy

dynamics of molecules – at different tot. energies or temperatures, differently extended regions of conformational space are sampled



complex energy landscape $E_{\rm pot}(x)$ blue and red – trajectories at different total energies – different phase-space densities

Continuous systems - canonical ensemble

- every point in phase space a certain value of energy composed of $E_{pot} = E_{pot}(\vec{r})$ (force field), $E_{kin} = E_{kin}(\vec{p})$
- continuous energy levels infinitesimally narrow spacing
- canonical probability distribution function
 - probability to find the system in state with *E*:

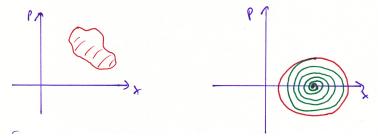
$$P(\vec{r}, \vec{p}) = \rho(\vec{r}, \vec{p}) = \frac{1}{Q} \cdot \exp\left[-\frac{E(\vec{r}, \vec{p})}{k_{\text{B}}T}\right]$$

- perform MD simulation with a correct thermostat, or MC sim.
 the goal is to obtain the right probability density ρ
- partition fction Q integral over phase space rather than sum

$$Q = \int \exp\left[-\frac{E(\vec{r}, \vec{p})}{k_{\rm B}T}\right] \, \mathrm{d}\vec{r} \, \mathrm{d}\vec{p}$$

Thermodynamic quantities - sampling

 $\rho(\vec{r}, \vec{p})$ – gives the probability of finding the system at (\vec{r}, \vec{p}) typically: system is sampling only a part of phase space $(P \neq 0)$:



sampling in MD or MC

undamped and damped classical HO

Continuous distribution of energy

thermodynamic quantities – weighted averages:

Thermodynamic quantities – sampling

$$\langle A \rangle = \frac{\int A \cdot \rho(\vec{r}, \vec{p}) \, d\vec{r} \, d\vec{p}}{\int \rho(\vec{r}, \vec{p}) \, d\vec{r} \, d\vec{p}}$$

why do MD? ... obtain the correct phase-space density ρ ergodic simulation – thermodynamic potentials U, H etc. are obtained as time averages

$$\langle A \rangle = rac{1}{t_1 - t_0} \int_{t_0}^{t_1} A(t) \, \mathrm{d}t$$

- valid only if the simulation has sampled the canonical ensemble
 → phase-space density is correct
- the density ρ is present in the the trajectory inherently \rightarrow e.g. structure with high ρ occur more often

Thermodynamic quantities - sampling

Thus, we have the following to do:

- perform MD simulation (with correct thermostat!)
 - \rightarrow trajectory in phase space (simulation has 'taken care' of the phase-space density)
- get time averages of desired thermodynamic properties

$$\langle A \rangle = \frac{1}{t_1 - t_0} \int_{t_0}^{t_1} A(t) dt$$

Thermodynamic quantities - sampling

we have to obtain the phase-space density with MD simulation

$$\rho(\vec{r}, \vec{p}) = \frac{\exp[-\beta E(\vec{r}, \vec{p})]}{Q} \qquad \vec{r} = \{r_1, \dots, r_{3N}\}, \vec{p} = \{p_1, \dots, p_{3N}\}$$

which is the probability of system occuring at point (\vec{r}, \vec{p})

How long an MD simulation can we perform?

1 ps ightarrow 1,000 points in trajectory

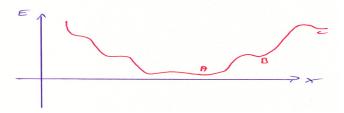
10 ns ightarrow 10M points – we cannot afford much more

example – we have 1,000 points

then, we have hardly sampled (\vec{r},\vec{p}) for which $\rho(\vec{r},\vec{p}) \leq \frac{1}{1000}$

ightarrow points with high energy will be never reached! (while low-energy region may be sampled well)

Missing high-energy points in sampling



High-energy points B and C may be sampled badly

- a typical problem in MD simulations of limited length
- the corresponding large energies are missing in averaging
- when does this matter?
- no serious error for the internal energy exponential dependence of phase-space density kills the contribution

$$ho(ec{r},ec{
ho}) = rac{\mathsf{exp}[-eta E(ec{r},ec{
ho})]}{Q}$$

Missing high-energy points in sampling

Free energies

- determine the spontaneity of process
- NVT canonical Helmholtz function F
- NPT 'canonical' Gibbs function G
- the relevant quantity always obtained depending on whether NVT or NPT simulation is performed
- much more pronounced sampling issues than e.g. for internal energy!

Missing high-energy points in sampling

$$F = -k_{\rm B}T \ln Q = k_{\rm B}T \ln \frac{1}{Q} =$$

$$= k_{\rm B}T \ln \frac{c^{-1} \cdot \iint \exp[\beta E(\vec{r}, \vec{p})] \cdot \exp[-\beta E(\vec{r}, \vec{p})] \, d\vec{r} \, d\vec{p}}{Q} =$$

$$= k_{\rm B}T \ln \iint \exp[\beta E(\vec{r}, \vec{p})] \cdot \rho(\vec{r}, \vec{p}) \, d\vec{r} \, d\vec{p} - \ln c$$

$$= k_{\rm B}T \cdot \ln \left\langle \exp\left[\frac{E}{k_{\rm B}T}\right] \right\rangle - \ln c$$

serious issue – the large energy values enter an exponential, and so the high-energy regions may contribute significantly! \rightarrow if these are undersampled, then free energies are wrong

calculation of free energies impossible! special methods needed!