Stochastic Methods wrap-up

10.34 Dec. 7, 2015

Multidimensional Integrals

- Many important multidimensional integrals have integrands of this form: p(x)f(x) where p is a probability density.
- Often an un-normalized p is known:
 p(x) = w(x)/∭w(x)dx
- Boltzmann distribution: w(x)=exp(-E(x)/kT)
- Bayesian analysis of experiments: $w(\underline{\theta} | data) = p_{prior}(\underline{\theta}) \exp(-\chi^2(\underline{\theta}, data)/2)$

These are candidates for Metropolis Monte Carlo

Metropolis Monte Carlo challenges

- Selecting step length in each dimension
 - want it same order of magnitude as the width of the high-w region
 - If too large, very few steps accepted
 - If too small, takes large number of steps to cover the high-w region.
- Achieving high accuracy requires very large N
 - adding one more sig fig requires 100x more samples than used so far.

Master Equations

- Often we have time-dependent probability of the system being in a discrete state: P_N(t).
 dP/dt = M * P
 P(t_o)=P_o
- Number of possible states usually enormous
 - e.g. problem 2 catalysis: 100 sites, each could be in any of 4 states: $4^{100} \sim 10^{60}$ distinct states
 - M is even bigger (4¹⁰⁰)²
- Different values of all these P's at each time step!

So usually we cannot compute all the elements of P, and we cannot sample even a small fraction of all the possible states <u>N</u>

- Common approach: sample from <u>P(t)</u> using Kinetic Monte Carlo (Gillespie algorithm).
 - Each trajectory requires computing $2*(t_f-t_o)/\Delta t$ random numbers.
 - Low-probability states will not be sampled at all.
 - $N^{-1/2}$ scaling: hard to achieve high precision
- Decide on which quantities <f> you are trying to compute before you start
 - Can compute several at same time "on the fly".
 - May not be able to store all the trajectories.

What is the initial condition \underline{P}_{o} ?

- Often have an idea of <<u>N</u>> but not full probability distribution of <u>N</u>.
- Common approach: assume N_A, N_B, etc. uncorrelated initially, and use a separate Poisson distribution for each one:

$$P_{poisson}(N) = \langle N \rangle^{N} \exp(-\langle N \rangle) / N!$$

 $\mathsf{P}_{\mathsf{o}}(\mathsf{N}_{\mathsf{A}},\mathsf{N}_{\mathsf{B}},\ldots)=\mathsf{P}_{\mathsf{poisson}}(\mathsf{N}_{\mathsf{A}})^*\mathsf{P}_{\mathsf{poisson}}(\mathsf{N}_{\mathsf{B}})^*\ldots$

When you start each KMC trajectory, you need to sample from a Poisson distribution for N_A , N_B , etc.

Which processes to include in Kinetic Monte Carlo?

- Accelerate calculation by making ∆t as long as possible, i.e. omit very fast processes (which will be "equilibrated" on long time scale).
- The low-probability processes are not going to be sampled adequately, so consider omitting them completely if it speeds calculation.
- "Adequate sampling": statistical sampling errors ~ sqrt(number of samples with positive result).

For example, in Problem 2....

- If the main reaction A+B is very fast, then you will get a lot of trajectories showing this reaction, and can get pretty good sampling statistics.
- On the other hand, if the coking reaction is slow, you may not sample enough trajectories that show a coking event to be able to reach any conclusions about it.
- If diffusion is too fast, ∆t will be very small and it will be very expensive to compute a trajectory.
 - Sometimes people assume diffusion is equilibrated and so create a different random distribution of A and B on surface at each time step corresponding to (slow) reactions

Molecular Dynamics

- What is it: Solving motions of atoms or clumps of atoms using Newton's equations of motion.
 - Usually fitted force fields but can be from DFT
 - Can include quantum effects e.g. by RPMD
 - Typically use Velocity Verlet algorithm to integrate the ODEs (conserves energy to high accuracy)
 - Typically use a "thermostat" to capture energy fluctuations due to contact with a thermal bath
- Alternative to Metropolis Monte Carlo for steady-state properties of molecules
- Can be used for computing time-dependent properties (it is a more-or-less exact simulation of what the molecules are really doing).

Limitations of Molecular Dynamics

- For molecules the stretching vibrations are very fast, so need to use very small Δt
- This usually limits (t_f-t_o) to nanoseconds
- So only can determine static quantities which have equilibrated faster than nanoseconds
- Only can follow dynamic processes which occur in nanoseconds.
- Issue with sampling over large number of possible initial conditions.

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10.34 Numerical Methods Applied to Chemical Engineering Fall 2015

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