Markov chain Monte Carlo and Perfect simulation Lecture at Aristotle University of Thessaloniki

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Introduction

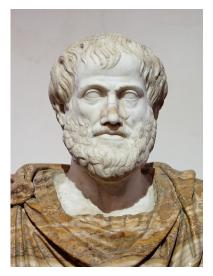


Figure 1: Αριστοτέλης 384–322 BCE

Aristotle: "The more you know, the more you know you don't know."

Sketch of MCMC (I)



Figure 2: Edward Teller (1908-2003)

The original Markov chain Monte Carlo method (MCMC) was introduced by Metropolis *et al.* (1953). The senior author was Edward Teller ("father of the H-bomb").

[Fermi once said,] Teller was the only monomaniac he knew who had several manias: see Brown & May (2004).

Sketch of MCMC (II)

- Markov chain basics:
 - Transition probabilities p(a,b) (or transition rates in continuous time: unified view using exponential distribution);
 - ► Equilibrium probabilities $\pi(a)$, balance, detailed balance $\pi(a)p(a,b) = \pi(b)p(b,a)$, reversibility;
 - ▶ Aperiodicity, recurrence (and uniform and geometric recurrence);
 - under detailed balance we can condition by forbidding transitions;
- We can modify any chain, transition probabilities p(a,b), to leave a specified target distribution $\pi(a)$ invariant, by censoring each possible transition $a \to b$ with probability $\alpha(a,b) \in [0,1]$ such that $\frac{\alpha(a,b)\pi(a)p(a,b)}{\alpha(a,b)\pi(a)p(a,b)} = \frac{\alpha(b,a)\pi(b)p(b,a)}{\alpha(a,b)\pi(a)}$;
- Common choice: Metropolis-Hastings $\alpha(a,b) = \min\{1, (\pi(b)p(b,a))/(\pi(a)p(a,b))\}.$
- If result still irreducible aperiodic, then $\pi(a)$ is its long-term equilibrium.
- This is MCMC, now of intense interest to statisticians.
- But, physicists always remind us, physicists got there fifty years earlier!



Sketch of MCMC (III)

Given the $\pi(a)$, how to design a Markov chain to have this as equilibrium?

- Independence sampler: draw from a fixed probability distribution, apply Metropolis-Hastings;
- Random walk Metropolis or RWM: propose move using a random walk, apply Metropolis-Hastings;
- **Metropolis-adjusted Langevin** or MALA: make a Gaussian jump shifted using gradient of $\log \pi$, apply Metropolis-Hastings.

Can mix-and-match! RWM is often favourite: flexible, not too complicated.

Issues:

- Burn-in: How long till approximate equilibrium?
- Scaling: How big should be the RWM jump?

Question (B) is about how to get fast mixing. There is a beautiful and useful theory, but that is for another day.

Question (A) is what this lecture is all about.



Sketch of MCMC (IV)

- MCMC practicalities: Burn-in: what to do about it?
 - ▶ Theory tends to be much too pessimistic. Example: Zanella (2015a, 2015b) developed statistical methods for Anglo-Saxon history: a simplified model appeared to converge approximately in 10⁵ steps (about 1 week on compute cluster), versus 10⁹ steps in theory (around 2 centuries);
 - ▶ Is (a) one long run better or (b) many short runs? (Option (b) requires starts of short runs spread "evenly" over the sample space almost as hard in high dimensions as the original problem!)
 - ▶ Diagnostics? (Meta-theorem: for any diagnostic technique there is a chain for which the technique is deceptive!)
 - ➤ Conclusion: effective MCMC requires very careful thought about appropriate length of run think deeply about the problem!
- Can there ever be a better way?

Perfect Simulation

- The idea of exact simulation / Coupling from the Past (CFTP) / perfect simulation Propp & Wilson (1996) (Persi Diaconis: "Like seeing the landscape of Mars for the first time");
- Ideas (of "classic CFTP"):
 - extending simulation backwards through time,
 - exploit monotonicity by coupling maximal and minimal processes,
 - seek coalescence;
- Details for *random-walk-CFTP*, which can be boosted as above to provide simple image reconstruction of an image using Ising model, Propp & Wilson (1996) show how to vary a clever algorithm to get exact samples for critical Ising model (this is what impressed Diaconis);
- "Perfect simulation" (WSK, 1998): because everyone knows it isn't going to be perfect, whereas people might imagine "exact simulation" would somehow miraculously defeat numerical approximation error :-).

An example and some theory

- An intensely visual example, which helps many people see intuitively
 what is going on here, is *DeadLeaves-CFTP* (WSK & Thönnes, 1999)
 (technically, *Occluded CFTP*);
- What about cases where monotonicity fails? or there isn't a sensible "maximal" process? WSK (1998):
 - cross-couple upper and lower envelope processes,
 - ▶ dominate by amenable "dominating process" (time-reversible, can draw from equilibrium, can couple target processes below dominating process);
- Theoretical limits: in principle
 - ► Classical CFTP equivalent to uniform ergodicity (Foss & Tweedie, 1998).
 - ▶ *Dominated CFTP* is achievable under geometric ergodicity (WSK, 2004).
 - ▶ It is even possible to carry out Dominated CFTP in some **non**-geometrically ergodicity cases [Connor & WSK (2007); *nb* corrigendum];
- We can use *Dominated CFTP* to carry out perfect simulation for stable point processes (WSK & Møller, 2000);
- Detailed expositions are given by WSK (2005), Huber (2015). WSK (2015) shows how to implement CFTP in R.

Applications to Queues and Epidemics



Figure 3: An illustration introducing both queues and epidemics!

Perfect Queues

The simplest queuing model (Poisson arrivals, exponential service times, single server) can be analyzed very throughly;

- Poisson arrivals are not unreasonable, but exponential service times are ludicrous. Fortunately the case of general service time can use the "embedded chain" (sample at instants of departure), if just 1 server;
- Multi-server case: computation of eg waiting-time distribution is out of reach so use simulation (and insights from Kiefer & Wolfowitz, 1955);
- Sigman (2011) shows how to do CFTP in the "super-stable" case (traffic so low that it could have been handled by just one server), using Dominated CFTP and comparing to a "Processor-sharing" discipline.
- Connor & WSK (2015) show how to extend Sigman (2011), showing how Dominated CFTP can be applied to simulate (sub-critical!) queues perfectly (and this has now been generalized by others to the case of non-Poissonian inter-arrival times). (Technical point: pathwise domination requires service times to be assigned in order of commencement of service!)

Perfect Epidemics

- Even the simple case of deterministic S-I-R permits only *partial* closed-form solution;
- Suppose one can observe **only** the "removal times"? Can one do perfect simulation of the infection times? (Given basic parameters.)
- YES: work in progress by Connor and Kendall. Here is a GIF illustrating this for a real-life small-pox epidemic;
- (be clear about assumptions!)
- (indicate how the perfect simulation algorithm can be used as a high-dimensional integration device to enable simulation-based Bayesian inference!).

Conclusion

- You don't always have to put up with burn-in issues when doing MCMC;
- CFTP works even for significantly complex and relevant models of real-life phenomena;
- Of course really detailed models are still going to resist perfect simulation: but it will always be helpful to compare with a simpler model (using fewer parameters!).
- CFTP is clearly an important tool to be considered by the investigator seeking to do accurate and informative MCMC.

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Queues	https://covidposters.github.io/	Open source
Epidemic	Result of code written by WSK	CC BY-SA 3.0

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Still this early version.

