Advanced computational methods X071521-Selected Topics: MCMC

In the following, we will focus on Markov Chain-Monte Carlo (MCMC) algorithms.

1 Motivation and foundation

1.1 Motivation

In many applications, we care about a probability distribution ν which cannot be written out or dealt with directly. For example, in Bayesian inference or data assimilation, we care about the posterior distribution P(u|y), where y is some observed data.

One possible approach is to use so-called empirical measure

$$\nu \approx \frac{1}{N} \sum_{n=1}^{N} \delta(x - X^i).$$

Such an approach is called **Monte Carlo** sampling $(X^i \sim \nu \text{ i.i.d})$. According to law of large numbers, as $N \to \infty$, the empirical measure can converge in weakly to the measure ν (In fact, also in some metric spaces, like Wasserstein spaces). Traditional sampling methods include rejection sampling (acceptance-rejection method), importance sampling.

However, often the exact sampling from ν is not possible. If this is the case, we can choose to generate a artificial Markov chain to get some approximate samples. Such a type of method will be called **Markov chain** Monte Carlo(MCMC).

- In data assimilation, we have some observations of a dynamical variables. We hope to know more of the dynamic variable using the observed data. As mentioned, MCMC can be used for sampling posterior distribution in data assimilation. This in fact corresponds to a class of methods, called smoothing algorithms in data assimilation. Often, people care about online algorithms and prediction, then another class of algorithms in data assimilation is the filtering algorithms. If we have time, we will touch the basic Kalman filter later.
- In Bayesian inference, it is assumed there is some hidden parameters θ to govern the distributions of data. We then use the observed data to get more estimates of θ using the posterior distribution. Clearly,

MCMC can also be used for Bayesian inference. Another popular class of algorithms is the variational inference, where we find a member from a family of distributions that is closet to the posterior distribution.

• Besides data assimilation and Bayesian inference, MCMC can also used for sampling some complicated probability measures, like the Gibbs measure $\exp(-V(x))$. This is particularly important in molecular dynamics, where the free energy surface V is unknown (we can compute, however, $-\nabla V(x)$ for each given x through some algorithms).

1.2 The foundation of MCMC

We aim to find a Markov chain such that the desired measure ν is invariant under the dynamics, and the law of the chain converges to ν . This is related to the so-called *ergodicity*. If we have ergodicity, then the running time average of the chain converges to average with respect to ν .

To be precise, we have

Theorem 1. Assume that ν is an invariant measure of a Markov chain X^n valued in \mathbb{R}^d with Lebesgue density ρ . If the chain is ergodic, then for every bounded continuous function φ , we have

$$\frac{1}{N}\sum_{n=1}^{N}\varphi(X^n) \to \mathbb{E}^{\nu}\varphi(X), \ a.s.$$

for a. s. $X^0 \sim \nu$. If moreover the Doeblin's condition holds: there exists a probability measure p such that for all $x \in \mathbb{R}^d$, and all Borel set A, we have $P(x, A) \geq \epsilon p(A)$, then

$$TV(P^n(x,\cdot),\nu) \le 2(1-\epsilon)^n.$$

Further,

$$\frac{1}{N}\sum_{n=1}^{N}\varphi(X^n) - \mathbb{E}^{\nu}\varphi(X) = K\xi_N N^{-1/2}$$

where ξ_N converges in law to $\mathcal{N}(0,1)$.

Above, P(x, A) is the transition measure. This theorem guarantees the convergence MCMC, and characterizes the convergence rate.

2 A basic MCMC: Metropolis-Hastings method

A very basic and widely used MCMC algorithm is the Metropolis-Hastings method. In this method, an accept/reject step is added to ensure the invariance with respect to ν .

In fact, a sufficient condition that enforces invariance is the **detailed** balance condition. Suppose p(x, z) is the transition density so that

$$P(x,A) = \int_A p(x,z) \, dz.$$

We say the detailed balance condition holds if there exists a density function ρ such that

$$\rho(x)p(x,y) = \rho(y)p(y,x), \ \forall x, y \in \mathbb{R}^d.$$

This basically means the mass moved from x to y equals the mass moved from y to x. Using this detailed balance condition, we find easily that ρ is the density of an invariant measure:

$$\rho(x) = \int \rho(x) p(x, y) \, dy = \int \rho(y) p(y, x) \, dy.$$

The idea of Metropolis-Hastings is then to guarantee the detailed balance.

Suppose we have a Markov chain with transition kernel q(x, y). Define

$$a(x,y) = 1 \land \frac{\rho(y)q(y,x)}{\rho(x)q(x,y)}$$

Then, the algorithm is as follows:

- 1. Choose $X^0 \in \mathbb{R}^d$.
- 2. For $n \ge 1$, do
 - Draw $Y^n \sim q(X^{n-1}, \cdot)$ by running the given Markov chain.
 - Choose $X^{(n)}$ such that $\mathbb{P}(X^n = Y^n) = a(X^{n-1}, Y^n)$ and $\mathbb{P}(X^n = X^{n-1}) = 1 a(X^{n-1}, Y^n)$.

Clearly, $\{X^n\}$ is a Markov chain and this will approximate the probability measure $\nu = \rho dx$.

We have the following

Lemma 1.

 $The \ Metropolis-Hastings \ MCMC \ ensures \ the \ detailed \ balance \ with \ transition \ kernel$

$$p(x, y) = q(x, y)a(x, y).$$

The claim is trivial and we omit. In fact, according to the detailed balance condition, the acceptance function should satisfy

$$\frac{a(x,y)}{a(y,x)} = \frac{\rho(y)q(y,x)}{\rho(x)q(x,y)}.$$

The function a is not unique. One has other choices.

Remark 1. Above, we mentioned the algorithm for continuous state space. If the state space is discrete, the algorithm can be modified correspondingly easily. The ergodicity can be verified by more easily checked conditions: aperiodicity and positive recurrence. Of course, the Doeblin's condition is still a sufficient condition for geometric ergodicity.

Remark 2. In many applications, we only know ρ up to a multiplicative factor, such as the Gibbs measure

$$\rho \propto \exp(-V(x)).$$

The M-H algorithm only requires the ratio so we do not have to compute the normalization constant.

The convergence of Metropolis-Hastings follows from the general theorem above. We need to verify the conditions. For details, one can refer to, for example, "Rates of convergence of the Hastings and Metropolis algorithms" by Mengersen and Tweedie.

2.1 An illustrating example

Suppose that we want to sample from the exponential distribution:

$$\rho(x) = 1_{x>0} \exp(-x).$$

Assume that we know already how to sample from a normal distribution. We then define the transition probability as

$$q(x,y) = \frac{1}{\sqrt{2\pi}} \exp(-\frac{(y-x)^2}{2}).$$

In other words,

$$Y^*|X^n \sim \mathcal{N}(X^n, 1) = X^n + \mathcal{N}(0, 1).$$

Since q(x, y) = q(y, x), we find

$$A = \frac{\rho(Y^*)}{\rho(X^n)}$$

Note that since X^n has already been accepted, $X^n > 0$.

Then, we generate $z \sim U(0, 1)$. If $z \leq A$, we set

$$X^{n+1} = Y^*;$$

otherwise

$$X^{n+1} = X^n$$

2.2 extensions

There are many extensions of M-H algorithms. For example, the slice sampling.

3 Gibbs sampler

This was introduced by brothers of Stuart and Geman in 1984. Gibbs sampler is commonly used for Bayesian inference. Compared with the traditional EM (expectation-maximization) algorithm, it is a stochastic one.

This is applicable when the joint distribution is not known or hard to sample but the conditional distribution (not marginal) is easier to sample from.

3.1 The basic version

Suppose we want to sample a multivariate distribution $\rho(Z) = \rho(Z_1, Z_2, \dots, Z_M)$, we know all the marginal distributions. Suppose we have already sampled Z^n , we do the following to get Z^{n+1} :

- $Z_1^{n+1} \sim p(z_1 | Z_2^n, Z_3^n, \dots, Z_M^n).$
- $Z_2^{n+1} \sim p(z_2 | Z_1^{n+1}, Z_3^n, \dots, Z_M^n).$
- $Z_3^{n+1} \sim p(z_3 | Z_1^{n+1}, Z_2^{n+1}, \dots, Z_M^n).$
- ...
- $Z_M^{n+1} \sim p(z_M | Z_1^{n+1}, \dots, Z_{M-1}^{n+1}).$

We first of all claim that the joint distribution is invariant. Consider the jump from (Z_j^m, Z_i^n) to (Z_j^m, Z_i^{n+1}) . The transition probability is given by

$$q(x, y|Z_j^m) = p(y|Z_j^m).$$

The desired distribution is the joint distribution $\rho(Z)$. The detailed balance hods for this ρ

$$\rho(Z_j^m, x)q(x, y|Z_j^m) = \rho(Z_j^m, y)q(y, x|Z_j^m).$$

This is found easily by the definition of the conditional distribution $p(y|Z_i^m)$.

With the detailed balance, we find that this basic version can be viewed as a special case of Metropolis-Hastings. In fact, the function $a(x, y|Z_j^m)$ is always 1 and the samples are always accepted. This can therefore be regarded as an MH algorithm. However, its extensions can be more general.

When n is large enough, the distribution of Z^n is close to the joint distribution. In practice, one may disregard some initial samples. Also, we may only want every n_1 th sample. The first is because the distribution will be close to the stationary distribution only after some steps. The second is considered because the adjacent data are usually not independent.

3.2 Some extensions

The obvious extension is to consider a stochastic version of the Gibbs sampler:

- We choose an index i randomly from $\{1, 2, \ldots, M\}$
- We sample $\xi \sim p(z|Z_{-i})$, where Z_{-i} means the vector by deleting Z_i .
- Set $Z_i = \xi$. In other words, the new random variable is

$$Z = (Z_1, \ldots, Z_{i-1}, \xi, Z_{i+1}, \ldots, Z_M).$$

Other variants include blocked Gibbs sampling, in which we may sample several Z_i 's by conditioning on others.

Another variant is the collapsed Gibbs sampling. Imagine that we want to sample from p(x, y|z). In the Gibbs sampler above, we can do $x^{n+1} \sim p(x|y^n, z)$ and $y^{n+1} \sim p(y|x^{n+1}, z)$. In the collapsed version, we do the following

$$y^{n+1} \sim p(y|z), \quad x^{n+1} \sim p(x|y^{n+1}, z).$$

Here, $p(y|z) = \int p(x, y|z) dx$. Note that the one with one variable integrated must be sample first. Otherwise, the distribution we are sampling from is not correct.

4 Hamiltonian MCMC

Clearly, in the running average, the two adjacent data are not independent. If we run Metropolis-Hastings, we need the time to be longer than the correlation time (the correlation between X^n and X^{n+m} decays exponentially with m. Hence, we need m bigger than some time that ensures the correlation to be smaller than a given value.)

The Hamiltonian MCMC is designed by Duane, Kennedy, et al in 1987 to reduce the correlation using a Hamiltonian evolution. It also allows a higher acceptance rate. Interestingly, the original name by the authors is 'Hybrid Monte Carlo', which is 'HMC' in short. Nowadays, it is better known as 'Hamiltonian Monte Carlo', which is also 'HMC' in short.

For more details, see review articles "MCMC using Hamiltonian dynamics", "A conceptual introduction to Hamiltonian Monte Carlo".

4.1 Motivation and idea

In statistical mechanics, it is well-known that a system with Hamiltonian H(p,q) that interacts with the surrounding heat bath could result in the the canonical distribution:

$$\rho(p,q) \propto \exp(-\beta H(p,q)),$$

in the thermoequilibrium. Often, the Hamiltonian is given by

$$H(p,q) = \frac{p^2}{2m} + U(q),$$

where the first term is the kinetic energy while the second is the potential energy. The marginal distribution on q is therefore

$$\exp(-\beta U(q)).$$

Hence, if we choose

$$U(q) = -\frac{1}{\beta} \log \rho(q),$$

we can run the dynamics of the system to obtain the desired distribution $\rho(q)$.

This motivates the HMC and Langevin MC. In HMC, the system is closed and evolves according to Hamilton ODEs while in Langevin dynamics, we have whilte noise and friction that describe the interaction with heat bath.

4.2 The Hamiltonian system and its discretization

In the Hamilton mechnics, H(p,q) is the energy functional. The dynamics is given by

$$\dot{q}_i = \frac{\partial H}{\partial p_i},$$
$$\dot{p}_i = -\frac{\partial H}{\partial q_i}$$

It is convenient to introduce the matrix

$$J = \left[\begin{array}{cc} 0 & I \\ -I & 0 \end{array} \right].$$

The dynamics is written as

$$\frac{dz}{dt} = J\nabla_z H,$$

where $z = [q, p]^T$.

Often, the Hamiltonian is given by

$$H(p,q) = U(q) + K(p),$$

and

$$K(p) = \frac{1}{2}p^T M^{-1}p,$$

and $M = diag(m_1, \ldots, m_d)$. There are several properties

- Reversible. According to the dynamics, if the state goes from (q_0, p_0) to (q_1, p_1) , we negate p and then have $(q_1, -p_1)$. From this state, after the same time, it will arrive at $(q_0, -p_0)$.
- The Hamiltonian is conserved. $\frac{d}{dt}H(p(t),q(t)) = 0.$
- Volume-preserving. The dynamics is $\frac{dz}{dt} = V(z) := J\nabla_z H$. We can verify easily that

$$\nabla_z \cdot V(z) = 0.$$

• Symplectic. Let the flow map be T_t and the corresponding Jacobi matrix be B_t . Then,

$$B_t^T J^{-1} B_t = J^{-1}.$$

To discretize, the naive idea is to use forward Euler,

$$p_{n+1} = p_n - \tau \nabla_q U(q),$$
$$q_{n+1} = q_n + \tau \frac{p_n}{m}$$

This scheme, however, behaves poorly in practice. The reason is that the volume in phase space is not preserved, and the Hamiltonian is not preserved.

A simple modified scheme

$$p_{n+1} = p_n - \tau \nabla_q U(q_n),$$
$$q_{n+1} = q_n + \tau \frac{p_{n+1}}{m_i}$$

behaves much better. The reason is that it preserves the volume. In fact, this is related to a large class of methods, called **symplectic methods** for Hamiltonian discretizations.

The following leapfrog method, which shares some flavor of Strang splitting, is as follows

$$\begin{split} p_{n+1/2} &= p_n - (\tau/2) \nabla_q U(q_n), \\ q_{n+1} &= q_n + \tau \frac{p_{n+1/2}}{m_i}, \\ p_{n+1} &= p_{n+1/2} - (\tau/2) \nabla_q U(q_{n+1}). \end{split}$$

This is a symplectic method.

4.3 Hamiltonian MC

The motivation comes from the canonical distribution in statistical mechanics. In the thermo-equilibrium, the probability that a configuration appears is

$$p(q,p) = \frac{1}{Z} \exp(-H(q,p)).$$

Of course, the canonical distribution is not for a Hamiltonian system, because for a closed system, the energy is conserved. This is for a system in a heat bath. The system has its own Hamitonian, but it also has interaction with the environment so that its energy can be changed. Anyway, this gives us the motivation to use the Hamilton system for sampling.

Suppose we want to sample from $\rho(q)$. Then, we construct a Hamiltonian

$$H(q, p) = -\log \rho(q) + K(p) =: U(q) + K(p).$$

Then, the marginal distribution of the canonical distribution is exactly what we want. Hence, we have the following HMC

- Sample $p_n \sim p(p|q_n) \propto \exp(-\frac{|p|^2}{2})$.
- Here, we apply Metropolis update. Run the Hamiltonian dynamics from state (q_n, p_n) to (q_{n+1}^*, p^*) using the leapfrog scheme, with step size τ for *L* steps. (It other words, we have 'time' consumed to be $L\tau$). Then, we negate *p* so that the proposed state is (q_{n+1}^*, p_{n+1}^*) . With probability

$$\min(1, \exp(-H(q_{n+1}^*, p_{n+1}^*) + H(q_n, p_n)))$$

we accept the state and then $q_{n+1} = q_{n+1}^*$. Otherwise, $q_{n+1} = q_n$.

- **Remark 3.** In the Metropolis update, since the dynamics is reversible, so q(x, y) = q(y, x) and hence, the function a is determined totally by the desired canonical distribution (Gibbs distribution).
 - Since the Hamiltonian is symmetric about the momentum, we do not really need to negate the momentum p^* .
 - The resampling of p at the beginning is very important. Otherwise, the distribution of q will not range over the desired distribution $\exp(-U(q))$

To verify that the method works, one should check that the canonical distribution is invariant under the process. This is straightforward to see.

A second thing is to gurantee the ergodicity. This is not very clear for HMC. In fact, in some cases, the chain is not aperiodic so that it is not ergodic. However, this is very rare to happen. However, to make sure the ergodicity, one may make the running time random. We will not go into details for these issues.

5 Langevin MCMC

In Hamiltonian MCMC, there is no noise. If we consider the dynamical system with white noise, we can have the following system of equation

$$dx = v \, dt,$$

$$m dv = -\gamma v \, dt - \nabla_x U(x) \, dt + \sqrt{2\gamma/\beta} dW,$$

where v = p/m. The distribution can be proven to converge to the Gibbs measure

$$\propto \exp(-\beta(U(x) + \frac{m|v|^2}{2}))$$

The marginal distribution is what we want.

We can also consider the overdamped regime:

$$dx = -rac{1}{\gamma}
abla_x U \, dt + \sqrt{rac{2}{\gammaeta}} dW$$

The invaraint measure of this SDE is

$$\propto \exp(-\beta U(x))$$

Hence, we can use both to do the MCMC.

For the exponential convergence of the overdampled case for sampling and the Metropolis-adjusted discretization, you may read "Exponential convergence of Langevin distributions and their discrete approximations" and "Sampling from a strongly log-concave distribution with the unadjusted Langevin algorithm". For the underdamped one (with momentum), the reference is "Underdamped Langevin MCMC: a nonasymptotic analysis".

5.1 A version of Langevin MCMC: the overdamped case

We take the temperature parameter to be $\beta = 1$, and consider the following SDE

$$dX = -\nabla U \, dt + \sqrt{2} \, dB.$$

Recall that

$$U = -\log \rho.$$

We hope U to be strongly convex so that we need ρ to be log concave.

A version of simple Langevin MC is therefore given by the Euler-Maruyama scheme

$$X^{n+1} = X^n - \gamma_{n+1} \nabla U(X^n) + \sqrt{2\gamma_{n+1}} Z_{n+1}$$

where $Z_{n+1} \sim \mathcal{N}(0, 1)$.