# Statistical Models & Computing Methods

## Lecture 6&7: Markov Chain Monte Carlo



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- ▶ Direct sampling in high-dimensional spaces is often infeasible, very hard to get rare events
- ► Rejection sampling, Importance sampling
  - ▶ Do not work well if the proposal q(x) is very different from f(x) or h(x)f(x).
  - Moreover, constructing appropriate q(x) can be difficult. Making a good proposal usually requires knowledge of the analytic form of the target distribution - but if we had that, we wouldn't even need to sample
- ▶ Intuition: instead of a fixed proposal q(x), what if we use an adaptive proposal?
- ► In this lecture, we are going to talk about one of the most popular sampling methods, Markov chain Monte Carlo.



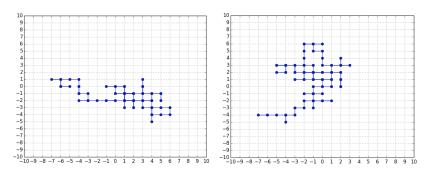
- ▶ Stochastic processes is a family of random variables, usually indexed by a set of numbers (time). A discrete time stochastic process is simply a sequence of random variables,  $X_0, X_1, \ldots, X_n$  defined on the same probability space
- ▶ One of the simplest stochastic processes (and one of the most useful) is the simple random walk
- Consider a simple random walk on a graph  $G = (\Omega, E)$ . The stochastic process starts from an initial position  $X_0 = x_0 \in \Omega$ , and proceeds following a simple rule:

$$p(X_{n+1}|X_n=x_n) \sim \text{Discrete}(\mathcal{N}(x_n)), \ \forall n \geq 0$$

where  $\mathcal{N}(x_n)$  denotes the neighborhood of  $x_n$ 



### Two random walks on a $20\times20$ grid graph



- ► The above simple random walk is a special case of another well-known stochastic process called *Markov chains*
- A Markov chain represents the stochastic movement of some particle in the state space over time. The particle initially starts from state i with probability  $\pi_i^{(0)}$ , and after that moves from the current state i at time t to the next state j with probability  $p_{ij}(t)$
- ► A Markov chain has three main elements:
  - 1. A state space S
  - 2. An initial distribution  $\pi^{(0)}$  over  $\mathcal{S}$
  - 3. Transition probabilities  $p_{ij}(t)$  which are non-negative numbers representing the probability of going from state i to j, and  $\sum_{j} p_{ij}(t) = 1$ .
- When  $p_{ij}(t)$  does not depend on time t, we say the Markov chain is time-homegenous

► Chain rule (in probability)

$$p(X_n = x_n, \dots, X_0 = x_0) = \prod_{i=1}^n p(X_i = x_i | X_{\le i} = x_{\le i})$$

► Markov property

$$p(X_{i+1} = x_{i+1}|X_i = x_i, \dots, X_0 = x_0) = p(X_{i+1} = x_{i+1}|X_i = x_i)$$

▶ Joint probability with Markov property

$$p(X_n = x_n, \dots, X_0 = x_0) = \prod_{i=1}^n p(X_i = x_i | X_{i-1} = x_{i-1})$$

fully determined by the transition probabilities

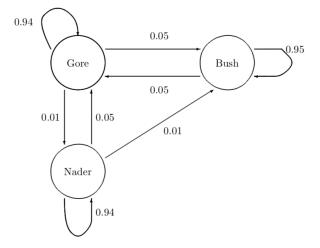


- ► Consider the 2000 US presidential election with three candidates: Gore, Bush and Nader (just an illustrative example and does not reflect the reality of that election)
- ▶ We assume that the initial distribution of votes (i.e., probability of winning) was  $\pi = (0.49, 0.45, 0.06)$  for Gore, Bush and Nader respectively
- ► Further, we assume the following transition probability matrix

	Gore	Bush	Nader
Gore	0.94	0.05	0.01
Bush	0.05	0.95	0
Nader	0.05	0.01	0.94



A probabilistic graph presentation of the Markov chain



▶ If we represent the transition probability a square matrix P such that  $P_{ij} = p_{ij}$ , we can obtain the distribution of states in step n,  $\pi^{(n)}$ , as follows

$$\pi^{(n)} = \pi^{(n-1)}P = \dots = \pi^{(0)}P^n$$

► For the above example, we have

$$\pi^{(0)} = (0.4900, 0.4500, 0.0600)$$

$$\pi^{(10)} = (0.4656, 0.4655, 0.0689)$$

$$\pi^{(100)} = (0.4545, 0.4697, 0.0758)$$

$$\pi^{(200)} = (0.4545, 0.4697, 0.0758)$$

- ► As we can see last, after several iterations, the above Markov chain converges to a distribution, (0.4545, 0.4697, 0.0758)
- ▶ In this example, the chain would have reached this distribution regardless of what initial distribution  $\pi^{(0)}$  we chose. Therefore,  $\pi = (0.4545, 0.4697, 0.0758)$  is the stationary distribution for the above Markov chain
- ▶ Stationary distribution. A distribution of Markov chain states is called to be stationary if it remains the same in the next time step, i.e.,

$$\pi = \pi P$$



- ▶ How can we find out whether such distribution exists?
- ▶ Even if such distribution exists, is it unique or not?
- ▶ Also, how do we know whether the chain would converge to this distribution?
- ► To find out the answer, we briefly discuss some properties of Markov chains



- ► Irreducible: A Markov chain is irreducible if the chain can move from any state to another state.
- ► Examples
  - ► The simple random walk is irreducible
  - ► The following chain, however, is reducible since Nader does not communicate with the other two states (Gore and Bush)

	Gore	Bush	Nader
Gore	0.95	0.05	0
Bush	0.05	0.95	0
Nader	0	0	1



- ▶ Period: the period of a state i is the greatest common divisor of the times at which it is possible to move from i to i.
- ► For example, all the states in the following Markov chain have period 3.

$$\begin{pmatrix} 0 & 1 & 0 \\ 0 & 0 & 1 \\ 1 & 0 & 0 \end{pmatrix}$$

▶ Aperiodic: a Markov chain is said to be aperiodic if the period of each state is 1, otherwise the chain is periodic.



▶ Recurrent states: a state i is called recurrent if with probability 1, the chain would ever return to state i given that it started in state i.

	Gore	Bush	Nader
Gore	0.94	0.05	0.01
Bush	0.05	0.95	0
Nader	0.05	0.01	0.94

- **Positive recurrent**: a recurrent state j is called positive recurrent if the expected amount of time to return to state j given that the chain started in state j is finite
- ► For a positive recurrent Markov chain, the stationary distribution exists and is unique



- ▶ Reversibility: a Markov chain is said to be reversible with respect to a probability distribution  $\pi$  if  $\pi_i p_{ij} = \pi_j p_{ji}$
- ▶ In fact, if a Markov chain is reversible with respect to  $\pi$ , then  $\pi$  is also a stationary distribution

$$\sum_{i} \pi_{i} p_{ij} = \sum_{i} \pi_{j} p_{ji}$$
$$= \pi_{j} \sum_{i} p_{ji}$$
$$= \pi_{j}$$

since  $\sum_{i} p_{ji} = 1$  for all transition probability matrices

► This is also known as *detailed balance condition* 



- ▶ We can define a Markov chain on a general state space  $\mathcal{X}$  with initial distribution  $\pi^{(0)}$  and transition probabilities p(x, A) defined as the probability of jumping to the subset A from point  $x \in \mathcal{X}$
- ► Similarly, with Markov property, we have the joint probability

$$p(X_0 \in A_0, \dots, X_n \in A_n) = \int_{A_0} \pi^{(0)}(dx_0) \dots \int_{A_n} p(x_{n-1}, dx_n)$$

Example. Consider a Markov chain with the real line as its state space. The initial distribution is  $\mathcal{N}(0,1)$ , and the transition probability is  $p(x,\cdot) = \mathcal{N}(x,1)$ . This is just a Brownian motion (observed at discrete time)



- ▶ Unlike the discrete space, we now need to talk about the property of Markov chains with a continuous non-zero measure  $\phi$ , on  $\mathcal{X}$ , and use sets A instead of points
- ▶ A chain is  $\phi$ -irreducible if for all  $A \subseteq \mathcal{X}$  with  $\phi(A) > 0$  and for all  $x \in \mathcal{X}$ , there exists a positive integer n such that

$$p^{n}(x, A) = p(X_{n} \in A | X_{0} = x) > 0$$

▶ Similarly, we need to modify our definition of period

 $\blacktriangleright$  A distribution  $\pi$  is a stationary distribution if

$$\pi(A) = \int_{\mathcal{X}} \pi(dx) p(x, A), \quad \forall A \subseteq \mathcal{X}$$

 $\blacktriangleright$  As for the discrete case, a continuous space Markov chain is reversible with respect to  $\pi$  if

$$\pi(dx)p(x,dy) = \pi(dy)p(y,dx)$$

- ▶ Similarly, if the chain is reversible with respect to  $\pi$ , then  $\pi$  is a stationary distribution
- ▶ Example. Consider a Markov chain on the real line with initial distribution  $\mathcal{N}(1,1)$  and transition probability  $p(x,\cdot) = \mathcal{N}(\frac{x}{2}, \frac{3}{4})$ . It is easy to show that the chain converges to  $\mathcal{N}(0,1)$  (Exercise)

- ightharpoonup Ergodic: a Markov chain is ergodic if it is both irreducible and aperiodic, with stationary distribution  $\pi$
- ▶ Ergodic Theorem. For an ergodic Markov chain on the state space  $\mathcal{X}$  having stationary distribution  $\pi$ , we have: (i) for all measurable  $A \subseteq \mathcal{X}$  and  $\pi$ -a.e.  $x \in \mathcal{X}$ ,

$$\lim_{t \to \infty} p^t(x, A) = \pi(A)$$

(ii)  $\forall f \text{ with } \mathbb{E}_{\pi}|f(x)| < \infty$ ,

$$\lim_{T \to \infty} \frac{1}{T} \sum_{t=1}^{T} f(X_t) = \int_{\mathcal{X}} f(x) \pi(x) dx, \quad \text{a.s.}$$

In particular,  $\pi$  is the unique stationary probability density function for the chain



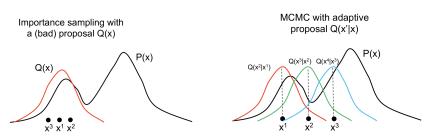
- Now suppose we are interested in sampling from a distribution  $\pi$  (e.g., the unnormalized posterior)
- Markov chain Monte Carlo (MCMC) is a method that samples from a Markov chain whose stationary distribution is the target distribution  $\pi$ . It does this by constructing an appropriate transition probability for  $\pi$
- ► MCMC, therefore, can be viewed as an inverse process of Markov chains







- ► The transition probability in MCMC resembles the proposal distribution we used in previous Monte Carlo methods.
- ► Instead of using a fixed proposal (as in importance sampling and rejection sampling), MCMC algorithms feature adaptive proposals



Figures adapted from Eric Xing (CMU)



- ▶ Suppose that we are interested in sampling from a distribution  $\pi$ , whose density we know up to a constant  $P(x) \propto \pi(x)$
- ▶ We can construct a Markov chain with a transition probability (i.e., proposal distribution) Q(x'|x) which is symmetric; that is, Q(x'|x) = Q(x|x')
- ▶ Example. A normal distribution with the mean at the current state and fixed variance  $\sigma^2$  is symmetric since

$$\exp\left(-\frac{(y-x)^2}{2\sigma^2}\right) = \exp\left(-\frac{(x-y)^2}{2\sigma^2}\right)$$

In each iteration we do the following

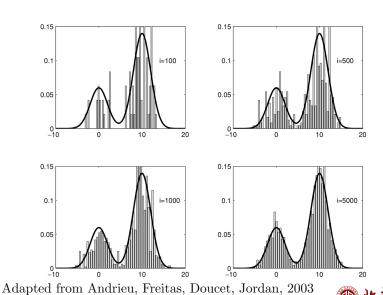
- ▶ Draws a sample x' from Q(x'|x), where x is the previous sample
- ► Calculated the acceptance probability

$$a(x'|x) = \min\left(1, \frac{P(x')}{P(x)}\right)$$

Note that we only need to compute  $\frac{P(x')}{P(x)}$ , the unknown constant cancels out

Accept the new sample with probability a(x'|x) or remain at state x. The acceptance probability ensures that, after sufficient many draws, our samples will come from the true distribution  $\pi(x)$ 





- ▶ How do we know that the chain is going to converge to  $\pi$ ?
- ▶ Suppose the support of the proposal distribution is  $\mathcal{X}$  (e.g., Gaussian distribution), then the Markov chain is irreducible and aperiodic.
- ▶ We only need to verify the detailed balance condition

$$\pi(dx)p(x,dx') = \pi(x)dx \cdot Q(x'|x)a(x'|x)dx'$$

$$= \pi(x)Q(x'|x)\min\left(1,\frac{\pi(x')}{\pi(x)}\right)dxdx'$$

$$= Q(x'|x)\min(\pi(x),\pi(x'))dxdx'$$

$$= Q(x|x')\min(\pi(x'),\pi(x))dxdx'$$

$$= \pi(x')dx' \cdot Q(x|x')\min\left(1,\frac{\pi(x)}{\pi(x')}\right)dx$$

$$= \pi(dx')p(x',dx)$$



▶ It turned out that symmetric proposal distribution is not necessary. Hastings (1970) later on generalized the above algorithm using the following acceptance probability for general Q(x'|x)

$$a(x'|x) = \min\left(1, \frac{P(x')Q(x|x')}{P(x)Q(x'|x)}\right)$$

▶ Similarly, we can show that detailed balanced condition is preserved

- ightharpoonup Under mild assumptions on the proposal distribution Q, the algorithm is ergodic
- ▶ However, the choice of Q is important since it determines the speed of convergence to  $\pi$  and the efficiency of sampling
- ▶ Usually, the proposal distribution depend on the current state. But it can be independent of current state, which leads to an independent MCMC sampler that is somewhat like a rejection/importance sampling method
- ▶ Some examples of commonly used proposal distributions
  - $ightharpoonup Q(x'|x) \sim \mathcal{N}(x,\sigma^2)$
  - $ightharpoonup Q(x'|x) \sim \text{Uniform}(x-\delta,x+\delta)$
- ▶ Finding a good proposal distribution is hard in general



► Recall the univariate Gaussian model with known variance

$$y_i \sim \mathcal{N}(\theta, \sigma^2)$$
$$p(y|\theta, \sigma) = \prod_{i=1}^{n} \frac{1}{\sqrt{2\pi\sigma}} \exp\left(-\frac{(y_i - \theta)^2}{2\sigma^2}\right)$$

- ▶ Note that there is a conjugate  $\mathcal{N}(\mu_0, \tau_0^2)$  prior for  $\theta$ , and the posterior has a close form normal distribution
- ▶ Now let's pretend that we don't know this exact posterior distribution and use a Markov chain to sample from it.

► We can of course write the posterior distribution up to a constant

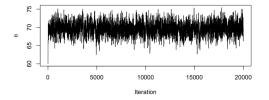
$$p(\theta|y) \propto \exp\left(\frac{(\theta - \mu_0)^2}{2\tau_0^2}\right) \prod_{i=1}^n \exp\left(-\frac{(y_i - \theta)^2}{2\sigma^2}\right) = P(\theta)$$

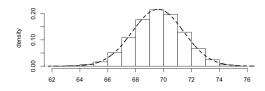
- We use  $\mathcal{N}(\theta^{(i)}, 1)$ , a normal distribution around our current state, to propose the next step
- ▶ Starting from an initial point  $\theta^{(0)}$  and propose the next step  $\theta' \sim \mathcal{N}(\theta^{(0)}, 1)$ , we either accept this value with probability  $a(\theta'|\theta^{(0)})$  or reject and stay where we are
- ▶ We continue these steps for many iterations



# Examples: Gaussian Model with Known Variance 30/75

► As we can see, the posterior distribution we obtained using the Metropolis algorithm is very similar to the exact posterior







▶ Now suppose we want to model the number of half court shots Stephen Curry has made in a game using Poisson model

$$y_i \sim \text{Poisson}(\theta)$$

- ► He made 0 and 1 half court shots in the first two games respectively
- ▶ We used Gamma(1.4, 10) prior for  $\theta$ , and because of conjugacy, the posterior distribution also had a Gamma distribution

$$\theta|y \sim \text{Gamma}(2.4, 12)$$

► Again, let's ignore the closed form posterior and use MCMC for sampling the posterior distribution



► The prior is

$$p(\theta) \propto \theta^{0.4} \exp(-10\theta)$$

► The likelihood is

$$p(y|\theta) \propto \theta^{y_1+y_2} \exp(-2\theta)$$

where  $y_1 = 0$  and  $y_2 = 1$ 

► Therefore, the posterior is proportional to

$$p(\theta|y) \propto \theta^{0.4} \exp(-10\theta) \cdot \theta^{y_1+y_2} \exp(-2\theta) = P(\theta)$$

► Symmetric proposal distributions such as

Uniform
$$(\theta^{(i)} - \delta, \theta^{(i)} + \delta)$$
 or  $\mathcal{N}(\theta^{(i)}, \sigma^2)$ 

might not be efficient since they do not take the non-negative support of the posterior into account.

- ▶ Here, we use a non-symmetric proposal distribution such as Uniform $(0, \theta^{(i)} + \delta)$  and use the Metropolis-Hastings (MH) algorithm instead
- We set  $\delta = 1$

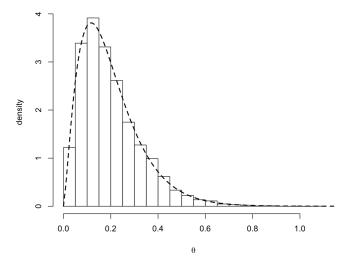
We start from  $\theta_0 = 1$  and follow these steps in each iteration

- ► Sample  $\theta'$  from  $\mathcal{U}(0, \theta^{(i)} + 1)$
- ► Calculate the acceptance probability

$$a(\theta'|\theta^{(i)}) = \min\left(1, \frac{P(\theta') \operatorname{Uniform}(\theta^{(i)}|0, \theta'+1)}{P(\theta^{(i)}) \operatorname{Uniform}(\theta'|0, \theta^{(i)}+1)}\right)$$

▶ Sample  $u \sim \mathcal{U}(0,1)$  and set

$$\theta^{(i+1)} = \begin{cases} \theta' & u < a(\theta'|\theta^{(i)}) \\ \theta^{(i)} & \text{otherwise} \end{cases}$$





- ▶ What if the distribution is multidimensional, *i.e.*,  $x = (x_1, x_2, \dots, x_d)$
- ▶ We can still use the Metropolis algorithm (or MH), with a multivariate proposal distribution, *i.e.*, we now propose  $x' = (x'_1, x'_2, \dots, x'_d)$
- ▶ For example, we can use a multivariate normal  $\mathcal{N}_d(x, \sigma^2 I)$ , or a d-dimensional uniform distribution around the current state

► Here we construct a banana-shaped posterior distribution as follows

$$y|\theta \sim \mathcal{N}(\theta_1 + \theta_2^2, \sigma_y^2), \quad \sigma_y = 2$$

We generate data  $y_i \sim \mathcal{N}(1, \sigma_y^2)$ 

▶ We use a bivariate normal prior for  $\theta$ 

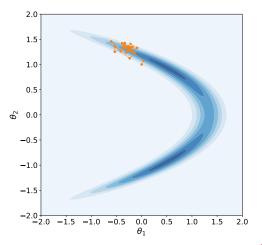
$$\theta = (\theta_1, \theta_2) \sim \mathcal{N}(0, I)$$

► The posterior is

$$p(\theta|y) \propto \exp\left(-\frac{\theta_1^2 + \theta_2^2}{2}\right) \cdot \exp\left(-\frac{\sum_i (y_i - \theta_1 - \theta_2^2)^2}{2\sigma_y^2}\right)$$

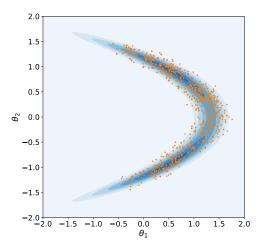
We use the Metropolis algorithm to sample from posterior, with a bivariate normal proposal distribution such as  $\mathcal{N}(\theta^{(i)}, (0.15)^2 I)$ 

The first few samples from the posterior distribution of  $\theta = (\theta_1, \theta_2)$ , using a bivariate normal proposal

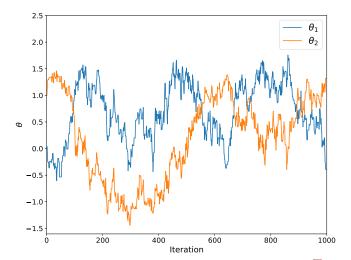




Posterior samples for  $\theta = (\theta_1, \theta_2)$ 



Trace plot of posterior samples for  $\theta = (\theta_1, \theta_2)$ 



- ▶ Sometimes, it is easier to decompose the parameter space into several components, and use the Metropolis (or MH) algorithm for one component at a time
- ▶ At iteration *i*, given the current state  $(x_1^{(i)}, \ldots, x_d^{(i)})$ , we do the following for all components  $k = 1, 2, \ldots, d$ 
  - Sample  $x'_k$  from the univariate proposal distribution  $Q(x'_k|\ldots,x_{k-1}^{(i+1)},x_k^{(i)},\ldots)$
  - ▶ Accept this new value and set  $x_k^{(i+1)} = x_k'$  with probability

$$a(x'_k|\ldots,x_{k-1}^{(i+1)},x_k^{(i)},\ldots)) = \min\left(1,\frac{P(\ldots,x_{k-1}^{(i+1)},x_k',\ldots)}{P(\ldots,x_{k-1}^{(i+1)},x_k^{(i)},\ldots)}\right)$$

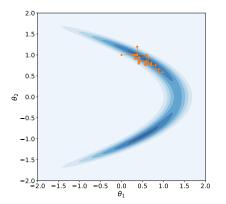
or reject it and set  $x_k^{(i+1)} = x_k^{(i)}$ 



- ▶ Note that in general, we can decompose the space of random variable into blocks of components
- ► Also, we can update the components sequentially or randomly
- ► As long as each transition probability individually leaves the target distribution invariant, their sequence would leave the target distribution invariant
- ▶ In Bayesian models, this is especially useful if it is easier and computationally less intensive to evaluate the posterior distribution when one subset of parameters change at a time



- ▶ In the example of banana-shaped distribution, we can sample  $\theta_1$  and  $\theta_2$  one at a time
- ▶ The first few samples from the posterior distribution of  $\theta = (\theta_1, \theta_2)$ , using a univariate normal proposal sequentially



- ▶ As the dimensionality of the parameter space increases, it becomes difficult to find an appropriate proposal distributions (e.g., with appropriate step size) for the Metropolis (or MH) algorithm
- ▶ If we are lucky (in some situations we are!), the conditional distribution of one component,  $x_j$ , given all other components,  $x_{-j}$  is tractable and has a close form so that we can sample from it directly
- ▶ If that's the case, we can sample from each component one at a time using their corresponding conditional distributions  $P(x_j|x_{-j})$



- ► This is known as the Gibbs sampler (GS) or "heat bath" (Geman and Geman, 1984)
- Note that in Bayesian analysis, we are mainly interested in sampling from  $p(\theta|y)$
- ► Therefore, we use the Gibbs sampler when  $P(\theta_j|y,\theta_{-j})$  has a closed form, e.g., there is a conditional conjugacy
- ▶ One example is the univariate normal model. As we will see later, given  $\sigma$ , the posterior  $P(\mu|y,\sigma^2)$  has a closed form, and given  $\mu$ , the posterior distribution of  $P(\sigma^2|\mu,y)$  also has a closed form

- ► The Gibbs sampler works as follows
- ▶ Initialize starting value for  $x_1, x_2, \ldots, x_d$
- ▶ At each iteration, pick an ordering of the *d* variables (can be sequential or random)
  - 1. Sample  $x \sim P(x_i|x_1, \dots, x_{i-1}, x_{i+1}, \dots, x_d)$ , *i.e.*, the conditional distribution of  $x_i$  given the current values of all other variables
  - 2. Update  $x_i \leftarrow x$
- ▶ When we update  $x_i$ , we immediately use it new value for sampling other variables  $x_j$



- ▶ Note that in GS, we are not proposing anymore, we are directly sampling, which can be viewed as a proposal that will always be accepted
- ► This way, the Gibbs sampler can be viewed as a special case of MH, whose proposal is

$$Q(x_i', x_{-i}|x_i, x_{-i}) = P(x_i'|x_{-i})$$

▶ Applying MH with this proposal, we obtain

$$a(x_i', x_{-i}|x_i, x_{-i}) = \min\left(1, \frac{P(x_i', x_{-i})Q(x_i, x_{-i}|x_i', x_{-i})}{P(x_i, x_{-i})Q(x_i', x_{-i}|x_i, x_{-i})}\right)$$

$$= \min\left(1, \frac{P(x_i', x_{-i})P(x_i|x_{-i})}{P(x_i, x_{-i})P(x_i'|x_{-i})}\right) = \min\left(1, \frac{P(x_i', x_{-i})P(x_i, x_{-i})}{P(x_i, x_{-i})P(x_i', x_{-i})}\right)$$

$$= 1$$

• We can now use the Gibbs sampler to simulate samples from the posterior distribution of the parameters of a univariate normal  $y \sim \mathcal{N}(\mu, \sigma^2)$  model, with prior

$$\mu \sim \mathcal{N}(\mu_0, \tau_0^2), \quad \sigma^2 \sim \text{Inv-}\chi^2(\nu_0, \sigma_0^2)$$

• Given  $(\sigma^{(i)})^2$  at the  $i^{\text{th}}$  iteration, we sample  $\mu^{(i+1)}$  from

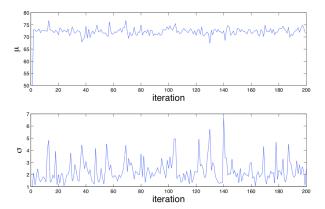
$$\mu^{(i+1)} \sim \mathcal{N}\left(\frac{\frac{\mu_0}{\tau_0^2} + \frac{n\bar{y}}{(\sigma^{(i)})^2}}{\frac{1}{\tau_0^2} + \frac{n}{(\sigma^{(i)})^2}}, \frac{1}{\frac{1}{\tau_0^2} + \frac{n}{(\sigma^{(i)})^2}}\right)$$

• Given  $\mu^{(i+1)}$ , we sample a new  $\sigma^2$  from

$$(\sigma^{(i+1)})^2 \sim \text{Inv-}\chi^2(\nu_0 + n, \frac{\nu_0 \sigma_0^2 + \nu n}{\nu_0 + n}), \quad \nu = \frac{1}{n} \sum_{j=1}^n (y_j - \mu^{(i+1)})^2$$



▶ The following graphs show the trace plots of the posterior samples (for both  $\mu$  and  $\sigma$ )



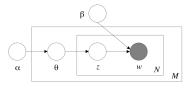


## Gibbs sampling algorithms have been widely used in **probabilistic graphical models**

- ► Conditional distributions are fairly easy to derive for many graphical models (e.g., mixture models, Latent Dirichlet allocation)
- ► Have reasonable computation and memory requirements, only needs to sample one random variable at a time
- ► Can be Rao-Blackwellized (integrate out some random variable) to decrease the sampling variance. This is called *collapsed Gibbs sampling*



► Generative model of documents (Blei, Jordan and Ng, 2003). Also broadly applicable to collaborative filtering, image retrieval, bioinformatics, etc.

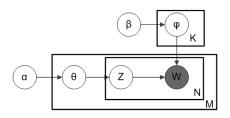


- choose a mixture of topics the document:  $\theta \sim \text{Dir}(\alpha)$
- ▶ choose a topic for each of the document:

$$z_n \sim \text{Multinomial}(\theta)$$

• choose word given the topic:  $w_n|z_n, \beta \sim p(w_n|z_n, \beta)$ 





- Use the probability model for LDA, with an additional Dirichlet prior on  $\phi$ .
- ► The complete probability model

$$w_i|z_i, \phi^{(z_i)} \sim \operatorname{Discrete}(\phi^{(z_i)})$$
  
 $\phi \sim \operatorname{Dirichlet}(\beta)$   
 $z_i|\theta^{(d_i)} \sim \operatorname{Discrete}(\theta^{(d_i)})$   
 $\theta \sim \operatorname{Dirichlet}(\alpha)$ 



► The joint probability is

$$p(w, z, \phi, \theta | \alpha, \beta) = \prod_{i} p(w_i | z_i, \phi^{(z_i)}) p(\phi | \beta) \cdot \prod_{i} p(z_i | \theta^{(d_i)}) p(\theta | \alpha)$$

▶ Due to conjugate priors, we can easily integrate out  $\phi$  and  $\theta$  (T. Griffiths & M. Steyvers, 2004)

$$p(w|z) = \left(\frac{\Gamma(V\beta)}{\Gamma(\beta)^V}\right)^K \prod_{j=1}^K \frac{\prod_w \Gamma(n_j^{(w)} + \beta)}{\Gamma(n_j^{(\cdot)} + V\beta)}$$
$$p(z) = \left(\frac{\Gamma(K\alpha)}{\Gamma(\alpha)^K}\right)^M \prod_{d=1}^M \frac{\prod_j \Gamma(n_j^{(d)} + \alpha)}{\Gamma(n_j^{(d)} + K\alpha)}$$

 $n_j^{(w)} \leftarrow \text{ number of times word } w \text{ assigned to topic } j$  $n_j^{(d)} \leftarrow \text{ number of times topic } j \text{ used in document } d$ 



- ▶ Need full conditional distributions for variables
- $\blacktriangleright$  We only sample z, whose conditional distributions is

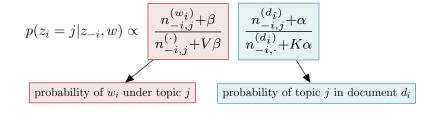
$$p(z_i = j | z_{-i}, w) \propto \left[ \frac{n_{-i,j}^{(w_i)} + \beta}{n_{-i,j}^{(\cdot)} + V\beta} \right] \left[ \frac{n_{-i,j}^{(d_i)} + \alpha}{n_{-i,i}^{(d_i)} + K\alpha} \right]$$

$$\frac{n_{-i,j}^{(d_i)} + \alpha}{n_{-i,\cdot}^{(d_i)} + K\alpha}$$

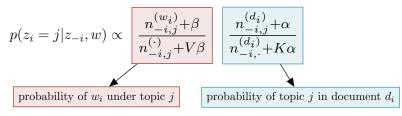
- ▶ Need full conditional distributions for variables
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$$p(z_{i} = j | z_{-i}, w) \propto \begin{bmatrix} \frac{n_{-i,j}^{(w_{i})} + \beta}{n_{-i,j}^{(\cdot)} + V\beta} \\ \\ \frac{n_{-i,j}^{(d_{i})} + \alpha}{n_{-i,j}^{(d_{i})} + K\alpha} \end{bmatrix}$$
probability of  $w_{i}$  under topic  $j$ 

- ▶ Need full conditional distributions for variables
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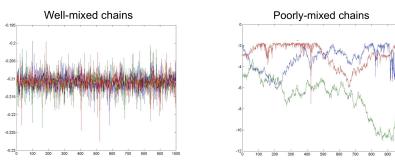
- ► This is nicer than your average Gibbs sampler:
  - ▶ memory: counts can be cashed in two sparse matrices
  - the distributions on  $\phi$  and  $\theta$  are analytic given z and w, and can later be found for each sample



- ► For more complex models, we might only have conditional conjugacy for one part of the parameters
- ► In such situations, we can combine the Gibbs sampler with the Metropolis method
- ► That is, we update the components with conditional conjugacy using Gibbs sampler and for the rest parameters, we use the Metropolis (or MH)



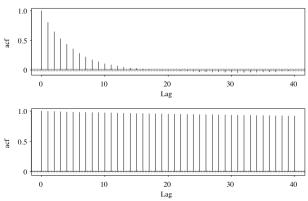
- ► MCMC would converge to the target distribution if run sufficiently long
- ► However, it is often non-trivial to determine whether the chain has converged or not in practice
- ▶ Also, how do we measure the efficiency of MCMC chains?
- ► In what follows, we will discuss some practical advice for coding MCMC algorithms



Monitor convergence by plotting samples from multiple MH runs (chains)

- ► If the chains are well-mixed (left), they are probably converged
- ► If the chains are poorly-mixed (right), we may need to continue burn-in





- ▶ An autocorrelation plot summarizes the correlation in the sequence of a Markov chain at different iteration lags
- A chain that has poor mixing will exhibit slow decay of the autocorrelation as the lag increases

- ▶ Since MCMC samples are correlated, effective sample size are often used to measure the efficiency when MCMC samples are used for estimation instead of independent samples
- ► The effective sample size (ESS) is defined as

$$ESS = \frac{n}{1 + 2\sum_{k=1}^{\infty} \rho(k)}$$

where  $\rho(k)$  is the autocorrelation at lag k

► ESS are commonly used to compare the efficiency of competing MCMC samplers for a given problem. Larger ESS usually means faster convergence



- ▶ One of the hardest problem to diagnose is whether or not the chain has become stuck in one or more modes of the target distribution
- ▶ In this case, all convergence diagnostics may indicate that the chain has converged, though it does not
- ► A partial solution: run multiple chains and compare the within- and between-chain behavior



- ► Auxiliary variable strategies can be used to improving mixing of Markov chains
- ▶ When standard MCMC methods mix poorly, one potential remedy is to augment the state space of the variable of interest
- ► This approach can lead to chains that mix faster and require less tuning than the standard MCMC methods
- ▶ Main idea: construct a Markov chain over (X, U) (U is the auxiliary variable) with stationary distribution marginalizes to the target distribution of X
- ► As we will see later, this includes a large family of modern MCMC methods



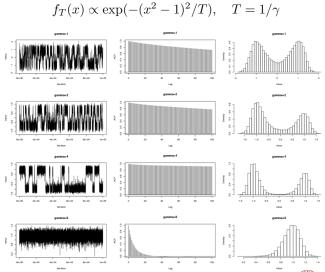
- ▶ Suppose that we have a challenging target distribution  $f(x) \propto \exp(-U(x))$
- ▶ We can introduce temperatures to construct a sequence of distributions that are easier to sample from

$$f_k(x) \propto \exp\left(-U(x)/T_k\right), \quad k = 0, \dots, K$$

where  $1 = T_0 < T_1 < \ldots < T_K$ 

- ▶ When simulating Markov chains with different temperature *T*, the chain with high temperature (hot chain) is likely to mix better than the chain with cold temperature (cold chain)
- ► Therefore, we can run parallel chains and swap states between the chains to improve mixing







We run parallel Markov chains for distributions with different temperatures. In each iteration

- ▶ Follow regular Metropolis steps in each chain to get new states  $x_0^{(t)}, \ldots, x_K^{(t)}$
- ▶ Select two temperatures, say (i, j), i < j, and swap the states

$$x_0^{(t)}, \dots, x_i^{(t)}, \dots, x_j^{(t)}, \dots x_K^{(t)} \to x_0^{(t)}, \dots, x_j^{(t)}, \dots, x_i^{(t)}, \dots x_K^{(t)}$$

► Accept the swapped new states with the following probability

$$\min\left(1, f_i(x_j^{(t)})f_j(x_i^{(t)})/f_i(x_i^{(t)})f_j(x_j^{(t)})\right)$$

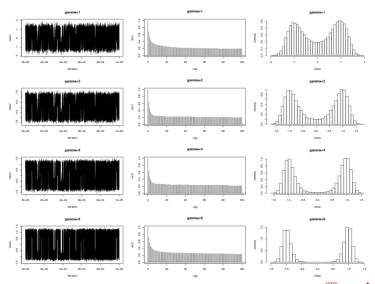


▶ Both the within-chain Metropolis updates and the between-chain swap preserves

$$p(x_0,\ldots,x_K) \propto f_0(x_0)f_1(x_1)\ldots f_K(x_K)$$

- ▶ Therefore, the joint distribution of  $(x_0^{(t)}, \ldots, x_K^{(t)})$  will converge to p(x), and the marginal distribution of  $x_0$  (cold chain) is the target distribution
- ► There are many ways to swap chains. For example, we can pick a pair of temperatures uniformly at random or only swap chains with successive temperatures
- ► The design of temperature levels could be crucial for the performance







- ► Slice sampling was introduced by Neal (2003) to accelerate mixing of Metropolis (or MH)
- ▶ It is essentially a Gibbs sampler in the augmented space (X, U) with density

$$f(x, u) = f(x)f(u|x)$$

where U is the auxiliary variable and f(u|x) is designed to be a uniform distribution  $\mathcal{U}(0, f(x))$ 

- ► For this purpose, slice sampling alternates between two steps:
  - ▶ Given the current state of the Markov chain, x, we uniformly sample a new point u from the interval (0, f(x))

$$U|x \sim \mathcal{U}(0, f(x))$$

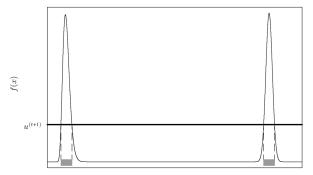
▶ Given the current value of u, we uniformly sample from the region  $S = \{x : f(x) > u\}$ , which is referred to as the *slice* defined by u

$$X|u \sim \mathcal{U}(S)$$

As mentioned by Neal (2003), in practice it is safer to compute  $g(x) = \log(f(x))$ , and use the auxiliary variable  $z = \log(u) = g(x) - e$ , where e has exponential distribution with mean one, and define the slice as  $S = \{x : z < g(x)\}$ 



- ► One advantage of slice sampling is for sampling from multimodal distributions
- ▶ Unlike standard Metropolis (or MH) that struggles between distant modes, sampling from the slice allows us to easily jump between different modes

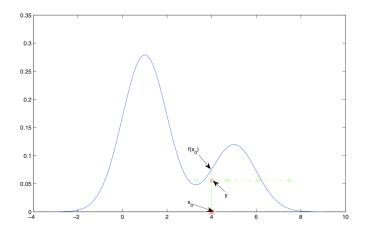




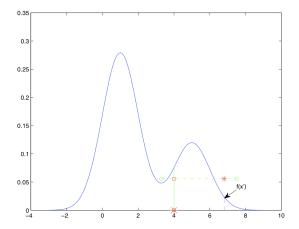
- ightharpoonup Sampling an independent point uniformly from S might be difficult. In practice, we can substitute this step by any update that leaves the uniform distribution over S invariant
- ▶ There are several methods to perform this task
- ▶ Here, we introduce a simple but effective procedure that consists of two phases:
  - ► Stepping-out. A procedure for finding an interval around the current point
  - ▶ *Shrinkage*. A procedure for sampling from the interval obtained
- ► For a detail description of these methods, see Neal (2003)



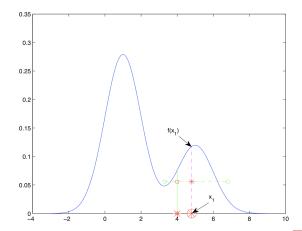
▶ Sampling  $u \sim \mathcal{U}(0, f(x_0))$  and stepping out (of size w) until we reach points outside the slice



▶ Shrinkage of interval to a point, x', which is sampled (uniformly) from the interval but it has f(x') < y



▶ Continue shrinkage until we reach a point  $x_1$  such that  $y < f(x_1)$ . We accept  $x_1$  as our new sample





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