Stat 535 C - Statistical Computing & Monte Carlo Methods

Arnaud Doucet

Email: arnaud@cs.ubc.ca

- Introduction to Markov chain Monte Carlo
- The Gibbs Sampler
- Examples

- Rejection Sampling and Importance Sampling are two general methods but limited to problems of moderate dimensions.
- "Problem": We try to sample all the components of a potentially high-dimensional parameter simultaneously.
- There are two ways to implement incremental strategies.
 - Iteratively: Markov chain Monte Carlo.
 - Sequentially: Sequential Monte Carlo.

• Multiple failures in a nuclear plant:

Pump	1	2	3	4	5	6	7	8	9	10
Failures	5	1	5	14	3	19	1	1	4	22
Times	94.32	15.72	62.88	125.76	5.24	31.44	1.05	1.05	2.10	10.48

- Model: Failures of the *i*-th pump follow a Poisson process with parameter λ_i ($1 \le i \le 10$). For an observed time t_i , the number of failures p_i is thus a Poisson $\mathcal{P}(\lambda_i t_i)$ random variable.
- The unknowns consist of $\theta := (\lambda_1, \ldots, \lambda_{10}, \beta)$.

• Hierarchical model

$$\lambda_i \stackrel{iid}{\sim} \mathcal{G}a(\alpha, \beta) \text{ and } \beta \sim \mathcal{G}a(\gamma, \delta)$$

with $\alpha = 1.8$ and $\gamma = 0.01$ and $\delta = 1$.

• The posterior distribution is proportional to

$$\prod_{i=1}^{10} \{ (\lambda_i t_i)^{p_i} \exp(-\lambda_i t_i) \lambda_i^{\alpha-1} \exp(-\beta\lambda_i) \} \beta^{10\alpha} \beta^{\gamma-1} \exp(-\delta\beta)$$

$$\propto \prod_{i=1}^{10} \{ \lambda_i^{p_i+\alpha-1} \exp(-(t_i+\beta)\lambda_i) \} \beta^{10\alpha+\gamma-1} \exp(-\delta\beta).$$

• This multidimensional distribution is rather complex. It is not obvious how the inverse cdf method, the rejection method or importance sampling could be used in this context. • The conditionals have a familiar form

$$\lambda_i | (\beta, t_i, p_i) \sim \mathcal{G}a(p_i + \alpha, t_i + \beta) \text{ for } 1 \le i \le 10,$$

$$\beta|(\lambda_1,\ldots,\lambda_{10})\sim \mathcal{G}a(\gamma+10\alpha,\delta+\sum_{i=1}^{10}\lambda_i).$$

• Instead of directly sampling the vector $\theta = (\lambda_1, \ldots, \lambda_{10}, \beta)$ at once, one could suggest sampling it iteratively, starting for example with the λ_i 's for a given guess of β , followed by an update of β given the new samples $\lambda_1, \ldots, \lambda_{10}$. • Given a sample, at iteration $t, \theta^t := (\lambda_1^t, \dots, \lambda_{10}^t, \beta^t)$ one could proceed as follows at iteration t + 1,

1.
$$\lambda_i^{t+1}|(\beta^t, t_i, p_i) \sim \mathcal{G}a(p_i + \alpha, t_i + \beta^t)$$
 for $1 \le i \le 10$,

2.
$$\beta^{t+1}|(\lambda_1^{t+1},\ldots,\lambda_{10}^{t+1}) \sim \mathcal{G}a(\gamma+10\alpha,\delta+\sum_{i=1}^{10}\lambda_i^{t+1}).$$

• Instead of directly sampling in a space with 11 dimensions, one samples in spaces of dimension 1

• Note that the deterministic version of such an algorithm would not generally converge towards the global maximum of the joint distribution.

- The structure of the algorithm calls for many questions:
 - Are we sampling from the desired joint distribution?
 - If yes, how many times should the iteration above be repeated?

• The validity of the approach described here stems from the fact that the sequence $\{\theta^t\}$ defined above is a Markov chain and some Markov chains have very nice properties.

• Markov chain: A sequence of random variables $\{X_n; n \in \mathbb{N}\}$ defined on $(\mathbb{X}, \mathcal{B}(\mathbb{X}))$ which satisfies the property, for any $A \in \mathcal{B}(\mathbb{X})$

$$\mathbb{P}(X_{n} \in A | X_{0}, ..., X_{n-1}) = \mathbb{P}(X_{n} \in A | X_{n-1}).$$

and we will write

$$P(x,A) = \mathbb{P}(X_n \in A | X_{n-1}).$$

• Markov chain Monte Carlo: Given a target π , design a transition kernel P such that asymptotically as $n \to \infty$

$$\frac{1}{N}\sum_{n=1}^{N}\varphi\left(X_{n}\right)\to\int\varphi\left(x\right)\pi\left(x\right)dx \text{ and/or } X_{n}\sim\pi.$$

• It should be easy to simulate the Markov chain even if π is complex.

• Consider the autoregression for $|\alpha| < 1$

$$X_n = \alpha X_{n-1} + V_n$$
, where $V_n \sim \mathcal{N}(0, \sigma^2)$.

• The limiting distribution is

$$\pi(x) = \mathcal{N}\left(x; 0, \frac{\sigma^2}{1 - \alpha^2}\right).$$

• To sample from π , we could just sample the Markov chain and asymptotically we would have $X_n \sim \pi$.

• Obviously, in this case this is useless because we can sample from π directly.

- Graphically, consider 1000 independent Markov chains run in parallel.
- We assume that the initial distribution of these Markov chains is $\mathcal{U}_{[0,20]}$. So initially, the Markov chains samples are not distributed according to π

From top left to bottom right: histograms of 1000 independent Markov chains with a normal distribution as target distribution.



• The target normal distribution seems to "attract" the distribution of the samples and even to be a fixed point of the algorithm.

• This is is what we wanted to achieve, *i.e.* it seems that we have produced 1000 independent samples from the normal distribution.

• In fact one can show that in many (all?) situations of interest it is not necessary to run N Markov chains in parallel in order to obtain 1000 samples, but that one can consider a unique Markov chain, and build the histogram from this single Markov chain by forming histograms from one trajectory.

2.8– Example: Mixture of Normals

1000 iterations



2.8– Example: Mixture of Normals

10000 iterations



• The estimate of the target distribution, through the series of histograms, improves with the number of iterations.

• Assume that we have stored $\{X_n, 1 \le n \le N\}$ for N large and wish to estimate $\int_X \varphi(x) \pi(x) dx$.

• In the light of the numerical experiments, one can suggest the estimator N

$$\frac{1}{N}\sum_{n=1}^{N}\varphi(X_n).$$

which is exactly the estimator that we would use if $\{X_n, 1 \le n \le N\}$ were independent.

• In fact, it can be proved, under relatively mild conditions, that such an estimator is consistent *despite the fact that the samples are NOT independent!* Under additional conditions, a CLT also holds with a rate of CV in $1/\sqrt{N}$.

To summarize, we are interested in Markov chains with transition kernel P which have the following three important properties observed above:

• The desired distribution π is a "fixed point" of the algorithm or, in more appropriate terms, an *invariant distribution* of the Markov chain, *i.e.* $\int_{\mathsf{X}} \pi(x) P(x, y) = \pi(y)$.

• The successive distributions of the Markov chains are "attracted" by π , or converge towards π .

• The estimator $\frac{1}{N} \sum_{n=1}^{N} \varphi(X_n)$ converges towards $E_{\pi}(\varphi(X))$ and asymptotically $X_n \sim \pi$

- Given $\pi(x)$, there is an infinite number of kernels P(x, y) which admits $\pi(x)$ as their invariant distribution.
- The "art" of MCMC consists of coming up with good ones.
- Convergence is ensured under very weak assumptions; namely irreducibility and aperiodicity.
- It is usually very easy to establish that an MCMC sampler converges towards π but very difficult to obtain rates of convergence.

• Consider the target distribution $\pi(\theta)$ such that $\theta = (\theta^1, \theta^2)$. Then the 2 component Gibbs sampler proceeds as follows.

Initialization:

• Select deterministically or randomly $\theta_0 = (\theta_0^1, \theta_0^2)$.

Iteration $i; i \ge 1$:

- Sample $\theta_i^1 \sim \pi \left(\left. \theta^1 \right| \theta_{i-1}^2 \right)$.
- Sample $\theta_i^2 \sim \pi \left(\left. \theta^2 \right| \theta_i^1 \right)$.
- Sampling from these conditional is often feasible even when sampling from the joint is impossible (e.g. nuclear pump data).

• Clearly $\left\{ \begin{pmatrix} \theta_i^1, \theta_i^2 \end{pmatrix} \right\}$ is a Markov chain and its transition kernel is $P\left(\left(\theta^1, \theta^2 \right), \left(\widetilde{\theta}^1, \widetilde{\theta}^2 \right) \right) = \pi \left(\left. \widetilde{\theta}^1 \right| \left. \theta^2 \right) \pi \left(\left. \widetilde{\theta}^2 \right| \widetilde{\theta}^1 \right).$

• Then
$$\int \int \pi (\theta^1, \theta^2) P\left(\left(\theta^1, \theta^2\right), \left(\widetilde{\theta}^1, \widetilde{\theta}^2\right)\right) d\theta^1 d\theta^2$$
 satisfies
 $\int \int \pi (\theta^1, \theta^2) \pi \left(\widetilde{\theta}^1 \middle| \theta^2\right) \pi \left(\widetilde{\theta}^2 \middle| \widetilde{\theta}^1\right) d\theta^1 d\theta^2$
 $= \int \pi (\theta^2) \pi \left(\widetilde{\theta}^1 \middle| \theta^2\right) \pi \left(\widetilde{\theta}^2 \middle| \widetilde{\theta}^1\right) d\theta^2$
 $= \int \pi \left(\widetilde{\theta}^1, \theta^2\right) \pi \left(\widetilde{\theta}^2 \middle| \widetilde{\theta}^1\right) d\theta^2$
 $= \pi \left(\widetilde{\theta}^1\right) \pi \left(\widetilde{\theta}^2 \middle| \widetilde{\theta}^1\right) = \pi \left(\widetilde{\theta}^1, \widetilde{\theta}^2\right)$

• This does not ensure that the Gibbs sampler does converge towards the invariant distribution!

• Additionally it is required to ensure irreducibility: loosely speaking the Markov chain can move to any set A such that $\pi(A) > 0$ for (almost) any starting point.

• This ensures that

$$\frac{1}{N}\sum_{n=1}^{N}\varphi\left(\theta_{n}^{1},\theta_{n}^{2}\right)\rightarrow\int\varphi\left(\theta^{1},\theta^{2}\right)\pi\left(\theta^{1},\theta^{2}\right)d\theta^{1}d\theta^{2}$$

but NOT that asymptotically $(\theta_n^1, \theta_n^2) \sim \pi$.

A distribution that can lead to a reducible Gibbs sampler.



• Consider a simple example where $\mathbb{X} = \{1, 2\}$ and P(1, 2) = P(2, 1) = 1. Clearly the invariant distribution is given by $\pi(1) = \pi(2) = \frac{1}{2}$.

• However, we know that if the chain starts in $X_0 = 1$, then $X_{2n} = 1$ and $X_{2n+1} = 0$ for any n.

• We have

$$\frac{1}{N}\sum_{n=1}^{N}\varphi\left(X_{n}\right)\rightarrow\int\varphi\left(x\right)\pi\left(x\right)dx$$

but clearly X_n is NOT distributed according to π .

• You need to make sure that you do NOT explore the space in a periodic way to ensure that $X_n \sim \pi$ asymptotically. Even when irreducibility and aperiodicity are ensured, the Gibbs sampler can still converge very slowly.



- If $\theta = (\theta_1, ..., \theta_p)$ where p > 2, the Gibbs sampling strategy still applies.
- Initialization:
 - Select deterministically or randomly $\theta_0 = (\theta_0^1, ..., \theta_0^p)$.
- Iteration $i; i \ge 1$:

For k = 1: p

• Sample $\theta_i^k \sim \pi \left(\left. \theta^k \right| \theta_i^{-k} \right)$.

where
$$heta_i^{-k} = \left(heta_i^1,..., heta_i^{k-1}, heta_{i-1}^{k+1},..., heta_{i-1}^p
ight).$$

- Initialization:
 - Select deterministically or randomly $\theta_0 = (\theta_0^1, ..., \theta_0^p)$.
- Iteration $i; i \ge 1$:
 - Sample $K \sim U_{\{1,\dots,p\}}$.
 - Set $\theta_i^{-K} = \theta_{i-1}^{-K}$.
 - Sample $\theta_i^K \sim \pi \left(\left. \theta^K \right| \theta_i^{-K} \right)$.

where
$$\theta_i^{-K} = \left(\theta_i^1, ..., \theta_i^{K-1}, \theta_i^{K+1}, ..., \theta_i^p\right)$$
.

- Try to have as few "blocks" as possible.
- Put the most correlated variables in the same block.
- If necessary, reparametrize the model to achieve this.
- Integrate analytically as many variables as possible: pretty algorithms can be much more inefficient than ugly algorithms.
- There is no general result telling strategy A is better than strategy B in all cases: you need experience.

• We select the following model

$$Y = \sum_{i=1}^{p} \beta_{i} X_{i} + \sigma V \text{ where } V \sim \mathcal{N}(0, 1)$$

where we assume
$$\mathcal{IG}\left(\sigma^2; \frac{\nu_0}{2}, \frac{\gamma_0}{2}\right)$$
 and for $\alpha^2 << 1$
 $\beta_i \sim \frac{1}{2} \mathcal{N}\left(0, \alpha^2 \delta^2 \sigma^2\right) + \frac{1}{2} \mathcal{N}\left(0, \delta^2 \sigma^2\right)$

• We introduce a latent variable $\gamma_i \in \{0, 1\}$ such that

$$\Pr\left(\gamma_{i}=0\right) = \Pr\left(\gamma_{i}=1\right) = \frac{1}{2},$$

$$\beta_{i}|\gamma_{i}=0 \sim \mathcal{N}\left(0, \alpha^{2}\delta^{2}\sigma^{2}\right), \quad \beta_{i}|\gamma_{i}=1 \sim \mathcal{N}\left(0, \delta^{2}\sigma^{2}\right).$$

- We have parameters $(\beta_{1:p}, \gamma_{1:p}, \sigma^2)$ and observe *n* observations $D = \{x_i, y_i\}_{i=1}^n$.
- A potential Gibbs sampler consists of sampling iteratively from $p(\beta_{1:p}|D, \gamma_{1:p}, \sigma^2)$ (Gaussian), $p(\sigma^2|D, \gamma_{1:p}, \beta_{1:p})$ (inverse-Gamma) and $p(\gamma_{1:p}|D, \beta_{1:p}, \sigma^2)$.
- In particular $p\left(\gamma_{1:p} \mid D, \beta_{1:p}, \sigma^{2}\right) = \prod_{i=1}^{p} p\left(\gamma_{i} \mid \beta_{i}, \sigma^{2}\right)$

and

$$p\left(\gamma_{i}=1|\beta_{i},\sigma^{2}\right)=\frac{\frac{1}{\sqrt{2\pi\delta\sigma}}\exp\left(-\frac{\beta_{i}^{2}}{2\delta^{2}\sigma^{2}}\right)}{\frac{1}{\sqrt{2\pi\delta\sigma}}\exp\left(-\frac{\beta_{i}^{2}}{2\delta^{2}\sigma^{2}}\right)+\frac{1}{\sqrt{2\pi\alpha\delta\sigma}}\exp\left(-\frac{\beta_{i}^{2}}{2\alpha^{2}\delta^{2}\sigma^{2}}\right)}.$$

• The Gibbs sampler becomes reducible as α goes to zero.

• This is the result of bad modelling and bad algorithm. You would like to put $\alpha \simeq 0$ and write

$$Y = \sum_{i=1}^{n} \gamma_i \beta_i X_i + \sigma V \text{ where } V \sim \mathcal{N}(0, 1)$$

where $\gamma_i = 1$ if X_i is included or $\gamma_i = 0$ otherwise. However this suggests that β_i is defined even when $\gamma_i = 0$.

• A neater way to write such models is to write

$$Y = \sum_{\{i:\gamma_i=1\}} \beta_i X_i + \sigma V = \beta_{\gamma}^{\mathrm{T}} X_{\gamma} + \sigma V$$

where, for a vector $\gamma = (\gamma_1, ..., \gamma_p), \beta_{\gamma} = \{\beta_i : \gamma_i = 1\}, X_{\gamma} = \{X_i : \gamma_i = 1\}$ and $n_{\gamma} = \sum_{i=1}^p \gamma_i$.

• Prior distributions

$$\pi_{\gamma} \left(\beta_{\gamma}, \sigma^{2} \right) = \mathcal{N} \left(\beta_{\gamma}; 0, \delta^{2} \sigma^{2} I_{n_{\gamma}} \right) \mathcal{IG} \left(\sigma^{2}; \frac{\nu_{0}}{2}, \frac{\gamma_{0}}{2} \right)$$
$$\pi \left(\gamma \right) = \prod_{i=1}^{p} \pi \left(\gamma_{i} \right) = 2^{-p}.$$

and

- We are interested in sampling from the trans-dimensional distribution $\pi(\gamma, \beta_{\gamma}, \sigma^2 | D)$
- However, we know that

$$\pi\left(\gamma,\beta_{\gamma},\sigma^{2} \mid D\right) = \pi\left(\gamma \mid D\right)\pi\left(\beta_{\gamma},\sigma^{2} \mid D,\gamma\right)$$

where

$$\pi \left(\left. \gamma \right| D \right) \propto \pi \left(\left. D \right| \gamma \right) \pi \left(\gamma \right)$$

and

$$\pi \left(D \right| \gamma \right) = \int \pi \left(D, \beta_{\gamma}, \sigma^{2} \right| \gamma \right) d\beta_{\gamma} d\sigma^{2}$$

$$\propto \Gamma \left(\frac{\nu_{0} + n}{2} + 1 \right) \delta^{-n_{\gamma}} |\Sigma_{\gamma}|^{1/2} \left(\frac{\gamma_{0} + \sum_{i=1}^{n} y_{i}^{2} - \mu_{\gamma}^{\mathrm{T}} \Sigma_{\gamma}^{-1} \mu_{\gamma}}{2} \right)^{-\left(\frac{\nu_{0} + n}{2} + 1\right)}$$

- Summary

- $\pi(\gamma | D)$ is a discrete probability distribution with 2^p potential values.
- We can use the Gibbs sampler to sample from it.
- Initialization:
 - Select deterministically or randomly $\gamma_0 = \left(\gamma_0^1, ..., \gamma_0^p\right)$.
- Iteration $i; i \ge 1$:

For k = 1: p

• Sample
$$\gamma_i^k \sim \pi \left(\gamma^k \middle| D, \gamma_i^{-k} \right)$$
.
where $\gamma_i^{-k} = \left(\gamma_i^1, \dots, \gamma_i^{k-1}, \gamma_{i-1}^{k+1}, \dots, \gamma_{i-1}^p \right)$.

• Optional step: Sample $(\beta_{\gamma,i}, \sigma_i^2) \sim \pi (\beta_{\gamma}, \sigma^2 | D, \gamma_i).$

- This very simple sampler is much more efficient than the previous one.
- However, it can also mix very slowly because the components are updated one at a time.

• Updating correlated components together would increase significantly the convergence speed of the algorithm at the cost of an increased complexity.

- The Gibbs sampler is a generic tool to sample approximately from high-dimensional distributions.
- Each time you face a problem, you need to think hard about it to design an efficient algorithm.
- Except the choice of the partitions of parameters, the Gibbs sampler is parameter free; this does not mean it is efficient.