

CPSC 535

Gibbs Sampling

AD

February 2007

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- **Problem:** We try to sample all the components of a potentially high-dimensional parameter simultaneously/sequentially and we can never correct for components already sampled.
- A powerful class of methods is available to deal with such methods: Markov chain Monte Carlo.

- Multiple failures in a nuclear plant

Pump i	1	2	3	4	5
# Failures p_i	5	1	5	14	3
Times t_i	94.32	15.72	62.88	125.76	5.24
Pump i	6	7	8	9	10
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- Model: Failures of the i -th pump follow a Poisson process with parameter λ_i ($1 \leq i \leq 10$). For an observed time t_i , the number of failures p_i is thus a Poisson $\mathcal{P}(\lambda_i t_i)$ random variable.

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- The unknown parameters consist of $\theta = (\lambda_1, \dots, \lambda_{10}, \beta)$.

- Hierarchical model

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

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

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- The posterior distribution is proportional to

$$\begin{aligned} & p(\lambda_{1:10}, \beta | p_{1:10}, t_{1:10}) \\ \propto & \prod_{i=1}^{10} \{ (\lambda_i t_i)^{p_i} \exp(-\lambda_i t_i) \lambda_i^{\alpha-1} \exp(-\beta \lambda_i) \} \beta^{10\alpha} \beta^{\gamma-1} \exp(-\delta\beta) \\ \propto & \prod_{i=1}^{10} \{ \lambda_i^{p_i+\alpha-1} \exp(-(t_i + \beta)\lambda_i) \} \beta^{10\alpha+\gamma-1} \exp(-\delta\beta). \end{aligned}$$

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- This multidimensional distribution is rather complex. It is not obvious how the rejection method or importance sampling could be used in this context.

- The conditionals have a familiar form

$$p(\lambda_{1:10} | p_{1:10}, t_{1:10}, \beta) = \prod_{i=1}^{10} p(\lambda_i | p_i, t_i, \beta)$$

where

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

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- Instead of directly sampling the vector $\theta = (\lambda_1, \dots, \lambda_{10}, \beta)$ 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 $\lambda_1, \dots, \lambda_{10}$.

My first Gibbs sampler

- 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 $1 \leq i \leq 10$,
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- 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 where sampling is replaced by maximization would not generally converge towards the global maximum of the joint distribution.

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 - 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.

Elements of Markov chains

- **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 $A \in \mathcal{B}(\mathbb{X})$

$$\mathbb{P}(X_n \in A | X_0, \dots, X_{n-1}) = \mathbb{P}(X_n \in A | X_{n-1}).$$

and we will write

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

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

$$\frac{1}{N} \sum_{n=1}^N \varphi(X_n) \rightarrow \int_{\mathbb{X}} \varphi(x) \pi(x) dx \text{ and/or } X_n \sim \pi.$$

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

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

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

then

$$P(x, dy) = P(x, y) dy = \frac{1}{\sqrt{2\pi}\sigma} \exp\left(-\frac{(y - \alpha x)^2}{2\sigma^2}\right) dx.$$

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

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- 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.]

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- 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

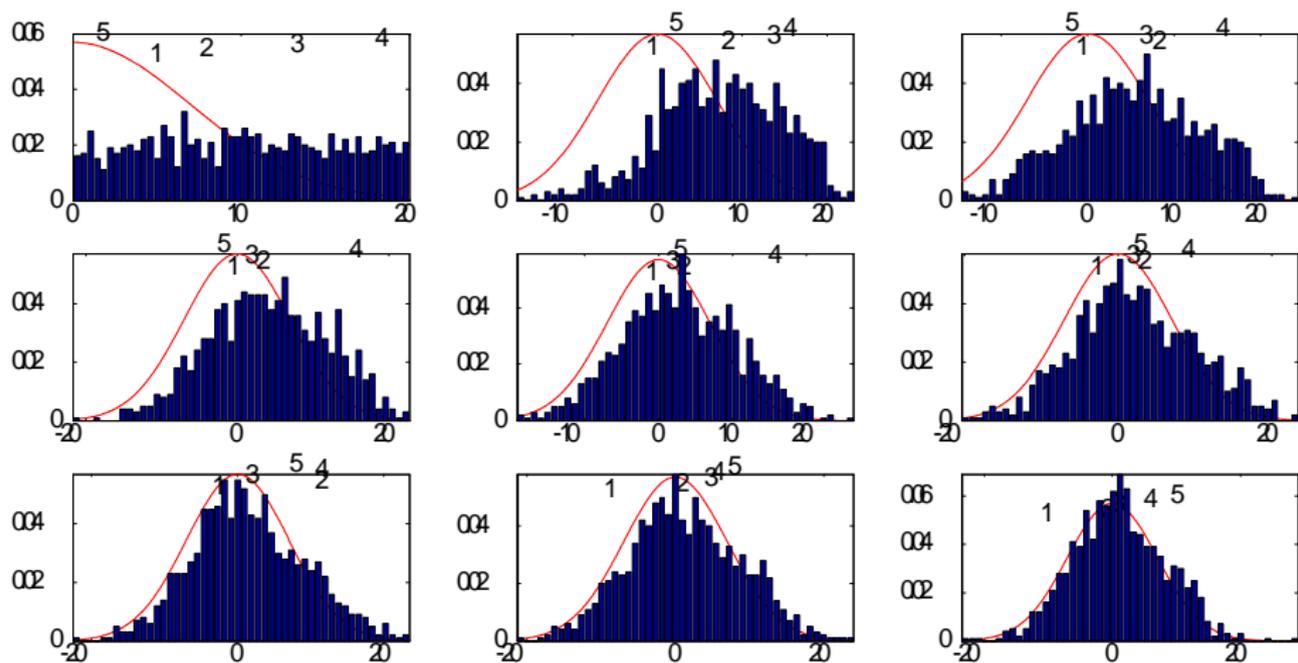


Figure: From top left to bottom right: histograms of 1000 independent Markov chains with a normal distribution as target distribution as n increases.

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- This 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.

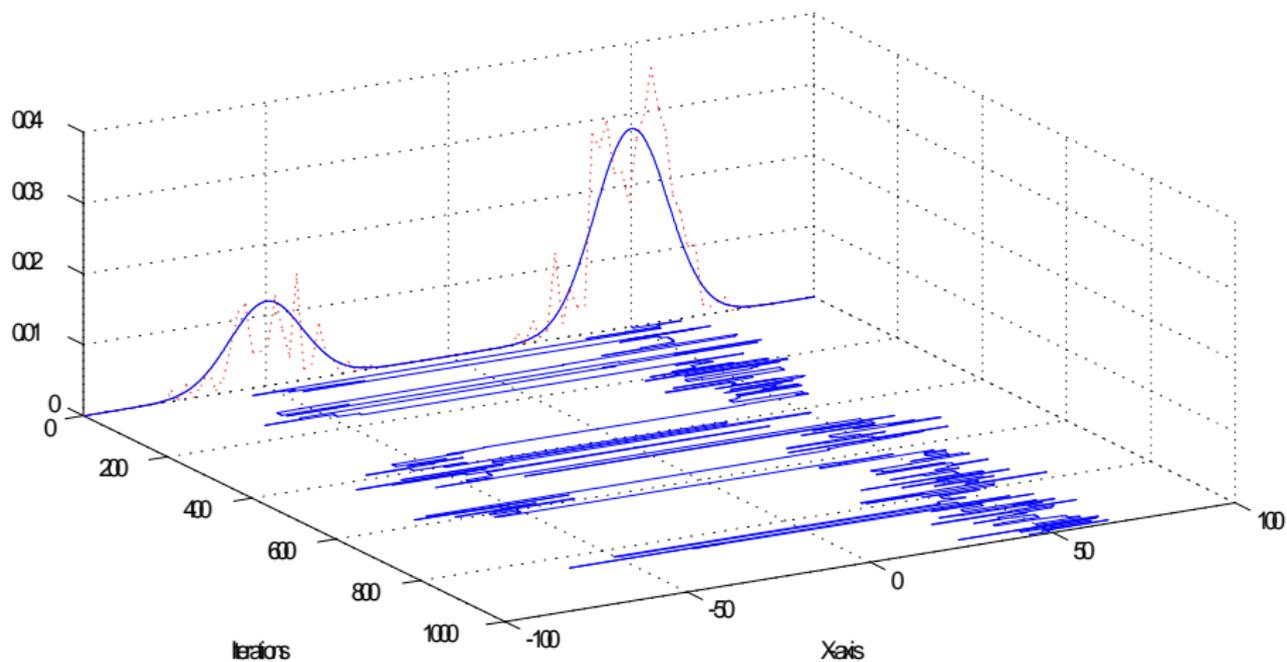


Figure: Bimodal target distributions and simulated Markov chain

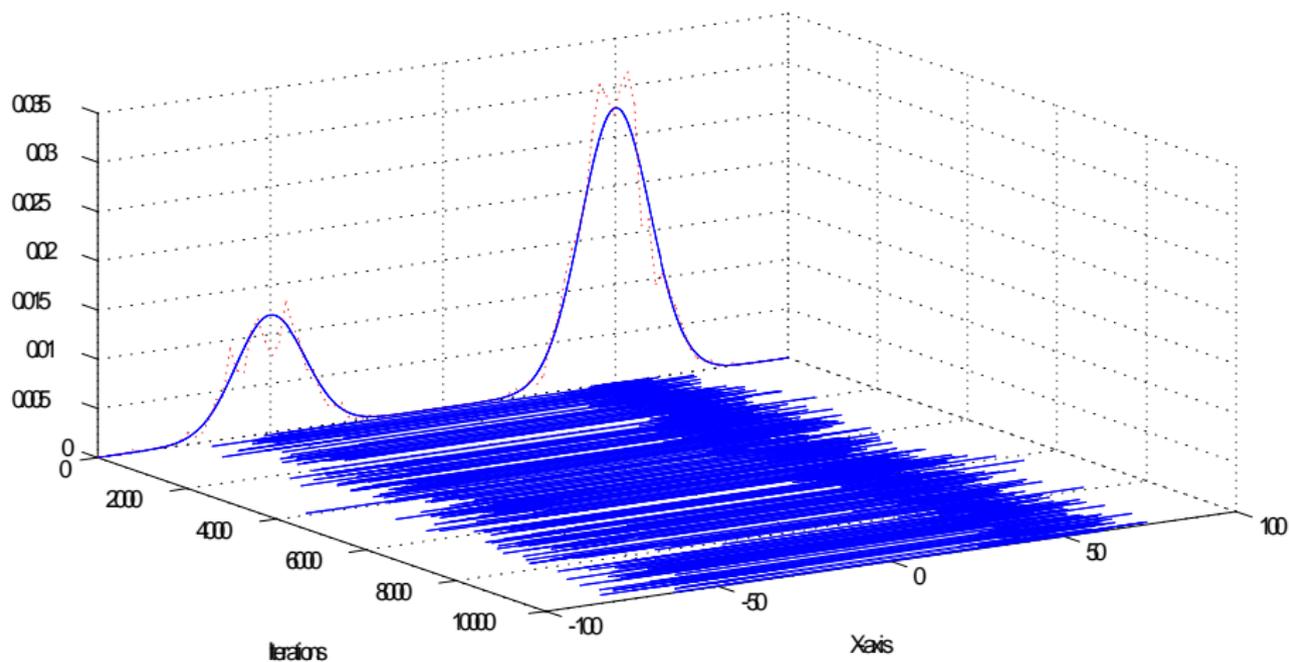


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- In fact, it can be proved, under relatively mild conditions, that such an estimator is consistent *despite the fact that the samples are NOT independent!* Under additional conditions, a CLT also holds with a rate of CV in $1/\sqrt{N}$.

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. $\int_{\mathbb{X}} \pi(x)P(x, y)dx = \pi(y)$.

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Markov chains for Monte Carlo

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- 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. $\int_{\mathbb{X}} \pi(x)P(x, y)dx = \pi(y)$.
- The successive distributions of the Markov chains converge towards π .
- The estimator $\frac{1}{N} \sum_{n=1}^N \varphi(X_n)$ converges towards $\mathbb{E}_{\pi}(\varphi(X))$ and asymptotically $X_n \sim \pi$

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- 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.

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

- Clearly $\{(\theta_i^1, \theta_i^2)\}$ is a Markov chain and its transition kernel is

$$P\left((\theta^1, \theta^2), (\tilde{\theta}^1, \tilde{\theta}^2)\right) = \pi\left(\tilde{\theta}^1 \mid \theta^2\right) \pi\left(\tilde{\theta}^2 \mid \tilde{\theta}^1\right).$$

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- Then $\int \int \pi(\theta^1, \theta^2) P\left((\theta^1, \theta^2), (\tilde{\theta}^1, \tilde{\theta}^2)\right) d\theta^1 d\theta^2$ satisfies

$$\begin{aligned} & \int \int \pi(\theta^1, \theta^2) \pi\left(\tilde{\theta}^1 \mid \theta^2\right) \pi\left(\tilde{\theta}^2 \mid \tilde{\theta}^1\right) d\theta^1 d\theta^2 \\ &= \int \pi(\theta^2) \pi\left(\tilde{\theta}^1 \mid \theta^2\right) \pi\left(\tilde{\theta}^2 \mid \tilde{\theta}^1\right) d\theta^2 \\ &= \int \pi\left(\tilde{\theta}^1, \theta^2\right) \pi\left(\tilde{\theta}^2 \mid \tilde{\theta}^1\right) d\theta^2 \\ &= \pi\left(\tilde{\theta}^1\right) \pi\left(\tilde{\theta}^2 \mid \tilde{\theta}^1\right) = \pi\left(\tilde{\theta}^1, \tilde{\theta}^2\right) \end{aligned}$$

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- 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

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

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

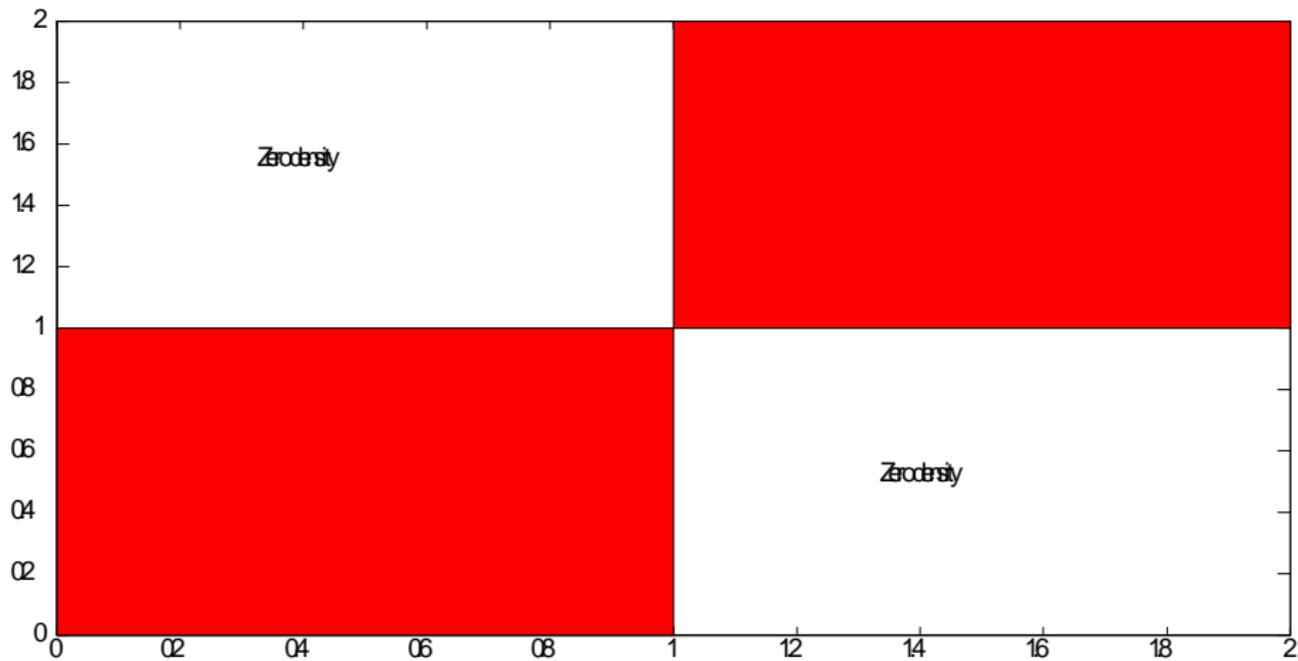


Figure: A distribution that can lead to a reducible Gibbs sampler.

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- You need to make sure that you do NOT explore the space in a periodic way to ensure that $X_n \sim \pi$ asymptotically.

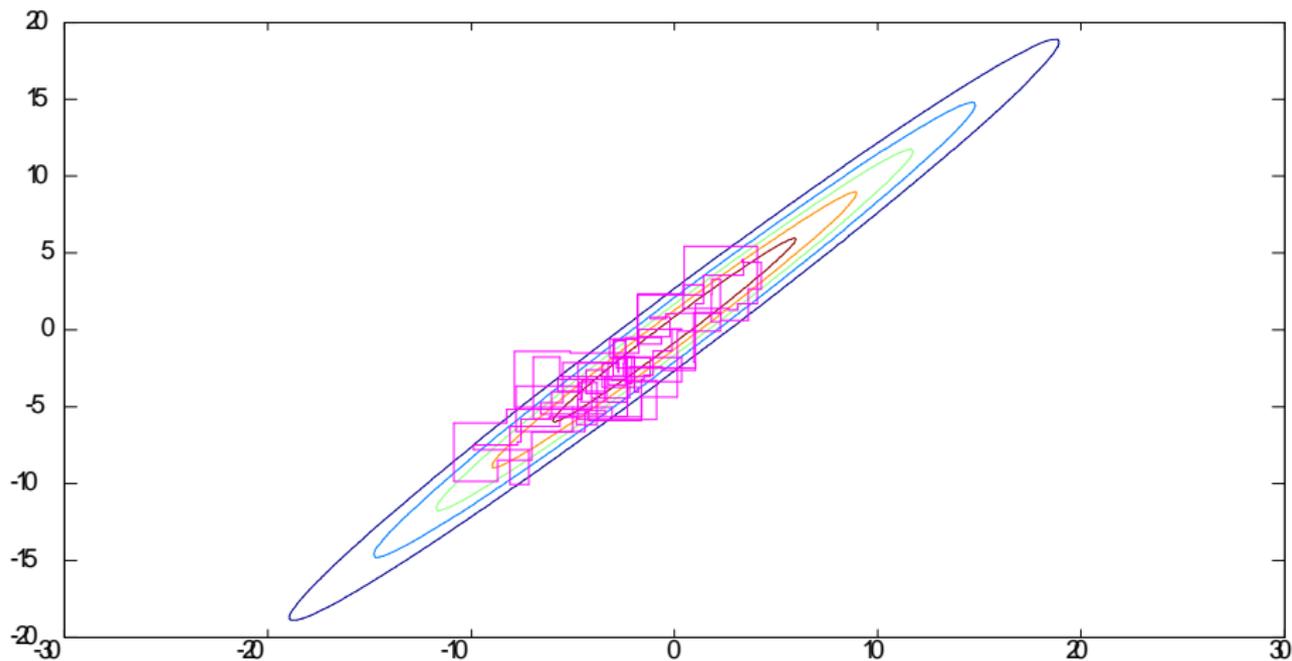


Figure: Even when irreducibility and aperiodicity are ensured, the Gibbs sampler can still converge very slowly.

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- Iteration i ; $i \geq 1$:
 - For $k = 1 : p$
 - Sample $\theta_i^k \sim \pi(\theta^k | \theta_i^{-k})$ where $\theta_i^{-k} = (\theta_i^1, \dots, \theta_i^{k-1}, \theta_{i-1}^{k+1}, \dots, \theta_{i-1}^p)$.

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 - Sample $\theta_i^K \sim \pi(\theta^K | \theta_i^{-K})$ where
 $\theta_i^{-K} = (\theta_i^1, \dots, \theta_i^{K-1}, \theta_i^{K+1}, \dots, \theta_i^p)$.

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- 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.

Application to Simulation of Fractal Images

- Consider a 2D black and white 'target' image. We define an distribution ν which assigns $1/P$ mass on each black point and zero on white points where P is the number of black points.

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- To find $\{w_i, A_i, b_i\}$, we write

$$\begin{aligned} \int \pi(x) P(x, y) f(y) dx dy &= \sum_{i=1}^k w_i \int f(A_i x + b_i) \pi(x) dx \\ &= \int f(x) \pi(x) dx \approx \int f(x) \nu(x) dx \end{aligned}$$

and solve approximately the equations for some functions f (linear or low order polynomials).

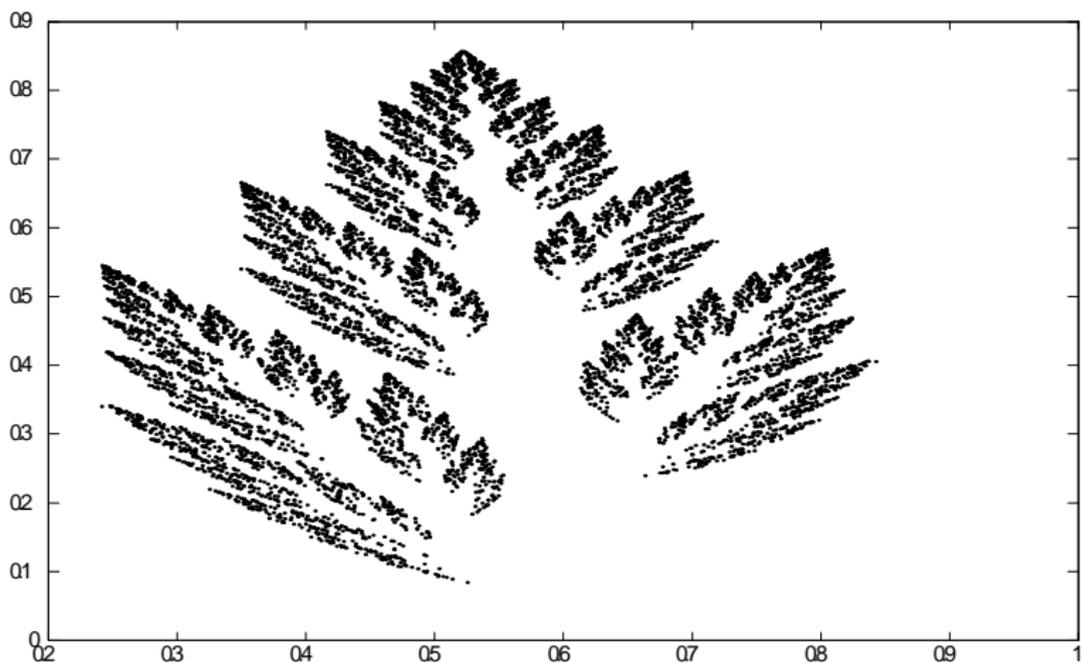


Figure: Fractal image generated using iterated random functions with $k = 2$ and $N = 10000$ samples

Gibbs Sampler for Bayesian Variable Selection

- 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 $\mathcal{IG}(\sigma^2; \frac{\nu_0}{2}, \frac{\gamma_0}{2})$ and for $\alpha^2 \ll 1$

$$\beta_i \sim \frac{1}{2} \mathcal{N}(0, \alpha^2 \delta^2 \sigma^2) + \frac{1}{2} \mathcal{N}(0, \delta^2 \sigma^2)$$

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

$$\Pr(\gamma_i = 0) = \Pr(\gamma_i = 1) = \frac{1}{2},$$
$$\beta_i | \gamma_i = 0 \sim \mathcal{N}(0, \alpha^2 \delta^2 \sigma^2), \quad \beta_i | \gamma_i = 1 \sim \mathcal{N}(0, \delta^2 \sigma^2).$$

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- 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 $p(\gamma_{1:p} | D, \beta_{1:p}, \sigma^2)$.

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- In particular

$$p(\gamma_{1:p} | D, \beta_{1:p}, \sigma^2) = \prod_{i=1}^p p(\gamma_i | \beta_i, \sigma^2)$$

and

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

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- 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

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

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- A neater way to write such models is to write

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

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

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- Prior distributions

$$\pi_\gamma(\beta_\gamma, \sigma^2) = \mathcal{N}(\beta_\gamma; 0, \delta^2 \sigma^2 I_{n_\gamma}) \mathcal{IG}(\sigma^2; \frac{\nu_0}{2}, \frac{\gamma_0}{2})$$

and $\pi(\gamma) = \prod_{i=1}^p \pi(\gamma_i) = 2^{-p}$.

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- However, we know that

$$\pi(\gamma, \beta_\gamma, \sigma^2 | D) = \pi(\gamma | D) \pi(\beta_\gamma, \sigma^2 | D, \gamma)$$

where

$$\pi(\gamma | D) \propto \pi(D | \gamma) \pi(\gamma)$$

and

$$\pi(D | \gamma) = \int \pi(D, \beta_\gamma, \sigma^2 | \gamma) d\beta_\gamma d\sigma^2.$$

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

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

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- Except the choice of the partitions of parameters, the Gibbs sampler is parameter free; this does not mean it is efficient.