Markov chain Monte Carlo methods

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We want to approximate complex integral: let $x_1,...,x_N\sim p(x)$, then for any test function h(x):

$$\mathbb{E}_{X \sim p}[h(X)] = \int h(x)p(x)dx \approx \frac{1}{N} \sum_{n=1}^{N} h(x_n).$$

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Here, x can be parameters or any latent variables of interest.

Importance sampling

Instead of sampling from p (hard to do because Z is unknown), sample $x \sim q$ and adjust for the difference between γ and q:

$$\int h(x)p(x)dx \approx \sum \bar{w}(x)h(x),$$

where
$$w(x) = \gamma(x)/q(x)$$
 and $\bar{w}(x) = w(x)/\sum_n w(x)$.

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If $\boldsymbol{x}=(x_1,...,x_D)$ is high-dimensional, we can sample each component sequentially:

- $x_d^n \sim q_d(x_d|x_{1:d-1})$.
- Interleave resampling step to maintain particle diversity and prune unpromising particles.

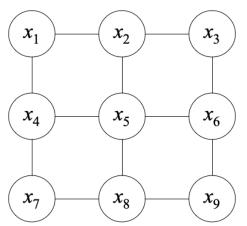
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So why do we need another algorithm/method?

- Large variance associated with choice of proposal distribution.
- Curse of dimensionality may still manifest and approximation can be poor.

Consider a $K \times K$ 2-dimensional lattice G = (V, E).



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- Spatial analysis: Each node represents a spatial coordinate (spatial statistics) and $X_v \in \{ \text{no gold}, \text{gold!} \}$ or $X_v \in \mathbb{R}^+$ some measurement of the amount of gold at location v.
- Image processing: Each node represents a pixel of an image. $X_v \in \{ \text{black}, \text{white} \}$ or gray scale $X_v \in [0,1]$ or RGB color.

Example: Ising model

Let $X=(X_v).$ The "energy" function for the Ising model is defined as:

$$H(x) = \sum_v \phi(x_v) + \sum_{(u,v) \in E} \psi(x_u,x_v),$$

- $(u, v) \in E$ denote neighbors (adjacent nodes),
- ϕ : unary potential,
- ψ : pairwise potential (measuring interaction strength).

Example: $\phi(x_v) = \beta x_v$ and $\psi(x_u, x_v) = \kappa x_u x_v$ for $\beta, \kappa \in \mathbb{R}$.

Example: Ising model

Tthe probability distribution on X is defined as,

$$p(x) = \frac{1}{Z} \exp(-H(x))$$

where

$$Z = \sum_{x_n: v \in V} \exp(-H(x)).$$

 ${\cal Z}$ in statistical physics is referred to as "partition function". Essentially a normalization constant.

SMC for Ising model?

- Not obvious what order to sample the variables.
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- Leads to poor approximation involving those sampled earlier on.
- \bullet For the Ising model, maybe it makese more sense to continually sample new values for x_v given x_{-v} until we are satisfied.

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- Compute acceptance probability:

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- Sample $u \sim U(0,1)$
- Set,

$$x_t = \left\{ \begin{array}{ll} x' & \text{if } u < A \\ x_{t-1} & \text{otherwise} \end{array} \right.$$

Why does MH work?

• If we take samples $x_1,...,x_N$ using MH algorithm, why is this equivalent to taking samples from the target distribution p(x)?

Markov chain

Markov chain $\{X_t\}$ is a stochastic process modeling a sequence of events where the probability of each event depends only on the previous event.

 $\bullet \ \ \text{Markov property:} \ p(x_t|x_{1:t-1}) = p(x_t|x_{t-1}).$

Markov chain

Given measurable space $(\mathcal{X},\mathcal{F})$,

$$K: \mathcal{X} \times \mathcal{F} \to [0,1]$$

is referred to as the Markov kernel (a probability measure).

- $\bullet \ \, \mathsf{Each} \,\, \mathsf{random} \,\, \mathsf{variable} \,\, X_t \in \mathcal{X} \\$
- K(x,A) specifies the probability of moving to a set $F \in \mathcal{F}$ given that the chain is in state $x \in \mathcal{X}$.

Markov chain: continuous state space

For continuous state space, $\mathcal{X}=\mathbb{R}$, the transition probability can be described using a density function $K(x_{t-1},x_t)=k(x_t|x_{t-1})$.

Markov chain: discrete state space

For discrete state space, the Markov chain is described using a transition matrix P, where P_{ij} represents the probability of transitioning from state $P(x_t=j|x_{t-1}=i)$.

Markov chain: stationary distribution

The Markov chain $\{X_t\}$ converges to unique **stationary** distribution as $t\to\infty$ if some conditions are satisfied.

A probability distribution π defined on $\mathcal X$ is invariant (stationary) under a Markov kernel K if for all $F\in\mathcal F$

$$\pi(A) = \int \pi(x)K(x,F)dx.$$

For discrete case: $\pi = \pi P$.

Markov chain: detailed balance (reversibility)

A Markov chain with kernel $K: \mathcal{X} \times \mathcal{F}$ satisfies the detailed balance condition with respect to a probability distribution π if,

$$\pi(x)k(x'|x) = \pi(x')k(x|x').$$

Reversibility: probability of being in state x and moving to x' from x is the same as being in state x' and moving to x.

• Note: detailed balance is a stronger condition than stationary condition: if detailed balance is satisfied, π is a stationary distribution of the Markov chain with kernel K.

Markov chain: Ergodicity

- Aperiodic: Markov chain does not return to the same state at some fixed interval.
- Positive recurrent: the expected number of steps for returning to the same state is finite.

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$$q(x'|x)A(x'|x)$$

stay at current state x with probability

$$q(x|x) + q(x'|x)(1 - A(x)).$$

To prove: p(x)k(x'|x) = p(x')k(x|x').

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Case 1:
$$A(x'|x) = \frac{p(x')q(x|x')}{p(x)q(x'|x)} < 1$$
.

$$p(x)q(x'|x)A(x'|x) = p(x)q(x'|x)\frac{p(x')q(x|x')}{p(x)q(x'|x)}$$
(1)

$$= p(x')q(x|x'). (2)$$

Case 2: $A(x'|x) \ge 1$.

$$p(x')q(x|x')A(x|x') = p(x')q(x|x')\frac{p(x)q(x'|x)}{p(x')q(x|x')}$$
(3)

$$= p(x)q(x'|x). (4)$$

Is MH an ergodic Markov chain?

Yes, as long as we choose our proposal carefully.

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- For discrete case, ensure q(x'|x) > 0 for all $x', x \in \mathcal{X}$.

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- Large global proposals tend to be rejected, causing the chain to get stuck at a point for long periods.

Gibbs sampling is an MCMC algorithm, which is well suited for high-dimensional distributions where sampling directly from the joint distribution is difficult.

• Initialize x^0 .

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- ② For t = 1, ..., T:
 - Iterate over each variable x_i :
 - \blacktriangleright Sample $x_i^t \sim p(x_i|x_{-i}^t)$, where x_{-i} refers to all other variables except $x_i.$

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Gibbs sampling is particularly effective when the conditional distributions $p(x_i|x_{-i})$ are easy to sample from.

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$$\begin{split} A(x'|x) &= \frac{p(x')q(x|x')}{p(x)q(x'|x)} \\ &= \frac{p(x_i'|x_{-i})p(x_{-i})q(x|x')}{p(x_i|x_{-i})p(x_{-i})q(x'|x)}. \end{split}$$

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Since $q(x'|x)=p(x_i'|x_{-i})$ and $q(x|x')=p(x_i|x_{-i})$, the acceptance probability simplifies to 1.

Gibbs for Ising model

For t = 1, ..., T:

 $\bullet \ \mbox{Sample} \ x_v \sim p(x_v|x_{-v}) \ \mbox{for each} \ v \in V.$

Sample each variable in turn, conditioned on the values of all of the other variables.

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What does $p(x_v|x_{-v})$ look like?

Gibbs sampling for Ising model

$$p(x_v|x_{-v}) = \frac{p(x_v, x_{-v})}{\sum_{x_v'} p(x_v', x_{-v})}$$
 (5)

$$\propto \exp(-\phi(x_v) - \sum_{(u,v) \in E} \psi(x_u,x_v)). \tag{6}$$

Example: image denoising

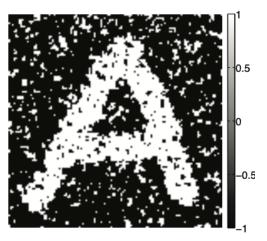


Figure 1: Fig 12.3 (a), PML 2

If all of the neighbors of x_v is white/black, x_v is likely to be white/black.

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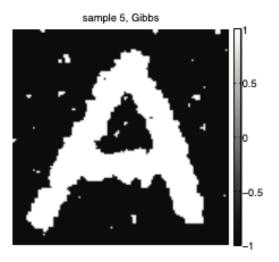


Figure 2: Fig 12.3 (b), PML 2

Example: image denoising

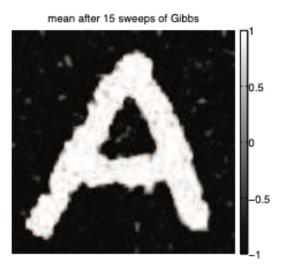


Figure 3: Fig 12.3 (c), PML 2

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- Each node $v \in V$ represents a random variable.
- ullet Edge between $u,v\in V$ is denoted (u,v). Presence of an edge indicates that there is a symmetric relationship between u and v but we cannot easily pinpoint directionality.

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- Example: The value taken at each pixel (random variable X_v) is related to the value taken by its neighbors but it is not causal.

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- \bullet Local Markov property: $X_u \bot X_{rest} | X_{nbr(u)}$, where $nbr(u) = \{v: (u,v) \in E\}.$
- Global Markov property:

Any two sets $A,B\subset V$, are conditionally independent given a separating set S, i.e., $X_A\bot X_B|X_S$, if S separates A and B in G.

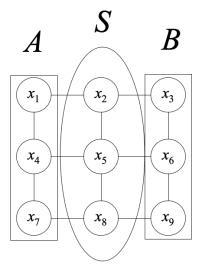


Figure 4: Global Markov property

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MB plays a central role in determining efficient inference algorithm. Example, Gibbs sampling update of a variable X_v is conditioned on its MB and nothing else.

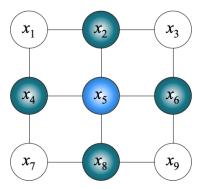


Figure 5: Markov blanket

Undirected graphical models: Hammersley-Clifford Theorem

A strictly positive probability distribution $p(x_V)$ satisfies the global Markov property with respect to G if and only if it can be factorized as,

$$p(x_V) = \frac{1}{Z} \prod_{C \in \mathcal{C}} \psi_C(x_C),$$

- ullet C denotes the set of (maximal) cliques,
- ψ_C denotes potential function for clicque C,
- ullet Z is normalization constant also referred to as partition function.

Clique $C\subseteq V$ of G=(V,E) is a fully connected subgraph of G such that every pair of nodes $u,v\in C$ are adjacent i.e., $\{u,v\}\in E$.

A clique ${\cal C}$ is maximal if adding a node to ${\cal C}$ does not preserve full connectivity.

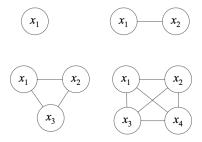


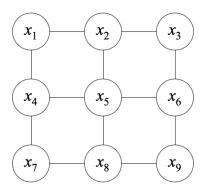
Figure 6: Example: Markov blanket

An edge $\{u,v\}$ is a clique. A fully connected set of nodes is a clique.

Computing partition function is a source of great computational challenge:

$$Z = \int_{\mathcal{X}_V} \prod_{C \in \mathcal{C}} \psi_C(x_C).$$

In most cases, the inference involving UGM requires approximate methods.



What are the maximal cliques in this graph?

Back to Gibbs sampling

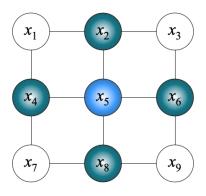
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Back to Gibbs sampling

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Determine the conditional $p(x_v|x_{mb(v)})$.

Back to Gibbs sampling



• Partition the nodes into disjoint sets $A,B\subset V$ such that

$$x_u \perp x_v | B, \quad u, v \in A$$

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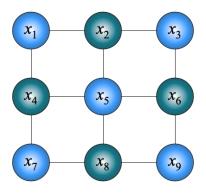
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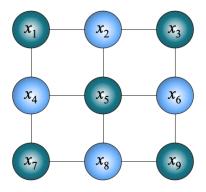
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At each iteration t = 1, ..., T:

- $\bullet \ \ \mathsf{Sample} \ p(x_A|x_{-A})\text{,}$
- Sample $p(x_B|x_{-B})$.





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- Gibbs sampling is efficient when conditional distributions are easy to sample from, leveraging local dependencies.
- MRFs serve as a foundation for probabilistic inference, particularly in structured probabilistic models.

Applications of UGMs

- Neuroscience and associative memory: Hopfield networks (1982, 1984).
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- Natural language processing and large language models (2017)
 - ▶ GPT-based large language models capture token dependencies.
 - ▶ The architecture is not a UGM (transformers use self-attention) but GPT learns long-range dependencies between tokens (subword) in non-sequential manner (not directional).