Efficient Sampling Techniques

Alexey Naumov

HDI Lab, HSE University



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Summary

- 1. Introduction
- 2. Monte-Carlo method
- 3. Rejection sampling
- 4. Importance sampling
- 5. Intro to Markov chains
- 6. MCMC
- 7. Analysis of LD and ULA
- 8. Variance of MCMC estimate

Density estimation

- Classical statistical problem:
 - 1. We have a sample $X_1, \ldots, X_n \in \mathbb{R}^d$ from a density $p_{data}(x)$.
 - 2. Aim: estimate $p_{data}(x)$ and sample from it
- Classical solution: kernel density estimation

$$\pi(x) = \frac{1}{n} \sum_{j=1}^n K_h(X_j - x),$$

where K_h – kernel, h – bandwidth.



• This approach work when d = 1, 2, 3.

Density estimation

- High dimension d > 3.
- Black and white pictures 1024×1024 pixels, dim $d = 2^{20} > 10^6$.
- Other object of interest: video, protein structure, ...
- We need other methods (e.g. GANs)
- How to sample from π ?

Motivation

Bayesian inference and learning. Let θ ∈ Θ be an unknown variable (parameter) and X = (X₁,..., X_N) ∈ X be a data.

1. Posterior distribution: given the prior $p_0(\theta)$ and likelihood $p(X_i|\theta)$

$$\pi(\theta|\mathbf{X}) = \frac{\prod_{i=1}^{N} p(X_i|\theta) p_0(\theta)}{\int\limits_{\Theta} \prod_{i=1}^{N} p(X_i|\theta) p_0(\theta) d\theta}$$

2. Expectation w.r.t. $\pi(\theta|\mathbf{X})$

$$\mathbb{E}_{\pi(\cdot|\mathbf{X})}[f(\theta)] = \int_{\Theta} f(\theta) \pi(\theta|\mathbf{X}) \mathrm{d}\theta$$

Statistical mechanics. Here, one needs to compute the partition function Z of a system with states s and Hamiltonian E(s)

$$Z = \sum_{s} \exp\left\{-\frac{E(s)}{kT}\right\},\,$$

where k is the Boltzmann's constant and T denotes the temperature of the system.

GANs framework

- Generator $G : \mathbb{R}^d \mapsto \mathbb{R}^D$: takes a latent variable z from a prior density $p_0(z), z \in \mathbb{R}^d$, produces $G(z) \in \mathbb{R}^D$ in the observation space;
- Discriminator D : ℝ^D → [0, 1]: takes a sample in the observation space, distinguishes between real examples and fake ones;

GAN training objective

$$L(g,D) := \mathbb{E}_{X \sim p_{\text{data}}}[\log(D(X))] + \mathbb{E}_{Z \sim p_0}[\log(1 - D(g(Z)))] \rightarrow \min_{g \in \mathcal{G}} \max_{D \in \mathcal{D}}.$$

Let $p_d(x)$ and $p_g(x)$ be the densities of real and fake observations;



GANs as an energy-based model

- Main drawback: information accumulated by discriminator is not used during the generation procedure;
- Let $d^*(x) = \text{logit } D^*(x)$, therefore:

$$\frac{p_d(x)}{p_d(x) + p_g(x)} = \frac{1}{1 + \frac{p_g(x)}{p_d(x)}} = \frac{1}{1 + \exp(-d^*(x))}$$

Hence, we can express

$$p_d(x) = p_g(x)e^{d^*(x)}.$$

Let us introduce d(x) = logit D(x) and consider the corresponding energy-based model

$$\hat{p}_d(x) = p_g(x)e^{d(x)}/Z_0\,,$$

where Z_0 is the normalizing constant. If $D(x) \approx D^*(x)$, $\hat{p}_d(x)$ is close to $p_d(x)$;

Sample from $\hat{p}_d(x)$ using MCMC.

GANs as an energy-based model

- Similar idea considered in Turner et al. [2019]; main issue: MCMC in pixel space is highly inefficient;
- ▶ Che et al. [2020] suggested latent-space sampling from the model

 $\hat{p}_d(x) = p_0(z) \exp\left\{ \text{logit}(D(G(z)) \right\}, z \in \mathbb{R}^d,$

where $p_0(z)$ is the generator's prior distribution in the latent space;

Sampling using Langevin-based algorithms, as suggested in Che et al. [2020], can be inefficient, especially if d is large.

Introduction

This Course

We aim at sampling from π and computing expectation

$$\pi(f) := \mathbb{E}[f(X)] = \int_X f(x)\pi(x) \,\mathrm{d}x, \quad f \in \mathsf{L}_2(\pi)$$

We discuss,

- Monte-Carlo method
- Rejection sampling
- Importance sampling
- MCMC
- Mixture of techniques

Monte-Carlo method

• Get an i.i.d. sample $(X_k)_{k=0}^{\infty}$ from π , estimate $\pi(f)$ by

$$\pi_n(f) := \frac{1}{n} \sum_{k=0}^{n-1} f(X_k),$$

Kolmogorov's strong law of large numbers: with probability 1

$$\lim_{n\to\infty}\pi_n(f)=\mathbb{E}[f(X_0)]=\pi(f)$$

- Advantage over deterministic integration: MC positions the integration grid (samples) in regions of high probability.
- Disadvantage: when π(x) has standard form, e.g. Gaussian, it is straightforward to sample from it using easily available routines. However, when this is not the case, we need to introduce more sophisticated techniques.

Monte-Carlo method

Variance:

$$\operatorname{Var}[\pi_n(f)] = \frac{1}{n^2} \sum_{k=0}^{n-1} \operatorname{Var}[f(X_k)] = \frac{\sigma_{\pi}^2(f)}{n}$$

where $\sigma_{\pi}^{2}(f) = \text{Var}[f(X_{0}] = \pi(f^{2}) - \pi^{2}(f).$

Central limit theorem (CLT)

$$\sqrt{n}(\pi_n(f) - \pi(f)) \xrightarrow{\text{Law}} N(0, \sigma_\pi^2(f)) \quad n \to \infty$$

Indeed,

$$\sqrt{n}(\pi_n(f) - \pi(f)) = \frac{\sum_{k=0}^{n-1} (f(X_k) - \mathbb{E}[f(X_k)])}{\sqrt{n}}$$

• Length of confidence interval for $\pi_n(f)$ proportional to $\frac{\sigma_{\pi}(f)}{\sqrt{n}}$

Rejection sampling

- Sample from a distribution π, which is known up to a proportionality constant, by sampling from another easy-to-sample proposal distribution g that satisfies π(x) ≤ Mg(x), M < ∞.</p>
- Algorithm:

Set k = 0; Repeat until k = n - 1

- 1. Sample $X_i \sim q$ and independent $U \sim \text{Uniform}[0, 1]$;
- 2. Accept X_i and set i := i + 1, if

$$U < \frac{\pi(X_i)}{Mg(X_i)}.$$

Otherwise, reject.



Rejection sampling

Advantage: simple

Disadvantage: impractical in high-dimensional scenarios. It is not always possible to bound π(x)/g(x) with a reasonable constant M over the whole space X. If M is too large,

$$\begin{split} \mathsf{P}(X_i \text{ accepted}) &= \mathsf{P}\bigg(U < \frac{\pi(X_i)}{Mg(X_i)}\bigg) = \mathbb{E}\bigg[\mathsf{P}\bigg(U < \frac{\pi(X_i)}{Mg(X_i)}\bigg)\bigg|X_i\bigg] \\ &= \mathbb{E}\bigg[\frac{\pi(X_i)}{Mg(X_i)}\bigg] = \int_{\mathsf{X}} \frac{\pi(x)}{Mg(x)}g(x)\mathrm{d}x = \frac{1}{M} \end{split}$$

will be too small (here we also assume $g(x) > 0, x \in X$)

Rejection sampling

We show that

$$\mathsf{P}\left(X_i \leq x | U < \frac{\pi(X_i)}{Mg(X_i)}\right) = \pi\{(-\infty, x]\}$$

Indeed, let $A = \{X_i \leq x\}, B = \left\{U < \frac{\pi(X_i)}{Mg(X_i)}\right\}$. Then

$$\mathsf{P}(A|B) = \mathsf{P}(B|A) \,\mathsf{P}(A) / \,\mathsf{P}(B).$$

We may check that

$$P(B|A) = \frac{P(A \cap B)}{G(x)} = \frac{1}{G(x)} \mathbb{E}[\mathbb{1}_{A \cap B}]$$

= $\frac{1}{G(x)} \mathbb{E}_{X_i}[\mathbb{1}_A] \mathbb{E}_U[\mathbb{1}_B] = \frac{1}{MG(x)} \mathbb{E}_{X_i}\left[\mathbb{1}_A \frac{\pi(X_i)}{g(X_i)}\right]$
= $\frac{\pi\{(-\infty, x]\}}{MG(x)}.$

Importance sampling

Make change of measure: replace π(x) by another easy-to-sample proposal distribution λ(x):

$$\pi(f) = \int_{\mathsf{X}} f(x)\pi(x) \mathrm{d}x = \int_{\mathsf{X}} f(x)w(x)\lambda(x) \mathrm{d}x,$$

where w(x) – importance weight (Radon-Nikodym derivative)

$$w(x) := \frac{\pi(x)}{\lambda(x)}$$

• Replace $\pi_n(f)$ by $\bar{\pi}_n(f)$,

$$\bar{\pi}_n(f):=\frac{1}{n}\sum_{k=0}^{n-1}f(X_i)w(X_i),$$

where $X_i \sim \lambda$.

Importance sampling

Variance

$${\sf Var}_\lambda[f(X_0)w(X_0)] = \mathbb{E}_\lambda[f^2(X_0)w^2(X_0)] - \pi^2(f)$$

By Jensen's inequality

$$\mathbb{E}_{\lambda}[f^2(X_0)w^2(X_0)] \geq (\mathbb{E}_{\lambda}[|f(X_0)|w(X_0)])^2 = \left(\int_X |f(x)|\pi(x)\mathrm{d}x\right)^2$$

Lower bound is attained for

$$\lambda^*(x) = \frac{|f(x)|\pi(x)|}{\int_X |f(x)|\pi(x)dx|}$$

High sampling efficiency is achieved when we focus on sampling from π in the importance regions where |f(x)|π(x) is relatively large.

Self-Normalized Importance Sampling

π is known up to a normalizing factor Z_π, π(dx) = π̃(dx)/Z_Π;
 Define *importance weights* as w̃(x) = π̃(x)/λ(x);
 Then

$$\pi(f) = \int f(x)\pi(x)dx = \mathsf{Z}_{\pi}^{-1} \int f(x)\tilde{w}(x)\lambda(x)dx$$
$$= \mathsf{Z}_{\pi}^{-1} \int f(x)\tilde{w}(x)\lambda(x)dx / \left\{ \mathsf{Z}_{\pi}^{-1} \int \tilde{w}(x)\lambda(x)dx \right\}$$

The self-normalized importance sampling (SNIS) estimator of π(f) is then given by

$$\widehat{\pi}_N(f) = \sum_{i=1}^N \omega_N^i f(X_i),$$

where

$$X_i \sim \lambda, \omega_N^i = rac{ ilde{w}(X_i)}{\sum_{j=1}^N ilde{w}(X_j)}, i \in \{1, \dots, N\}.$$

MCMC

- What can be done if drawing i.i.d. samples from π is not an option?
- ▶ If we run the (ergodic) Markov chain $(Z_k)_{k\geq 0}$ for a long time (started from anywhere), then for large *N* the distribution of Z_N will be approximately invariant: Law $(Z_N) \approx \pi$. We can then set $X_1 = Z_N$, and then restart and rerun the Markov chain to obtain X_2, X_3, \ldots , and then do estimates as in MC,

$$\pi_n(f) = \frac{1}{n} \sum_{k=0}^{n-1} f(X_k)$$

Important question

How to construct P(x, A) such that the distribution of X_n converges to invariant distribution π as quickly as possible for arbitrary initial distribution ξ ?

What to read?

For more details see Douc et al. [2018]

Define a Markov chain (i.e., discrete time).

Ingredients of the definition:

- ▶ X state space (e.g. $X \subset \mathbb{R}^d$), $\mathcal{X} \sigma$ -algebra of X
- Initial distribution $X_0 \sim \xi$;

▶ Transition kernel P(x, A), where $x \in X, A \in \mathcal{X}$:

$$\mathsf{P}(X_{n+1} \in A | X_n = x) = \mathsf{P}(x, A)$$

• Markov property: X_{n+1} depends only on X_n ;

Example: Model $X_0 \sim \xi$ and for $n \geq 1$

$$X_n = F(X_{n-1}, \varepsilon_n)$$

where $(\varepsilon_n)_{n\geq 1}$ is an i.i.d. sequence independent of $\sigma\{X_k, 0 \leq k \leq n-1\}$ and F is some function, $F: X \times \mathbb{R}^{d'} \to X$

Markov chains: gym

- More about MK kernels
- Ergodicity (finite case)
- Ergodicity (not in this course:()
- Ready for MCMC

Action on measures

Let $\boldsymbol{\mu}$ be a probability measure on X

$$\mu P(A) = \int_{\mathsf{X}} \mu(\mathrm{d} x) \, \mathsf{P}(x, A)$$

Action on functions

$$\mathsf{P} f(x) = \int_{\mathsf{X}} f(y) \mathsf{P}(x, \mathrm{d}y)$$

Composition of kernels

$$\mathsf{P}^n(x,A) = \int_{\mathsf{X}} \mathsf{P}(x,\mathrm{d} y) \, \mathsf{P}^{n-1}(y,A)$$

(Kolmogorov-Chapman equation)

Tensor product (kernel \otimes kernel)

$$\mathsf{P} \otimes \mathsf{P} f(x) = \int_{\mathsf{X}} \mathsf{P}(x, \mathrm{d}y) \int_{\mathsf{X}} f(y, z) \mathsf{P}(y, \mathrm{d}z)$$
$$= \int_{\mathsf{X} \times \mathsf{X}} f(y, z) \mathsf{P}(x, \mathrm{d}y) \mathsf{P}(y, \mathrm{d}z)$$

Take $f(y, z) = 1(y \in A, z \in B)$. Then

$$\mathsf{P}\otimes\mathsf{P}\,f(x)=\mathsf{P}(X_1\in A,X_2\in B|X_0=x)=\mathsf{P}^{\otimes 2}(x,A imes B)$$

Tensor product (measure \otimes kernel)

$$\begin{split} \xi \otimes \mathsf{P} f &= \int_{\mathsf{X}} \xi(\mathrm{d} y) \int_{\mathsf{X}} f(y, z) \, \mathsf{P}(y, \mathrm{d} z) \\ &= \int_{\mathsf{X} \times \mathsf{X}} f(y, z) \xi(\mathrm{d} y) \, \mathsf{P}(y, \mathrm{d} z) \end{split}$$

Invariant distribution

Distrbution π is invariant w.r.t. P if

$$\pi P = \pi$$

Theorem

Let $(X_k)_{k=0}^{\infty}$ be a MC with initial distribution π and kernel P. $(X_k)_{k=0}^{\infty}$ is stationary iff π is invariant.

Proof.

Let $(X_k)_{k=0}^{\infty}$ be stationary. Then $Law(X_1) = Law(X_0)$. Hence, $\pi P(A) = P_{\pi}(X_1 \in A) = P(X_0 \in A) = \pi(A)$. If π is invariant, then the distribution of (X_n, \ldots, X_{n+k}) is $\pi P^n \otimes P^{\otimes k} = \pi \otimes P^{\otimes k}$ is independent of n

Reversibility

Distribution $\boldsymbol{\xi}$ is reversible w.r.t. P if

$$\xi \otimes \mathsf{P}(A \times B) = \xi \otimes \mathsf{P}(B \times A)$$

If X is countable,

$$\xi(x) \mathsf{P}(x, x') = \xi(x') \mathsf{P}(x', x)$$

Detailed balance equation.

$$\begin{split} \mathbb{E}_{\xi}[f(X_0, X_1)] &= \int_{\mathsf{X} \times \mathsf{X}} \xi(\mathrm{d} x_0) \, \mathsf{P}(x_0, \mathrm{d} x_1) f(x_0, x_1) \\ &= \int_{\mathsf{X} \times \mathsf{X}} \xi(\mathrm{d} x_0) \, \mathsf{P}(x_0, \mathrm{d} x_1) f(x_1, x_0) = \mathbb{E}_{\xi}[f(X_1, X_0)] \end{split}$$

Hence, $\mathsf{Law}(X_0, X_1) = \mathsf{Law}(X_1, X_0)$

Theorem

Let P be a MK. If ξ is reversible w.r.t. P then ξ is invariant.

Proof.

$$\begin{split} \xi \, \mathsf{P}(\mathcal{A}) &= \xi \otimes \mathsf{P}(\mathsf{X} \times \mathsf{A}) = \xi \otimes \mathsf{P}(\mathsf{A} \times \mathsf{X}) \\ &= \int_{\mathsf{X}} \xi(\mathrm{d} x) \, \mathsf{P}(x,\mathsf{X}) \mathbf{1}_{\mathsf{A}}(x) = \xi(\mathsf{A}) \end{split}$$

Let X be finite, $\mathsf{X} = [1, \dots, r]$

Total variation distance (finite case)

Let μ, ξ be probability measures on X. Define

$$\mathsf{d}_{\mathsf{TV}}(\xi,\mu) := \frac{1}{2} \sum_{\mathsf{i}=1}^{\mathsf{r}} |\mu(\mathsf{i}) - \xi(\mathsf{i})| = \sum_{\mathsf{i}:\mu(\mathsf{i}) > \xi(\mathsf{i})} (\mu(\mathsf{i}) - \xi(\mathsf{i}))$$

Clearly, $d_{TV} \leq 1$.

Denote J := {i : μQ(i) > ξQ(i)}. Let Q be an arbitrary MK. Then for any μ, ξ

$$d_{\mathsf{TV}}(\mu \mathsf{Q}, \xi \mathsf{Q}) = \sum_{j \in J} (\mu \mathsf{Q}(j) - \xi \mathsf{Q}(j))$$

= $\sum_{j \in J} \sum_{i \in \mathsf{X}} (\mu(i) \mathsf{Q}(i, j) - \xi(i) \mathsf{Q}(i, j))$
 $\leq \sum_{i: \mu(i) > \xi(i)} (\mu(i) - \xi(i)) \sum_{j \in J} \mathsf{Q}(i, j) \leq \mathsf{d}_{\mathsf{TV}}(\mu, \xi)$ (2)

▶ Let $Q(i,j) \ge a > 0$ for any $i, j \in X$. Then $\exists j' \notin J$ and hence for any $i \in X$ $\sum_{i \in J} Q(i,j) < 1 - a$

Eq. (2) my be improved:

$$\mathsf{d}_{\mathsf{TV}}(\mu\mathsf{Q},\xi\mathsf{Q}) < (1-\mathsf{a})\mathsf{d}_{\mathsf{TV}}(\mu,\xi)$$

Assume

$$\exists s: \mathsf{P}^{s}(x, x') > 0 \text{ for any } x, x' \in \mathsf{X}$$
(3)

• Let us fix arbitrary distribution μ_0 and denote $\mu_n = \mu_0 P^n$. Then

$$d_{\mathsf{TV}}(\mu_{n},\mu_{n+k}) = d_{\mathsf{TV}}(\mu_{0} \mathsf{P}^{n},\mu_{0} \mathsf{P}^{n+k})$$

$$\leq (1-a)d_{\mathsf{TV}}(\mu_{0} \mathsf{P}^{n-s},\mu_{0} \mathsf{P}^{n+k-s})$$

$$\leq (1-a)^{m}d_{\mathsf{TV}}(\mu_{0} \mathsf{P}^{n-ms},\mu_{0} \mathsf{P}^{n+k-ms}),$$
(4)

where $m: 0 < n - ms \le s$. Take *n* large such that $(1 - a)^m < \varepsilon$. Then $\{\mu_n\}_{n\ge 1}$ is a Cauchy sequence.

Set

$$\pi:=\lim_{n\to\infty}\mu_n.$$

Then

$$\pi \mathsf{P} = \lim_{n \to \infty} \mu_n \mathsf{P} = \lim_{n \to \infty} \mu_0 \mathsf{P}^{n+1} = \pi$$

• Uniqueness: Assume $\pi_1 \neq \pi_2$ such that $\pi_1 P = \pi_1, \pi_2 P = \pi_2$. Then $\pi_i = \pi_i P^s, i = 1, 2$ and

$$\mathsf{d}_{\mathsf{TV}}(\pi_1,\pi_2) \leq (1-\mathsf{a})\mathsf{d}_{\mathsf{TV}}(\pi_1,\pi_2)$$

Hence, $\pi_1 = \pi_2$.

$$d_{\mathsf{TV}}(\mu_0 \mathsf{P}^n, \pi) = d_{\mathsf{TV}}(\mu_0 \mathsf{P}^n, \pi \mathsf{P}^n) \le (1-\mathsf{a})^m d_{\mathsf{TV}}(\mu_0 \mathsf{P}^{n-\mathsf{ms}}, \pi \mathsf{P}^{n-\mathsf{ms}})$$
$$\le (1-\mathsf{a})^m \le (1-\mathsf{a})^{n/s-1} = (1-\mathsf{a})^{-1}\beta^n,$$
(5)

where $\beta = (1 - a)^{1/s} < 1$.

Theorem

Assume (3) and let π be an invariant distribution. Then for any $f : X \to \mathbb{R}$, with probability 1:

$$\lim_{n\to\infty}\frac{1}{n}\sum_{k=0}^{n-1}f(X_k)=\pi(f)$$

Compare with SLLN for i.i.d. sequence.

MCMC

- What can be done if drawing i.i.d. samples from π is not an option?
- ▶ If we run the (ergodic) Markov chain $(Z_k)_{k\geq 0}$ for a long time (started from anywhere), then for large *N* the distribution of Z_N will be approximately invariant: Law $(Z_N) \approx \pi$. We can then set $X_1 = Z_N$, and then restart and rerun the Markov chain to obtain X_2, X_3, \ldots , and then do estimates as in MC,

$$\pi_n(f) = \frac{1}{n} \sum_{k=0}^{n-1} f(X_k)$$

Important question

How to construct P(x, A) such that the distribution of X_n converges to invariant distribution π as quickly as possible for arbitrary initial distribution ξ ?

Example: Metropolis-Hastings algorithm

Let $Q(x, A) = \int_A q(x, y) dy$ be some MK (e.g. Gaussian)

- 1. Choose X_0 .
- 2. Given X_k , a candidate move Y_{k+1} is sampled from $Q(X_k, \cdot)$
- 3. $X_{k+1} = Y_{k+1}$ with probability $\alpha(X_k, Y_{k+1})$, otherwise $X_{k+1} = X_k$, where acceptance ratio

$$\alpha(x,y) = \min\left\{1, \frac{\pi(y)q(y,x)}{\pi(x)q(x,y)}\right\}$$

Example: Random walk MH Take $q(x, y) = \overline{q}(y - x)$, where $\overline{q}(x) = \overline{q}(-x)$. Then

$$Y_{k+1} = X_k + Z_{k+1}, \quad Z_{k+1} \sim \overline{q}$$

In this case

$$\alpha(x,y) = \min\left\{1,\frac{\pi(y)}{\pi(x)}\right\}$$

Example: Langevin Dynamics

Langevin Dynamics Itô SDE:

$$\mathrm{d}\theta_t = -\nabla U(\theta_t) \,\mathrm{d}t + \sqrt{2} \mathrm{d}W_t,$$

Invariant measure: $\pi(\theta) = e^{-U(\theta)}$ and $Law(\theta_t) \to \pi$ as $t \to \infty$.

1. Take
$$\pi(\theta) = (2\pi)^{-1/2} e^{-\theta^2/2}$$

- 2. SDE: $d\theta_t = \theta_t dt + \sqrt{2} dW_t$, θ_0 is independent of W. This is Ornstein–Uhlenbeck process
- 3. Apply Ito's formula to obtain

$$\theta_t = \theta_0 \mathrm{e}^{-t} + \sqrt{2} \int_0^t e^{-(t-s)} \mathrm{d}W_s$$

 Since the Itô integral of deterministic integrand is normally distributed, we readily have

$$\mathsf{Law}(\theta_t) = \mathcal{N}(\theta_0 \mathrm{e}^{-t}, 1 - \mathrm{e}^{-2t}) \to \mathcal{N}(0, 1)$$

Example: Langevin Dynamics

Itô SDE:

$$\mathrm{d}\theta_t = -\nabla U(\theta_t) \,\mathrm{d}t + \sqrt{2} \mathrm{d}W_t,$$

Invariant measure: $\pi(\theta) = e^{-U(\theta)}$

1. First-order discretization (Unadjusted Langevin Algorithm, ULA):

$$Y_{k+1} = Y_k - \gamma \nabla U(Y_k) + \sqrt{2\gamma} Z_{k+1}, \quad i.i.d. \ Z_k \sim \mathsf{N}(\mathsf{0},\mathsf{I}_d)$$

Equivalently, $Y_{k+1} \sim \mathcal{N}(Y_k - \gamma \nabla U(Y_k), 2\gamma I)$

- Metropolis-adjusted Langevin Algorithm (MALA): ULA + Metropolis-Hastings correction;
- 3. Demo: https://chi-feng.github.io/mcmc-demo
- 4. If we can't calculate ∇U replace it by its estimate over batch (SGLD, SGLD-FP, SAGA etc)

SGLD

1. Posterior distribution:

$$\pi(\theta|\mathbf{X}) = \frac{\prod_{i=1}^{N} p(X_i|\theta) \pi_0(\theta)}{\int\limits_{\mathbb{R}^d} \prod_{i=1}^{N} p(X_i|\theta) \pi_0(\theta) \, d\theta} \propto e^{-U(\theta)},$$

where $U = \log \pi_0(\theta) + \sum_{i=1}^N \log p(X_i|\theta);$

- 2. A computational bottleneck: calculating the full gradient ∇U scaling proportionally to N can be very time consuming in the "big data" limit;
- 3. Replace $\nabla U(\theta)$ by an unbiased estimate. This gives rise to the SGLD algorithm, where the parameters are updated according to

$$\theta_{k+1} = \theta_k - \gamma G(\theta_k, S_{k+1}) + \sqrt{2\gamma} \xi_{k+1},$$

$$G(\theta, S) = \nabla U_0(\theta) + K M^{-1} \sum_{i \in S} \nabla U_i(\theta),$$
(6)

where each S_{k+1} is a random batch taking values in S_M (here S_M is the set of all subsets S of $\{1, \ldots, N\}$ with |S| = M) which is sampled from a uniform distribution over S_M independently of \mathfrak{F}_k (here $(\mathfrak{F}_k)_{k\geq 0}$ is the filtration generated by $\{(\theta_\ell, S_\ell)\}_{\ell\geq 0}$).

4. Note that $\mathbb{E}[G(\theta_k, S_{k+1})|\mathfrak{F}_k] = \nabla U(\theta_k)$ and therefore $G(\theta_k, S_{k+1})$ is an unbiased estimate of $\nabla U(\theta_k)$.

Transition kernel of MH algorithm

Let $Q(x, A) = \int_A q(x, y) dy$ be some MK (e.g. Gaussian) 1. Choose X_0 .

- 2. Given X_k , a candidate move Y_{k+1} is sampled from $Q(X_k, \cdot)$
- 3. $X_{k+1} = Y_{k+1}$ with probability $\alpha(X_k, Y_{k+1})$, otherwise $X_{k+1} = X_k$, where acceptance ratio

$$\alpha(x,y) = \min\left\{1, \frac{\pi(y)q(y,x)}{\pi(x)q(x,y)}\right\}$$

MH transition kernel

$$\mathsf{P}(x,A) = \int_{A} \alpha(x,y) q(x,y) \mathrm{d}y + \overline{\alpha}(x) \delta_{x}(A),$$

where

$$\overline{\alpha}(x) = \int_{\mathsf{X}} (1 - \alpha(x, y)) q(x, y) \mathrm{d}y.$$

Invariance of π

Theorem

Distribution π is reversible w.r.t. P.

Proof.

We need to show that for any ${\it C} \in {\it X} \times {\it X}$

$$\int_{\mathsf{X}\times\mathsf{X}} \pi(x) \mathrm{d}x \,\mathsf{P}(x,\mathrm{d}y) \mathbf{1}_{\mathsf{C}}(x,y) = \int_{\mathsf{X}\times\mathsf{X}} \pi(y) \mathrm{d}y \,\mathsf{P}(y,\mathrm{d}x) \mathbf{1}_{\mathsf{C}}(x,y)$$

For any $x, y \in X$

$$\pi(x)\alpha(x,y)q(x,y) = \{\pi(x)q(x,y)\} \lor \{\pi(y)q(y,x)\} = \pi(y)\alpha(y,x)q(y,x)$$

Moreover,

$$\int_{X \times X} \pi(x) dx \delta_x(dy) \overline{\alpha}(x) \mathbf{1}_{\mathcal{C}}(x, y) = \int_X \pi(x) dx \overline{\alpha}(x) \mathbf{1}_{\mathcal{C}}(x, x)$$
$$= \int_X \pi(y) dy \overline{\alpha}(y) \mathbf{1}_{\mathcal{C}}(y, y) = \int_{X \times X} \pi(y) dy \delta_y(dx) \overline{\alpha}(y) \mathbf{1}_{\mathcal{C}}(x, y)$$

Analysis of ULA

• Let
$$\pi(x) = Z_d^{-1} e^{-U(x)}$$
;

L-smooth potential

U is L–smooth is $U\in \mathcal{C}^2(\mathbb{R}^d)$ and there exists L>0 such that

$$\|\nabla U(x) - \nabla U(y)\| \le L \|x - y\|$$

for any $x, y \in \mathbb{R}^d$.

Unadjusted Langevin Algorithm, ULA:

$$X_{k+1} = X_k - \gamma \nabla U(Y_k) + \sqrt{2\gamma} Z_{k+1}, \quad i.i.d. \ Z_k \sim \mathsf{N}(\mathsf{0},\mathsf{I}_d)$$

• Denote $\mathsf{P}_{\gamma}(x, \cdot) = \mathcal{N}(x - \gamma \nabla U(x), 2\gamma \mathsf{I}).$

Kantorovich-Wasserstein distance

Kantorovich-Wasserstein distance

For λ, ν , we denote their coupling set by $\Pi(\lambda, \nu)$, i.e. $\xi \in \Pi(\lambda, \nu)$ is the measure on X × X satisfying for all $A \in \mathcal{B}(X)$, $\xi(A, X) = \lambda(A)$ and $\xi(X, A) = \nu(A)$. For $p \ge 1$ and λ, ν , let

$$W_{p,d}(\lambda,\nu) := \inf_{\Pi(\lambda,\nu)} \left\{ \int_{\mathsf{X}\times\mathsf{X}} \mathsf{d}^p(x,y) \,\xi(\mathrm{d} x,\mathrm{d} y) \right\}^{1/p}$$

be the Kantorovich–Wasserstein distance of order p between λ and ν .

Analysis of ULA

A1

U is L-smooth and m-strongly convex:

$$\langle \nabla U(x) - \nabla U(y), x - y \rangle \ge m \|x - y\|^2.$$

Theorem

For any $\gamma \in (0, m/L^2)$ there exists invariant distribution π_{γ} :

$$W_2^2(\delta_x \mathsf{P}_{\gamma}^k, \pi_{\gamma}) \leq (1 - m\gamma)^k \int ||x - y||^2 \pi_{\gamma}(\mathrm{d}y)$$

Analysis of ULA

Fix $x, \tilde{x} \in \mathbb{R}^d$. Synchronous coupling:

$$\begin{aligned} X_{k+1} &= X_k - \gamma \nabla U(X_k) + \sqrt{2\gamma} Z_{k+1}, \\ \tilde{X}_{k+1} &= \tilde{X}_k - \gamma \nabla U(\tilde{X}_k) + \sqrt{2\gamma} Z_{k+1} \end{aligned}$$

Then

$$egin{aligned} \|X_{k+1}- ilde{X}_{k+1}\|^2 &= \|X_k- ilde{X}_k\|^2 \ \gamma^2 \|
abla \mathcal{U}(X_k)-
abla \mathcal{U}(ilde{X}_k)\|^2 \ &-2\gamma \langle X_k- ilde{X}_k,
abla \mathcal{U}(X_k)-
abla \mathcal{U}(ilde{X}_k)[] \end{aligned}$$

Use A1:

$$\begin{split} \|X_{k+1} - ilde{X}_{k+1}\|^2 &\leq (1 + \gamma^2 L^2 - 2\gamma m) \|X_k - ilde{X}_k\|^2 \ &\leq (1 - \gamma m) \|X_k - ilde{X}_k\|^2. \end{split}$$

Hence

$$W_2^2(\delta_x \mathsf{P}_\gamma^k, \delta_{\widetilde{x}} \mathsf{P}_\gamma^k) \leq (1 - m\gamma)^k W_2^2(\delta_x, \delta_{\widetilde{x}})$$

• We may show that $(\lambda P_{\gamma}^{k})_{k \in \mathbb{N}}$ is a Cauchy sequence and there exists $\pi_{\gamma}^{\lambda} = \pi_{\gamma}$, moreover $\pi_{\gamma} P_{\gamma} = \pi_{\gamma}$.

Variance of MCMC estimate

Let π be an invariant distribution. Assume $X_0 \sim \pi,$ i.e. we start from the invariant distribution. Then

$$\operatorname{Var}_{\pi}\left[n^{-1}\sum_{k=0}^{n-1}f(X_{k})\right] = \frac{\operatorname{Var}_{\pi}[f]}{n} + \frac{1}{n^{2}}\sum_{i\neq j}\mathbb{E}_{\pi}\left[(f(X_{i}) - \pi(f))(f(X_{j}) - \pi(f))\right] = \frac{\rho^{(f)}(0)}{n} + \frac{2}{n}\sum_{k=1}^{n-1}\left(1 - \frac{k}{n}\right)\rho^{(f)}(k) \neq \frac{\operatorname{Var}_{\pi}[f]}{n}$$

where

$$\rho^{(f)}(k) = \mathbb{E}_{\pi} \left[(f(X_0) - \pi(f))(f(X_k) - \pi(f)) \right]$$

Variance of MCMC estimate

Under appropriate conditions (e.g. \u03c6-irreducibility + apereodicity + existence of solution of Poisson eq.) CLT holds:

$$\frac{1}{\sqrt{n}}\sum_{i=0}^{n-1}[f(X_i)-\pi(f)]\xrightarrow{Law}\mathcal{N}(0,V_{\infty}(f)),$$

where
$$V_{\infty}(f) := \lim_{n \to \infty} \operatorname{Var}_{\pi} \left[\frac{1}{\sqrt{n}} \sum_{i=0}^{n-1} (f(X_i) - \pi(f)) \right]$$

• Length of confidence interval for $\pi_n(f)$ proportional to $\frac{\sqrt{V_{\infty}(f)}}{\sqrt{n}}$

Ex²MCMC: Sampling through Exploration Exploitation

Importance Sampling procedure

• Aim: sample from π and estimate $\pi(f) = \int_{\mathbb{R}^D} f(x) \pi(dx)$;

- π is known up to a normalizing factor Z_{Π} , $\pi(dx) = \tilde{\pi}(dx)/Z_{\Pi}$;
- Importance Sampling (IS) consists of re-weighting samples from a proposal distribution λ.
- Define importance weights as $\tilde{w}(x) = \tilde{\pi}(x)/\lambda(x)$;
- The self-normalized importance sampling (SNIS) estimator of π(f) is then given by

$$\widehat{\pi}_N(f) = \sum_{i=1}^N \omega_N^i f(X^i) \,,$$

where

$$X^{1:N} \sim \lambda, \omega_N^i = rac{ ilde{w}(X')}{\sum_{j=1}^N ilde{w}(X^j)}, i \in \{1, \dots, N\}.$$

From IS to SIR

- Sampling counterpart of the IS procedure is known as Sampling Importance Resampling (SIR; Rubin [1987]);
- Sample X¹,..., X^N i.i.d. from λ and compute the importance weights ω¹_N,..., ω^N_N;
- Sample Y¹,..., Y^M from X¹,..., X^N with replacement, and with probabilities proportional to the weights ω¹_N,..., ω^N_N. That is, we sample from the empirical distribution

$$\hat{\pi}(\mathrm{d} x) = \sum_{i=1}^{N} \omega_{N}^{i} \delta_{X^{i}}(\mathrm{d} x),$$

where $\delta_y(dx)$ denotes the Dirac mass at y.

- As $N \to \infty$, $Y^1, \ldots, Y^M \sim \hat{\Pi}$ will be distributed according to π .
- Main drawback: the described procedure is only asymptotically valid.

Iterated SIR (i-SIR) algorithm

Iterating samples from λ , we arrive at iterated SIR algorithm (i-SIR, Andrieu et al. [2010], and Andrieu et al. [2018]).

Algorithm 1: Single stage of i-SIR algorithm

Input : Sample Y_j from previous iteration Output: New sample Y_{j+1} 1 Set $X_{j+1}^1 = Y_j$ and draw $X_{j+1}^{2:N} \sim \lambda$. 2 for $i \in [N]$ do 3 compute the normalized weights $\omega_{i,j+1} = \tilde{w}(X_{j+1}^i) / \sum_{k=1}^N \tilde{w}(X_{j+1}^k)$. 4 Set $I_{j+1} = \operatorname{Cat}(\omega_{1,j+1}, \dots, \omega_{N,j+1})$. 5 Draw $Y_{j+1} = X_{j+1}^{l_{j+1}}$.

The Markov chain $\{Y_k, k \in \mathbb{N}\}$ generated by i-SIR has the following Markov kernel

$$\mathsf{P}_{N}(x,\mathsf{A}) = \int \delta_{x}(\mathrm{d}x^{1}) \sum_{i=1}^{N} \frac{\tilde{w}(x^{i})}{\sum_{j=1}^{N} \tilde{w}(x^{j})} \mathbb{1}_{\mathsf{A}}(x^{i}) \prod_{j=2}^{N} \lambda(\mathrm{d}x^{j}).$$
(7)

i-SIR algorithm

Provided also that $|\tilde{w}|_{\infty} < \infty$, it was shown in Andrieu et al. [2018] that the Markov kernel P_N is uniformly geometrically ergodic. Namely, for any initial distribution ξ on (X, \mathcal{X}) and $k \in \mathbb{N}$,

$$\|\xi \mathsf{P}_{N}^{k} - \pi\|_{\mathrm{TV}} \le \kappa_{N}^{k},\tag{8}$$

with $\epsilon_N = \frac{N-1}{2L+N-2}$, $L = |\tilde{w}|_{\infty}/\lambda(\tilde{w})$ and $\kappa_N = 1 - \epsilon_N$.

- Note that the bound (8) relies significantly on the restrictive condition that weights are uniformly bounded |w̃|∞ < ∞.</p>
- Moreover, even when this condition is satisfied, the rate κ_N can be close to 1 when the dimension *d* is large.
- Indeed, consider a simple scenario π(x) = Π^d_{i=1} p(x_i) and λ(x) = Π^d_{i=1} q(x_i) for some densities p(·) and q(·) on ℝ. Then it is easy to see that L = (sup_{y∈ℝ} p(y)/q(y))^d grows exponentially with d.

i-SIR algorithm

To illustrate this phenomenon, we consider a simple problem of sampling from the standard normal distribution $\mathcal{N}(0, I_d)$ with the proposal $\mathcal{N}(0, 2 I_d)$ in increasing dimensions d up to 300.



Figure: Sampling from $\mathcal{N}(0, I_d)$ with the proposal $\mathcal{N}(0, 2I_d)$. We display confidence intervals for i-SIR and Ex²MCMC obtained from 100 independent runs as blue and red regions, respectively. Ex²MCMC helps to achieve efficient sampling even in high dimensions.

- Main i-SIR drawback: absence of local exploration moves;
- Idea: apply a local MCMC kernel R (*rejuvenation kernel*) after each i-SIR step;
- R has π as invariant distribution;
- Here comes Ex²MCMC : Exploration steps through i-SIR, Exploitation steps through R(x, ·);
- As our default choice we consider MALA as rejuvenation, but other ones (HMC, NUTS) are also possible.

Algorithm 2: Single stage of Ex^2MCMC algorithm with independent proposals

1 **Procedure** $Ex^2MCMC(Y_i, \Lambda, R)$: **Input** : Previous sample Y_i ; proposal distribution Λ ; rejuvenation kernel R; **Output:** New sample Y_{i+1} ; Set $X_{i+1}^1 = Y_j$, draw $X_{i+1}^{2:N} \sim \lambda$; 2 for $i \in [N]$ do 3 compute the normalized weights 4 $\omega_{i,i+1} = \tilde{w}(X_{i+1}^{i}) / \sum_{k=1}^{N} \tilde{w}(X_{i+1}^{k});$ Set $I_{i+1} = Cat(\omega_{1,i+1}, ..., \omega_{N,i+1});$ 5 Draw $Y_{i+1} \sim \mathsf{R}(X_{i+1}^{I_{i+1}}, \cdot)$. 6

V-geometric ergodicity

A Markov kernel Q with invariant probability measure π is *V*-geometrically ergodic if there exist constants $\rho \in (0, 1)$ and $M < \infty$ such that, for all $x \in X$ and $k \in \mathbb{N}$,

$$\|\mathbf{Q}^k(\mathbf{x},\cdot)-\pi\|_V \leq M \{V(\mathbf{x})+\pi(V)\}\rho^k.$$

Assumptions

A1

(i) R has π as its unique invariant distribution; (ii) There exists a function $V: X \to [1, \infty)$, such that for all $r \ge r_R > 1$ there exist $\lambda_{R,r} \in [0, 1)$, $b_{R,r} < \infty$, such that $RV(x) \le \lambda_{R,r}V(x) + b_{R,r}\mathbb{1}_{V_r}$, where $V_r = \{x: V(x) \le r\}$;

A2

(i) For all
$$r \ge r_{\mathsf{R}}$$
, $\tilde{w}_{\infty,r} := \sup_{x \in V_r} \{\tilde{w}(x)/\lambda(\tilde{w})\} < \infty$;
(ii) $\operatorname{Var}_{\lambda}[\tilde{w}]/\{\lambda(\tilde{w})\}^2 < \infty$.

Theorem

Let A1 and A2 hold. Then, for all $x \in X$ and $k \in \mathbb{N}$,

$$\|\mathsf{K}_{\mathsf{N}}^{k}(x,\cdot)-\pi\|_{\mathsf{V}} \leq c_{\mathsf{K}_{\mathsf{N}}}\{\pi(\mathsf{V})+\mathsf{V}(x)\}\tilde{\kappa}_{\mathsf{K}_{\mathsf{N}}}^{k},\qquad (9)$$

where c_{K_N} , $\tilde{\kappa}_{K_N} \in [0, 1)$ are some constants. In addition, $c_{K_N} = c_{K_{\infty}} + O(N^{-1})$ and $\tilde{\kappa}_{K_N} = \tilde{\kappa}_{K_{\infty}} + O(N^{-1})$.

Toy example



Figure: Single chain mixing visualization. – Blue color levels represent the target 2d density. Random chain initialization is noted in black, 100 steps are plotted per sampler: the size of each red dot corresponds to the number of consecutive steps the walkers remains at a given location. For MALA, we generate 300 samples and choose each 3-rd one for comparability. Note that the variance of the global proposal (dotted countour lines) should be relatively large to cover well all the modes. The step size of MALA also can not be increased much to keep reasonable acceptance ratio.

Adaptive proposals

- Consider family of proposals {λ_θ}, θ ∈ ℝ^D, chosen to match the target distribution π̃;
- Let $T : \mathbb{R}^d \to \mathbb{R}^d$ be smooth and invertible. Denote by $T \# \Lambda$ the distribution of Y = T(X) with $X \sim \lambda$;
- The corresponding density is given by λ_T(y) = λ(T⁻¹(y)) J_{T⁻¹}(y), where J_T denotes the Jacobian determinant of T;

Adaptive proposals: learning procedure

 Disperancy measure: linear combination of forward and backward KL divergence (generalizations to [Papamakarios et al., 2021] possible);

Forward and backward KL:

$$\mathcal{L}^{f}(\theta) = \int \log \frac{\pi(x)}{\lambda_{\theta}(x)} \pi(x) \mathrm{d}x ,$$

$$\mathcal{L}^{b}(\theta) = \int \log \frac{\lambda(x)}{\pi(T_{\theta}(x)) \mathsf{J}_{T_{\theta}}(x)} \lambda(x) \mathrm{d}x .$$

• Given a sample $Y_k \sim \pi$ and $Z_k \sim \lambda$ for $k \in [K]$, by

$$\begin{split} \widehat{\nabla \mathcal{L}^{f}}(Y^{1:K},\theta) &= -\frac{1}{K}\sum_{k=1}^{K} \nabla \log \lambda_{\theta}(Y_{k}), \\ \widehat{\nabla \mathcal{L}^{b}}(Z^{1:K},\theta) &= -\frac{1}{K}\sum_{k=1}^{K} \nabla \log \big(\tilde{\pi}(T_{\theta}(Z_{k}) \, \mathsf{J}_{T_{\theta}}(Z_{k}) \big). \end{split}$$

Following Gabrié et al. [2021], we consider

$$\widehat{\mathcal{L}}(Y^{1:K}, Z^{1:K}, \theta) = \alpha \widehat{\mathcal{L}^f}(Y^{1:K}, \theta) + \beta \widehat{\mathcal{L}^b}(Z^{1:K}, \theta).$$

FIEx²MCMC algorithm with adaptive proposals

Algorithm 3: Single stage of FIEx²MCMC. Steps of Ex²MCMC are done in parallel with common values of proposal parameters θ_j . Step 4 updates the parameters using the gradient estimate obtained from all the chains.

Input : weights θ_j , batch $Y_j^{1:K}$ Output: new weights θ_{j+1} , batch $Y_{j+1}^{1:K}$ 1 for $k \in [K]$ do 2 $\lfloor Y_{j+1,k} = \text{Ex}^2 \text{MCMC} (Y_{j,k}, T_{\theta_j} \# \Lambda, R)$ 3 Draw $\overline{Z}^{1:K} \sim \lambda$.

4 Update
$$heta_{j+1} = heta_j - \gamma \widehat{
abla \mathcal{L}}(Y_{j+1}, \overline{Z}, heta_j).$$

Practical note

In our experiments: T_{θ} is modelled as a normalizing flow based on RealNVP architecture (Dinh et al. [2017]).

Take-home Messages & Future Works

- We know basics of MC, rejection sampling, importance samling, MCMC, normalizing flows
- ▶ To become world expert in Markov chains read Douc et al. [2018]
- We are ready for 'real' projects (join HDI Lab team)

Thank you!

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