Geometric Data analysis Random walks, Sampling, Volume

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Introduction

- Introductory Notions
- Exact sampling

- Motivation
- Sampling algorithms
- MCMC Diagnostics
- - Reduction to Multiphase Monte Carlo
 - Simulated annealing for cooling convex bodies
- - Cutting planes
 - Simulated Annealing



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- We represent a distribution Q with a Probability Density Function (PDF) π(x), x ∈ K, where K is the support of π(x).
- The support is the subset of \mathbb{R}^n which $\pi(x)$ does not map to zero.



Here the support is the polytope P and the distribution Q is the uniform distribution over P, i.e. $\pi(x) = 1/\text{vol}(P)$.

• When a random variable follows X a distribution Q with PDF $\pi(x)$ then,

$$\Pr[X \in A] = Q(A) = \int_A \pi(x) dx$$
, where $A \subseteq K$.

- $\int_{K} \pi(x) dx = 1.$
- A function $f : \mathbb{R}^d \to \mathbb{R}_+$ induces a PDF $\pi(x) \propto f(x)$ when there is (possibly unknown) normalizing constant C such that $\pi(x) = f(x)/C$.



• When $\pi(x) \propto f(x)$ we say that $\pi(x)$ is proportional to f(x).

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Uniform Sampling from the hypersphere

- To sample uniformly from the boundary of a hypersphere with radius r:
 - 1. Sample *d* numbers g_1, \ldots, g_d from $\mathcal{N}(0, 1)$.
 - 2. The point $v = r(g_1, \ldots, g_d) / \sqrt{\sum g_i^2}$ is uniformly distributed on the surface of the *d*-dimensional hypersphere with radius *r* and centered at the origin.
- To sample uniformly from the interior of a hypersphere with radius *r*:
 - 1. Sample a point $v \sim \mathcal{U}(\partial B_d)$ and $u \sim \mathcal{U}(0, 1)$.
 - 2. The point $p = ru^{1/d}v$ is uniformly distributed in the interior of the hypersphere with radius r and centered at the origin.

To pick a random direction through point $p \in \mathbb{R}^d$ we sample from the surface of a hypersphere centered at p.



Uniform Sampling from the simplex

- 1. [Smith and Tromble'04]:
 - Generate distinct: $x_0 < x_1 < \cdots < x_{d+1} \in \mathbb{N}^*$. Return y:

$$y_i = \frac{x_i - x_{i-1}}{M}$$
, $i = 1, \dots, d+1$. M: largest integer.

- To guarantee distinct choice we use a variation of Bloom filter.
- Sampling one point takes $O(d \log d)$.
- 2. [Rubinstein and Melamed'98]:
 - Generate independent unit-exponential random variables, X_1, \dots, X_{d+1} . Return $Y \in \mathbb{R}^{d+1}$: $Y_i = X_i / \sum_{i=1}^{d+1} X_i$.
 - Sampling one point takes O(d).



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The problem is the computation of a multidimensional definite integral,

$$I = \int_P f(\mathbf{x}) d\mathbf{x}$$

Given $\mathbf{x}_1, \ldots, \mathbf{x}_N$ uniformly distributed samples from P,

$$R_N = V \frac{1}{N} \sum_{i=1}^N f(\mathbf{x}_i), \ V = \operatorname{vol}(P) = \int_P d\mathbf{x}$$

Then $\lim_{N\to\infty} R_N = I$

In general for $\mathbb{E}[f(\mathbf{x})] = \int_P f(\mathbf{x}) \pi_P(\mathbf{x}) d\mathbf{x}$ sample N i.i.d. points from π_P and $\mathbb{E}[f(\mathbf{x})] \approx \frac{1}{N} \sum_{i=1}^N f(\mathbf{x}_i)$.



Some easy cases

Some elementary polytopes have determinantal formulas.



$$\begin{vmatrix} 1 & 2 & 1 \\ 3 & 6 & 1 \\ 6 & 1 & 1 \end{vmatrix} / 2! = 11$$
$$\begin{vmatrix} 2 & 5 \\ 4 & 0 \end{vmatrix} = 20$$

Geometric Data analys

Problem: Minimize a linear function $f(\mathbf{x}) = \mathbf{c} \cdot \mathbf{x}$ in body K. Answer: Sample from $\pi_T(\mathbf{x}) \propto e^{-\mathbf{c} \cdot \mathbf{x}/T}$, for $T = T_0 > \cdots > T_I$.



A sample from π_{T_I} is ϵ -close to the optimal solution with high probability.

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- Let $\pi(x) = f(x)/C$, $x \in \mathbb{R}^d$, where f(x) is an *unnormalized* density and $C \in \mathbb{R}$ a normalizing constant.
- Let h(x) a PDF that can be simulated by some known method and $f(x) \le kh(x)$, where $k \in \mathbb{R}$ is a constant.

To obtain a random variate from $\pi(x)$,

- 1. Generate a candidate Z from h(x) and a value u from U(0, 1), the uniform distribution on (0, 1).
- 2. If $u \leq f(Z)/kh(Z)$ return Z.
- 3. Otherwise goto 1.

Acceptance-rejection sampling Drawbacks

• Sampling/rejections techniques (sample from bounding box) fail in high dimensions



$$\frac{\operatorname{vol}(unitball)}{\operatorname{vol}(unitcube)} = O((1/d)^d)$$

Markov Chain Monte Carlo sampling

A MCMC sampling algorithm is applied on a continuous state space $K \subseteq \mathbb{R}^d$. The algorithm,

- Starts at a point $x_0 \in K$.
- When being at the point x_i moves to the next point x_{i+1} according to a transition kernel p_x(A).
- The transition kernel of a MCMC algorithm gives the probability to jump from x to a set A ⊆ K.
- For example $p_x(K) = 1$.



To sample from a density $\pi(x)$ define a random walk on a continuous state space with a transition kernel $p_x(A)$ such that,

1. [Convergence]

$$\int_{P} p_{x}(A)\pi(x)dx = \int_{A} \pi(y)dy$$

Then $\pi(x)$ is called target density.

2. [Uniqueness] $\lim_{n\to\infty} p_x^n(A) = \int_A \pi(y) dy$, where

$$p_x^n(A) = \int_P p_x^{n-1}(y) p_y(A) dy,$$

the transition kernel of the n-th iteration.

[Understanding the Metropolis-Hastings Algorithm, '95].

Ball walk

Ball Walk(P, p, δ, f): Polytope $P \subset \mathbb{R}^d$, point $p \in P$, radius $\delta, f : \mathbb{R}^d \to \mathbb{R}_+$

- 1. Pick a uniform random point x from the ball of radius δ centered at p
- 2. return x with probability min $\left\{1, \frac{f(x)}{f(p)}\right\}$; return p with the remaining probability.



- When the density is not restricted to a body then the algorithm is known as the **Metropolis-Hastings** algorithm.
- Task: write the pseudocode for the special case of uniform sampling.

Hit and Run(P, p, f): Polytope $P \subset \mathbb{R}^d$, point $p \in P$, $f : \mathbb{R}^d \to \mathbb{R}_+$

- 1. Pick a line ℓ through p.
- 2. **return** a random point on the chord $\ell \cap P$ chosen from the distribution $\pi_{\ell,f}$ restricted in *P*.



- When the density is not restricted to a body then the algorithm samples from $\pi_{\ell,f}.$
- Task: write the pseudocode for the special case of uniform sampling.
- **Q**: How can we compute the $\ell \cap P$?

- 1. Generate the length of the trajectory $L = -\tau \ln \eta$, $\eta \sim U(0, 1)$.
- 2. Pick a uniform direction v to define the trajectory.
- 3. When the trajectory meets a boundary with internal normal s, ||s|| = 1, the direction is changed as $v \leftarrow v 2 < v, s > s$.
- 4. **return** the end of the trajectory as p_{i+1} . If the number of reflections exceeds *R* **return** $p_{i+1} = p_i$.



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- Uniform sampling from the hypercube $[-1,1]^{200}$ and projection to \mathbb{R}^3 .
- Rows: Ball Walk, Coordinate Directions Hit and Run, Random Directions Hit and Run, Billiard Walk.
- Columns: walk length, {1, 50, 100, 150, 200}

Limitations of BW and HnR

- Their mixing time is $O^*(d^3)$ for log-concave distributions.
- Their performance is crucially affected by the starting point.
- Typically a warm start is required. A distribution S is said to be *M*-warm with respect to the distribution Q if,

$$M = \sup_{A \in P} \frac{S(A)}{Q(A)}$$

• They perform better when the distribution is (approximately) isotropic. A distribution Q is said to be isotropic if

$$\mathbb{E}_Q[X] = 0$$
, and $\mathbb{E}_Q[XX^T] = I_d$

Limitations of BW and HnR

- They spend many steps around the mode of the distribution.
- Consider the spherical Gaussian centered at the origin with $\sigma^2=0.1.$
- The mode of the PDF in two dimensions it is 1/9 and in three dimensions it is only 1/27 of the volume of the cube.



• Q: In 20 dimensions?

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 The problem that motivates us is to compute the expectation of a function *f*, say E_π[*f*], which reduces to the integral,

$$\mathbb{E}_{\pi}[f] = \int_{P} f(x)\pi(x)dx \tag{1}$$

- In high dimensions a probability density, π(x), will concentrate around its mode (local maximum of π(x)).
- Contributions to the expectation are determined by the product of density and volume, π(x)dx.
- The points with dominant contribution to (1) concentrate in a neighborhood called the typical set.



- We can interpret the mode of the target density as a massive planet and the gradient of the target density as that planet's gravitational field.
- The typical set becomes the space around the planet through which we want a satellite to orbit .

- HMC defines trajectories that guide the walk inside the typical set.
- The choice of the derivative of π(x) at the current point p of the walk would be wrong as it points directly towards its mode.
- The Hamiltonian dynamics behind HMC operate on a position vector **p** and a velocity **v**.

 \bullet The system is described by a function of ${\bf p}$ and ${\bf v}$ known as the Hamiltonian,

$$H(\mathbf{p}, \mathbf{v}) = U(\mathbf{p}) + K(\mathbf{v}) = -\log(\pi(\mathbf{p})) + \frac{1}{2}|\mathbf{v}|^2.$$

To sample from π , one has to solve the following system of Ordinary Differential Equations (ODE):

$$\frac{d\mathbf{p}}{dt} = \frac{\partial H(\mathbf{p}, \mathbf{v})}{\partial \mathbf{v}} \qquad \Rightarrow \qquad \begin{cases} \frac{d\mathbf{p}(t)}{dt} = \mathbf{v}(t) \\ \frac{d\mathbf{v}}{dt} = -\frac{\partial H(\mathbf{p}, \mathbf{v})}{\partial \mathbf{p}} \qquad \Rightarrow \qquad \begin{cases} \frac{d\mathbf{v}(t)}{dt} = \mathbf{v}(t) \\ \frac{d\mathbf{v}(t)}{dt} = \nabla \log(\pi(\mathbf{p})) \end{cases}$$
(2)

• When the density is restricted in a Polytope *P* then HMC walks on trajectories inside *P*.



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How can we evaluate the quality of a sample obtained by a random walk?

- [Convergence diagnostics for Markov chain Monte Carlo, Vivekananda Roy, '19].
- [Revisiting the Gelman-Rubin Diagnostic, Dootika Vats, Christina Knudson, '20].

A MCMC convergence diagnostic can also be used as a termination criterion for sampling.
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- open source, written in C++
 https://github.com/GeomScale/volume_approximation
- R package in CRAN https://CRAN.R-project.org/package=volesti.
- Python interface.
- since 2018.

Problem	currently	soon	description
volume computation	\checkmark		8 algo. / thousands of dimensions / fastest practical estimation
sampling distributions			
uniform / gaussian/ Exp	\checkmark		4 algo. / thousands of dimensions
log-concave densities	√		HMC / Langevin Diffusion
convex optimization Semidefinite Programming Linear Programming	✓		beating SDPA / working to improve goal: best open source
multivariate integration simple MC integration importance sampling	 ✓ 		hundreds of dimensions goal: best open source approximation
Preprocessing	\checkmark		3 rounding algo. / 4 MCMC diagnostics

GeomScale/volesti on Google Summer of Code 2020 Mentoring organization



- https://geomscale.github.io/
- https://summerofcode.withgoogle.com/organizations/5673184117915648/

Three student projects this year:

- 1. Sampling log-concave densities.
- 2. Convex optimization.
- 3. Uniform sampling / metabolic networks in biology.



- stan is a platform for statistical modeling.
- Provides HMC implementations.
- https://mc-stan.org/.



- cobra is the state-of-the-art package for the analysis of metabolic networks.
- Provides three random walks for uniform / Gaussian sampling from convex polytopes.
- https://github.com/opencobra.

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Computing the exact volume of P,

- is #P-hard for all the representations [DyerFrieze'88]
- is open if both H- and V- representations available
- is APX-hard (oracle model) [Elekes'86]

Theorem

[Dyer, Frieze, Kannan'91] For any convex body P and any $0 \le \epsilon$, $\delta \le 1$, there is a randomized algorithm which computes an estimate V s.t. with probability $1 - \delta$ we have $(1 - \epsilon)vol(P) \le V \le (1 + \epsilon)vol(P)$, and the number of oracle calls is $poly(d, 1/\epsilon, log(1/\delta))$.

• Using randomness, we can go from an exponential approximation to an arbitrarily small one.

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Let a sequence of functions $\{f_0, \ldots, f_m\}$, $f_i : \mathbb{R}^d \to \mathbb{R}$. Then,

$$\operatorname{vol}(P) = \int_{P} dx = \int_{P} f_{m}(x) dx \ \frac{\int_{P} f_{m-1}(x) dx}{\int_{P} f_{m}(x) dx} \cdots \frac{\int_{P} f_{0}(x) dx}{\int_{P} f_{1}(x) dx} \ \frac{\int_{P} dx}{\int_{P} f_{0}(x) dx}$$

Then select f_i s.t.,

- The number of phases, *m*, is as small as possible.
- Each integral ratio can be efficiently estimated by sampling from $\pi \propto f_i$ restricted to *P* (using geometric random walks).
- There is a closed formula for $\int_P f_m(x) dx$.

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complexity = #phases $\times \#$ points per phase \times cost per point

Authors-Year	Complexity (oracle calls)	fi	random walk
[Dyer, Frieze, Kannan'91]	$O^{*}(d^{23})$	Indicator function of a ball	grid walk
[Kannan, Lovasz, Simonovits'97]	$O^*(d^5)$	Indicator function of a ball	ball walk
[Lovasz, Vempala'03]	$O^{*}(d^{4})$	Exponential	hit-and-run
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Practical algorithms:

• Follow theory but make practical adjustments (experimental).

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- [Emiris, Fisikopoulos'14] Sequence of balls + coordinate hit-and-run.

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- Multiphase Monte Carlo algorithm with statistical tests.
- Sequence of scaled copies of any convex body (generalization of sequence of balls).
- Faster practical algorithm for zonotopes and V-polytopes besides H-polytopes. Performs computations in:
 - thousands of dimensions for H-polytopes in few hours.
 - 100 dimensions for zonotopes and V-polytopes in \leq 1hour.

Multiphase Monte Carlo

 Let C_m ⊆ · · · ⊆ C₁ a sequence of concentric balls intersecting P, s.t. C_m ⊆ P ⊆ C₁.



• Construct a sequence of balls intersecting *P*, then:

$$\mathsf{vol}(P) = \mathsf{vol}(P \cap C_m) \frac{\mathsf{vol}(P \cap C_{m-1})}{\mathsf{vol}(P \cap C_m)} \cdots \frac{\mathsf{vol}(P \cap C_1)}{\mathsf{vol}(P \cap C_2)} \frac{\mathsf{vol}(P)}{\mathsf{vol}(P \cap C_1)}$$

Ratio estimation

• Estimate $r_i = \frac{\text{vol}(P \cap C_{i+1})}{\text{vol}(P \cap C_i)}$ within some target relative error ϵ_i .

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Ratio estimation

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• Keep each ratio bounded, then $N = O(1/\epsilon_i^2)$ points suffices.

Given convex bodies $P_1 \supseteq P_2$, we define two statistical tests:

 $\begin{array}{l} [\textbf{U-test}(P_1, P_2)] \ H_0: \ vol(P_2)/vol(P_1) \ge r + \delta \\ [\textbf{L-test} \ (P_1, P_2)] \ H_0: \ vol(P_2)/vol(P_1) \le r \end{array}$

- The **U-test** and **L-test** are successful iff both H₀ are rejected.
- If both **U-test** and **L-test** are successful then $r_i = \operatorname{vol}(P_{i+1})/\operatorname{vol}(P_i) \in [r, r+\delta]$, with high probability.

How to fix the sequence of balls

Let C_d the unit ball. Given $C_i = q_i C_d$, $C_m = q_m C_m$, with $q_i > q_m$,

Problem: Compute a new ball
$$C_{i+1}$$
 with radius $q_m < q_{i+1} < q_i$,
s.t. $\frac{\operatorname{vol}(P \cap C_{i+1})}{\operatorname{vol}(P \cap C_i)} \in [r, r+\delta]$.

Answer: binary search to compute $q_{i+1} \in [q_m, q_i]$ until both U-test $(P \cap C_i, P \cap C_{i+1})$ and L-test $(P \cap C_i, P \cap C_{i+1})$ are successful.



Apostolos Chalkis (Athens, Greece)

Geometric Data analysis

- Our algorithm terminates with constant probability.
 - Bound the probability that the construction of the sequence of bodies in MMC fails.

• #phases
$$m = O\left(log(vol(P)/vol(P \cap C_m))\right)$$
.

• If the body we use in MMC is a good fit to P the $vol(P \cap C_m)$ increases and the number of phases m decreases.

• Use the generators of a zonotope *P* to define a centrally symmetric H-polytope that is a good fit to *P*.



 $r = 0.8, r + \delta = 0.85.$ #phases: Left m = 5. Right m = 1.

Comparison with other software



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Problem

Given *P* a convex body in \mathbb{R}^n :

• minimize a convex function *f* in *P* (convex optimization).



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Goal: Randomized approximation algorithms based on sampling from ${\it P}$ with geometric random walks.



- The objective function is linear $f(\mathbf{x}) = \mathbf{c} \cdot \mathbf{x}$.
- The body is given as an intersection of *m* half-spaces.

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H-polytope : $P = \{x \mid Ax \leq b, A \in \mathbb{R}^{m \times n}, b \in \mathbb{R}^m\}$



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Spectrahedron : $P = \{x \mid A_0 + x_1A_1 + \dots + x_dA_d \succeq 0\}$, where A_i : symmetric matrices, $B \succeq 0$: B is positive semidefinite



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 - Exact sampling

2 MCMC sampling

- Motivation
- Sampling algorithms
- MCMC Diagnostics
- 3 Software
- Volume approximation
 - Reduction to Multiphase Monte Carlo
 - Simulated annealing for cooling convex bodies
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• Let $rB_d \subseteq K \subseteq RB_d$.

 \bullet The expected number of phases s.t. $|f_I-f^*|<\varepsilon$ is,

$$I = \left\lceil \frac{1}{\ln(N+1)} d \ln(R/\epsilon) \right\rceil = O^*(d)$$

• Total number of uniform points minimized for N = 1.

Total cost,

$$\left[d \ln(R/\epsilon) \right] imes \ ext{cost per point}$$

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Exponential sampling and Simulated Annealing Kalai, Vempala, 06'

Problem: Minimize a linear function $f(\mathbf{x}) = \mathbf{c} \cdot \mathbf{x}$ in body K. Answer: Sample from $\pi_T(\mathbf{x}) \propto e^{-\mathbf{c} \cdot \mathbf{x}/T}$, for $T = T_0 > \cdots > T_I$.



A sample from π_{T_l} is ϵ -close to the optimal solution with high probability.

Simulated Annealing

Fix the sequence of Temperatures



• The sequence $T_0 > \cdots > T_I$ is fixed s.t. the L_2 norm of π_{T_i} w.r.t. $\pi_{T_{i+1}}$ is bounded by a constant,

$$\|\pi_{T_{i}}/\pi_{T_{i+1}}\| = \mathbb{E}_{\pi_{T_{i}}}\left[\frac{d\pi_{T_{i}}}{d\pi_{T_{i+1}}}\right] = \int_{\mathcal{K}} \frac{\pi_{T_{i}}(x)}{\pi_{T_{i+1}}(x)} \pi_{T_{i}}(x) dx = O(1)$$

• Then π_{T_i} is a warm start for $\pi_{T_{i+1}}$ (Hit-and-Run).

Convergence to the optimal solution

• Starting with $T_0 = R$ (uniform distribution is a warm start).

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Simulated Annealing

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• Knowing that for a temp. T,

$$\mathbb{E}_{\pi_{\mathcal{T}}}[\mathbf{c}\cdot\mathbf{x}] \leq dT + \min_{x\in\mathcal{K}}\mathbf{c}\cdot\mathbf{x}$$

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- $I = O^*(\sqrt{d})$ phases suffices to obtain a solution $|f_I f^*| \le \epsilon$.
- No sequence of distributions $\propto f_i(\mathbf{c} \cdot \mathbf{x})$ can, in general, solve the problem in less than $\Omega(\sqrt{d})$ phases.