

# Sampling and Diffusion Model

Handstein Wang

Institute of Computational Mathematics and Scientific/Engineering Computing  
Academy of Mathematics and Systems Science  
Chinese Academy of Sciences, China

November 7, 2025

# Outline

---

1. Two Settings of Sampling with Applications
2. Density Estimation
3. Generative Model
4. Score Matching
5. Diffusion Model

# Outline

---

## 1. Two Settings of Sampling with Applications

## 2. Density Estimation

## 3. Generative Model

## 4. Score Matching

## 5. Diffusion Model

# Sampling

---

**Target:** We want to sample from a distribution  $\mu$  on  $(\mathbb{R}^d, \mathcal{B}(\mathbb{R}^d))$  which admits a density w.r.t. Lebesgue measure  $m$ , also denoted by  $\mu(x)$ .

There are two settings of sampling problems:

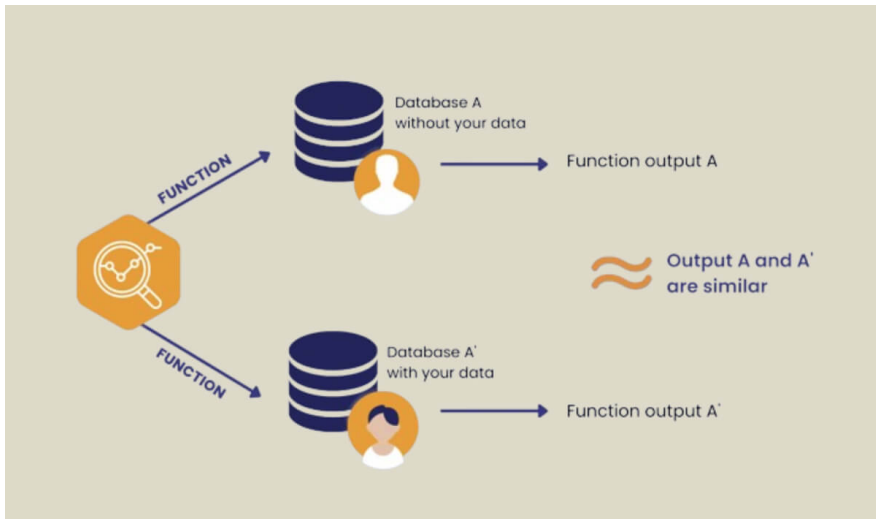
**Setting 1:**  $\mu$  is given in explicit form up to a normalization constant.

Applications: Bayesian inference, inverse problem, finance, computation of high dimensional integral, differential privacy, approximate computation, uncertainty quantification, etc.

**Setting 2:**  $\mu$  is given with a collection of i.i.d. samples.

Applications: generative models (diffusion models, GANs, etc).

# Differential Privacy



# Differential Privacy

- Let  $(\mathcal{X}, \mathcal{X})$  be a measurable space of individual records. For a fixed  $n \in \mathbb{N}$ , datasets live in  $(\mathcal{X}^n, \mathcal{X}^{\otimes n})$ .
- Fix a adjacency relation  $\sim \subseteq \mathcal{X}^n \times \mathcal{X}^n$  specifying which dataset pairs differ by “one individual”.
- Let  $(Y, \mathcal{Y})$  be a measurable output space equipped with a  $\sigma$ -finite base measure  $\mu$ . A randomized algorithm (mechanism) is a Markov kernel

$$K : \mathcal{X}^n \times \mathcal{Y} \rightarrow [0, 1], \quad (D, A) \mapsto K(D, A),$$

i.e., for each  $D$ ,  $K(D, \cdot)$  is a probability measure on  $(Y, \mathcal{Y})$ , and for each  $A$ , the map  $D \mapsto K(D, A)$  is  $\mathcal{X}^{\otimes n}$ -measurable.

## Definition ( $\epsilon$ -DP)

The mechanism  $K$  is  $\epsilon$ -differentially private (w.r.t.  $\sim$ ) if for all adjacent datasets  $D \sim D'$  and all measurable events  $A \in \mathcal{Y}$ ,

$$K(D, A) \leq e^\epsilon K(D', A).$$

# Differential Privacy

- A *score function* is a measurable function  $u : \mathcal{X}^n \times \mathcal{Y} \rightarrow \mathbb{R}$ .
- The *global sensitivity* of  $u$  w.r.t.  $\sim$  is

$$\Delta u := \sup_{D \sim D'} \operatorname{ess\,sup}_{y \in \mathcal{Y}} |u(D, y) - u(D', y)| \in (0, \infty).$$

## Definition (Exponential Mechanism)

Fix  $\varepsilon > 0$  and set  $\alpha := \varepsilon / (2\Delta u)$ . For each dataset  $D \in \mathcal{X}^n$ , the *Exponential Mechanism* is the Markov kernel  $K_{\text{exp}}$  given by

$$K_{\text{exp}}(D, A) := \frac{\int_A \exp(\alpha u(D, y)) \mu(dy)}{\int_{\mathcal{Y}} \exp(\alpha u(D, y)) \mu(dy)}, \quad A \in \mathcal{Y}.$$

Equivalently,  $K_{\text{exp}}(D, \cdot)$  has density proportional to  $\exp(\alpha u(D, \cdot))$  w.r.t.  $\mu$ .

# Differential Privacy

---

## Theorem

*The exponential mechanism  $K_{\text{exp}}$  is  $\varepsilon$ -differentially private, namely, for all adjacent datasets  $D \sim D'$  and all measurable events  $A \in \mathcal{Y}$ ,*

$$K_{\text{exp}}(D, A) \leq e^{\varepsilon} K_{\text{exp}}(D', A).$$

By the theorem above, it suffices to sample from the distribution

$$\pi(dy) \propto \exp(\alpha u(D, y)) \mu(dy)$$

which falls under Setting 1.



# Approximate Computation

---

## Example

Let  $K \subset \mathbb{R}^n$  be a bounded convex set given via a *membership oracle*, i.e., for any query point  $x \in \mathbb{R}^n$  we can decide whether  $x \in K$ . The task is to approximate  $\text{Vol}(K)$ .

- It is known that, under a membership oracle, exact volume computation is computationally intractable: computing the volume of a polytope is NP-hard [Dyer and Frieze, 1988].
- Formulate this as sampling from the uniform law on  $K$  with unnormalized density

$$\mu(x) \propto \mathbf{1}_K(x),$$

by using some sampling technique, the approximate volume computation achieves constant error with  $O(n^3)$  complexity [Jia et al., 2021].

# Outline

---

1. Two Settings of Sampling with Applications

**2. Density Estimation**

3. Generative Model

4. Score Matching

5. Diffusion Model

# Reducing the Problem in Setting 2 to Setting 1

---

In setting 2,  $\mu$  is given with a collection of i.i.d. samples  $x_1, x_2, \dots, x_N$ .

## Density Estimation

We can estimate an explicit expression of  $\mu$  from the i.i.d. samples  $x_1, x_2, \dots, x_N$  by *density estimation*.

There are many methods for density estimation, see [Scott, 2015] for an overview. By density estimation, we can reduce the Problem in Setting 2 to Setting 1.

**But density estimation can NOT avoid the curse of dimensionality!**

# Curse of Dimensionality of Density Estimation

Let  $n \in \mathbb{N}$  be the dimension and  $\beta > 0$  the smoothness parameter. Write  $\beta = \alpha + m$  with  $m \in \mathbb{N}$  and  $\alpha \in (0, 1]$ . Define the class of probability densities  $\mathcal{P}_\beta$  on  $[0, 1]^n$  by those  $f$  satisfying:

- (1)  $f$  is a probability density on  $[0, 1]^n$  and is bounded, say 2.
- (2) The  $m$ -th derivative  $f^{(m)}$  is  $\alpha$ -Hölder continuous, i.e., for all  $x, y \in [0, 1]^n$ ,

$$|f^{(m)}(x) - f^{(m)}(y)| \leq \|x - y\|^\alpha.$$

## Theorem ([Tsybakov, 2009])

Given  $N$  i.i.d. samples  $x_1, \dots, x_N$  from a pdf  $f \in P_\beta$ , the minimax risk of an estimation  $\hat{f}$  of  $f$  under the quadratic loss function  $\ell(\hat{f}, f) := \|\hat{f} - f\|_2^2 = \int_{[0,1]^n} (f(x) - \hat{f}(x))^2 dx$  satisfies

$$\inf_{\hat{f}} \sup_{f \in P_\beta} \|\hat{f} - f\|_2^2 \gtrsim N^{-\frac{2\beta}{n+\beta}}.$$

The infimum is taken over all estimators  $\hat{f}$  built on the data  $x_1, \dots, x_N$ .

# Outline

---

1. Two Settings of Sampling with Applications
2. Density Estimation
- 3. Generative Model**
4. Score Matching
5. Diffusion Model

## Solving Sampling Problems in Setting 2

---

In 2021, Song and Ermon [Song and Ermon, 2019] categorized the existing generative modeling techniques into two major categories: **likelihood-based models** and **implicit generative models**.

**Likelihood-based models**, which directly models the distribution's probability density function via (approximate) maximum likelihood. That is, we model

$$\mu(x) = p_{\theta}(x),$$

where  $p_{\theta}(x)$  is the probability density parametrized by some parameter  $\theta$ . We learn  $\theta$  by maximizing the log-likelihood of the data

$$\max_{\theta} \sum_{i=1}^N \log p_{\theta}(x_i).$$

This category includes autoregressive models, normalizing flow models, energy-based models, and variational auto-encoders (VAEs).

## Solving Sampling Problems in Setting 2

---

**Implicit generative models**, which implicitly represent the probability distribution via a model of sampling process. That is, we say the target distribution is close to a transformation of a Gaussian

$$g(Z), \quad Z \in \mathcal{N}(0, I_m),$$

where  $g : \mathbb{R}^m \rightarrow \mathbb{R}^n$  is a measurable mapping to be learned. Then with a distance between two measures  $\text{dist}(\cdot, \cdot)$ , we want to find  $g$  to minimize

$$\text{dist}\left(\text{Law}(g(Z)), \frac{1}{N} \sum_{i=1}^n \delta_{x_i}\right).$$

The most prominent example is generative adversarial networks (GANs), where new samples are synthesized by transforming a random Gaussian vector with a neural network  $g$ . The parameters of the neural network  $g$  are learned via minimizing the adversarial loss between newly generated images and the empirical measure.

## Solving Sampling Problems in Setting 2

---

- **Likelihood-based models:** Either impose strong architectural restrictions to make the normalizing constant tractable for likelihood computation, or rely on surrogate objectives that only approximate maximum likelihood.
- **Implicit generative models:** Typically require adversarial training, which is notoriously unstable and prone to mode collapse.

Here we introduce another way to represent probability distributions that

- is an iterative sampling process, which does not rely on a good distance between two measures,
- models the score function, rather than the likelihood, avoiding the normalizing constant.

The key idea is to model the gradient of the log probability density function, a quantity often known as the **(Stein) score function**. Such score-based models are not required to have a tractable normalizing constant, and can be directly learned by score matching.



# Outline

---

1. Two Settings of Sampling with Applications
2. Density Estimation
3. Generative Model
- 4. Score Matching**
5. Diffusion Model

# Score Matching

## Definition (Score function)

Let  $\mu$  be a probability measure on  $\mathbb{R}^d$  with density also denoted by  $\mu(x)$  with respect to Lebesgue measure satisfying  $\mu \in W_{\text{loc}}^{1,1}(\mathbb{R}^d)$  and  $\mu(x) > 0$  a.e. Then the *score* of  $\mu$  is the vector field defined by

$$s(x) := \nabla \log \mu(x) \quad \text{a.e.},$$

where  $\nabla \log \mu$  is to be interpreted in the sense of *weak derivatives*.

Equivalently,  $s(x) = (s_1(x), \dots, s_d(x))$  is the unique vector field satisfying, for all  $\varphi \in C_c^\infty(\mathbb{R}^d)$  and each  $i = 1, \dots, d$ ,

$$\int_{\mathbb{R}^d} \partial_i \varphi d\mu = - \int_{\mathbb{R}^d} \varphi s_i d\mu.$$

Even if we only know  $\mu$  up to a normalizing constant,  $\mu \propto e^{-f}$ , we can compute the score function without knowing the normalizing constant

$$\nabla \log \mu(x) = \nabla \log e^{-f} = -\nabla f(x).$$

# Score Matching

---

Now we assume the score of  $\mu$  exists and  $\nabla \log \mu \in L^2(\mu)$ .

To estimate the score function, it is natural to parametrize the score function by a parameter  $\theta$  via  $s_\theta(x)$ , and try to minimize its distance to the true score function over data sampled from the target measure:

$$\min_{\theta} \mathbb{E} \|\nabla \log \mu(X) - s_\theta(X)\|_2^2, \quad X \sim \mu. \quad (\text{SM})$$

In general, it is difficult to evaluate the objective function above when  $\mu$  is represented via  $N$  samples.  $\nabla \log \mu(x)$  is difficult to evaluate with only data samples, for this, a family of methods called score matching is introduced.

# Denoising Score Matching

Instead of solving score matching problem (SM), we consider the following denoising score matching problem

$$\min_{\theta} \text{DSM}(s_{\theta}) := \mathbb{E} \left\| \nabla \log \mu_Y(Y) - s_{\theta}(Y) \right\|_2^2, \quad , \quad (\text{DSM})$$

where  $Y = aX + \sigma Z$ ,  $a \in \mathbb{R}$ ,  $\sigma > 0$ ,  $X \sim \mu$ ,  $Z \sim N(0, I_d)$ ,  $Z$  is independent of  $X$  and  $\mu_Y$  is the density of  $Y$  w.r.t. Lebesgue measure .

## Theorem

*Under the above assumptions and define*

$$\text{DSM}'(s_{\theta}) = \mathbb{E} \left[ \left\| s_{\theta}(Y) + \frac{1}{\sigma} Z \right\|_2^2 \right].$$

*Then*

$$\text{DSM}'(s_{\theta}) = \text{DSM}(s_{\theta}) + \frac{1}{\sigma^2} \mathbb{E} \|Z - \mathbb{E}[Z|Y]\|_2^2.$$

# Denoising Score Matching

The main idea of the proof is using the Stein's lemma to show the following lemma.

## Lemma

$$\nabla \log \mu_Y(Y) = -\frac{1}{\sigma} \mathbb{E}[Z|Y] \quad a.s.$$

By the previous theorem,

$$\arg \min_{\theta} DSM(s_{\theta}) = \arg \min_{\theta} DSM'(s_{\theta}) = \arg \min_{\theta} \mathbb{E} \left[ \left\| s_{\theta}(Y) + \frac{1}{\sigma} Z \right\|_2^2 \right].$$

Then, the objective in the right hand side can be replaced with an empirical version

$$\min_{\theta} \frac{1}{N} \sum_{i=1}^N \left\| s_{\theta}(ax_i + \sigma z_i) + \frac{1}{\sigma} z_i \right\|_2^2$$

**Remark:** The score we learned here is the score of  $\mu_Y = a_{\#} \mu * N(0, \sigma^2 I_d)$  rather than the score of  $\mu$ , which provides a good estimate of  $\mu$  when  $\sigma$  is small enough.

# Score-based Langevin Monte Carlo Algorithm

---

Recall the Langevin Monte Carlo algorithm for sampling  $\mu \propto e^{-f}$ ,

$$X_{k+1} = X_k - h \nabla f(X_k) + \sqrt{2h} \xi_k,$$

where  $h > 0$  is step-size and  $\xi_k \stackrel{\text{i.i.d.}}{\sim} N(0, I_d)$ . Replacing the  $-\nabla f(\cdot)$  with estimated score leads to the following.

## Score-based Langevin Monte Carlo Algorithm

$$X_{k+1} = X_k + h s_\theta(X_k) + \sqrt{2h} \xi_k,$$

where  $s_\theta(x)$  is an estimate of score function  $\nabla \log \mu(x) = -\nabla f(x)$ .

As long as the score estimate is good ( $s_\theta \approx \nabla \log \mu(\cdot)$ ) and  $h$  is small, we expect that the law of  $\mu_K$  to be close to  $\mu$  for  $K$  large enough.

# Main Challenges

- Estimated score function is inaccurate in low density regions.

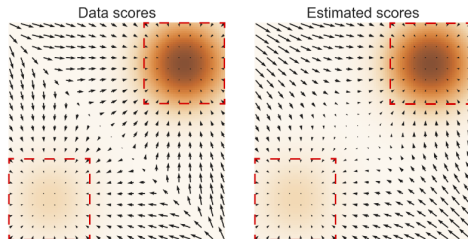


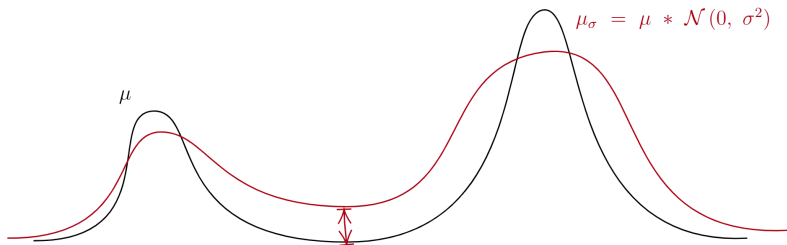
Figure 2: **Left:**  $\nabla_{\mathbf{x}} \log p_{\text{data}}(\mathbf{x})$ ; **Right:**  $\mathbf{s}_{\theta}(\mathbf{x})$ . The data density  $p_{\text{data}}(\mathbf{x})$  is encoded using an orange colormap: darker color implies higher density. Red rectangles highlight regions where  $\nabla_{\mathbf{x}} \log p_{\text{data}}(\mathbf{x}) \approx \mathbf{s}_{\theta}(\mathbf{x})$ .

- Langevin algorithms mix slowly when faced a bottleneck.

# Annealed Langevin Algorithm

A key observation here is that once we smooth  $\mu$  by convolving it with large Gaussian noise,  $\mu_\sigma := \mu * \mathcal{N}(0, \sigma^2 I_d)$ , then both problems are gone for the problem of sampling  $\mu_\sigma$ . However, to sample from  $\mu$ , it is not enough to say that we can sample from  $\mu_\sigma$  with large enough noise  $\sigma$ . [Song and Ermon, 2020] proposed to improve Langevin algorithms with estimated score by

1. perturbing the data using various levels of noise
2. estimating score and run Langevin algorithms, at all noise levels.





# Annealed Langevin Algorithm

---

Inspired by the idea of annealing, the annealed Langevin algorithm is proposed. Set  $L$  levels of noise from large to small,  $\sigma_1, \dots, \sigma_L$ , and  $h_L > 0$

- For  $i$  from 1 to  $L$  run
  - Set step-size  $h_i = h_L \sigma_i^2 / \sigma_L^2$ .
  - Run unadjusted Langevin algorithm for  $K$  steps

$$X_{k+1} = X_k + h_i s_\theta(X_k, \sigma_i) + \sqrt{2h_i} Z_k,$$

where  $Z_k$  is independent standard Gaussian noise, and  $s_\theta(X_k, \sigma_i)$  is the estimated score function for  $\mu * N(0, \sigma_i^2 I_d)$ .

## Questions:

- How to set noise levels  $\sigma_1, \dots, \sigma_L$  in practice?
- Does the annealed Langevin algorithm converges to the target measure as our intuition suggest?

# Outline

---

1. Two Settings of Sampling with Applications
2. Density Estimation
3. Generative Model
4. Score Matching
- 5. Diffusion Model**

# Diffusion Model: Forward Process

## Forward Process

The forward process is specified via a stochastic differential equation (SDE).

$$dX_t = -X_t dt + \sqrt{2} dB_t, \quad X_0 \sim q_0 := \mu, \quad (\text{forward process})$$

where  $(B_t)_{t \geq 0}$  is a standard Brownian motion in  $\mathbb{R}^d$ .

- The solution of this SDE is also called the Ornstein-Uhlenbeck (OU) process.

(1) The solution is

$$X_t = e^{-t} X_0 + \sqrt{2} \int_0^t e^{-(t-s)} dB_s.$$

(2)  $\mathbb{E} X_t = e^{-t} \mathbb{E} X_0$ ,  $\text{Var}(X_t) = 1 - e^{-2t}$ , hence we can write

$$X_t = e^{-t} X_0 + (1 - e^{-2t})^{1/2} Z_t, \text{ where } Z_t \sim N(0, I_d).$$

(3)  $N(0, I_d)$  is an invariant measure of OU process.

- In practice, one may consider the time-rescaled OU process :

$$d\bar{X}_t = -g(t)^2 \bar{X}_t dt + \sqrt{2} g(t) dB_t, \text{ with a positive smooth function } g : \mathbb{R}_+ \rightarrow \mathbb{R}_+.$$

## Diffusion Model: Reverse Process

---

In general, suppose we have an SDE of the form

$$dX_t = a(X_t, t)dt + b_t dB_t.$$

Under mild conditions on the process, the process can be reversed, and the reverse process also admits an SDE description. Fix terminal time  $T > 0$ , define the reverse process

$$X_t^{\leftarrow} := X_{T-t}, \quad \text{for } t \in [0, T],$$

then the process  $(X_t^{\leftarrow})_{t \in [0, T]}$  satisfies the following reverse SDE

$$dX_t^{\leftarrow} = a^{\leftarrow}(X_t^{\leftarrow}, t)dt + b_{T-t}dW_t, \quad X_0^{\leftarrow} \sim q_T,$$

where  $W_t$  is the reversed Brownian motion, for simplicity, we don't distinguish  $B_t$  and  $W_t$ . We need to choose the reverse drift  $a^{\leftarrow}(x, T - t)$  such that

$$\text{Law}(X_t^{\leftarrow}) = \text{Law}(X_{T-t})$$

By **Fokker-Planck equation**, we can choose

$$a(x, t) + a^{\leftarrow}(x, T - t) = b_t b_t^{\top} \nabla \log q_t, \quad \text{where } q_t := \text{law}(X_t).$$

# Diffusion Model: Reverse Process and Score Matching

---

Applying the result to the forward process, we obtain the reverse process in DDPM

## Reverse Process

$$dX_t^{\leftarrow} = [X_t^{\leftarrow} + 2\nabla \log q_{T-t}(X_t^{\leftarrow})] dt + \sqrt{2} dB_t, \quad X_0^{\leftarrow} \sim q_T, \quad (\text{reverse process})$$

where  $(B_t)_{t \in [0, T]}$  is the reversed Brownian motion.

Since  $q_0$  is not explicitly known and is only known via its samples  $x_1, \dots, x_N$ , in order to implement the reverse process, we need to estimate the score function of  $q_t$  at any time  $t \in [0, T]$  via the samples.

## Diffusion Model: Score Matching

---

By the properties of OU process, we know for any given  $t$ ,  $X_t$  can be written as a linear combination of  $X_0$  and independent noise

$$X_t = e^{-t}X_0 + (1 - e^{-2t})^{1/2}Z_t, \quad Z_t \sim N(0, I_d).$$

By the theorem of Donising Score Matching, the score matching problem for  $X_t$

$$\min_{s_t \in \mathcal{F}} \mathbb{E} \left\| \nabla \log q_t(X_t) - s_t(X_t) \right\|_2^2$$

is equivalent to

$$\min_{s_t \in \mathcal{F}} \mathbb{E} \left[ \left\| s_t(X_t) + \frac{1}{\sqrt{1 - e^{-2t}}} Z_t \right\|_2^2 \right],$$

where  $\mathcal{F}$  could be, e.g., a class of neural networks.

## Diffusion Model: Score Matching

---

The objective can be replaced with an empirical version and estimated on the basis of samples  $x_0^{(1)}, \dots, x_0^{(N)}$  from  $q_0$ , leading to the finite-sample problem

$$\min_{s_t \in \mathcal{F}} \frac{1}{N} \sum_{i=1}^N \left\| s_t(x_t^{(i)}) + \frac{1}{\sqrt{1 - \exp(-2t)}} z_t^{(i)} \right\|^2, \quad (\text{DDPM-SM})$$

where  $(z_t^{(i)})_{i \in [N]}$  are i.i.d. standard Gaussian samples independent of the data  $(x_0^{(i)})_{i \in [N]}$ .

Hence, we can learn the score of  $q_t$  for all  $t \in [0, T]$ , and we assume

$$\mathbb{E}[\|s_t(X_t) - \nabla \log q_t(X_t)\|_2^2] \leq \varepsilon_{\text{score}}^2.$$

# Diffusion Model

---

## Forward process:

$$dX_t = -X_t dt + \sqrt{2} dB_t, \quad X_0 \sim q_0 := \mu$$

## Reverse process:

$$dX_t^{\leftarrow} = [X_t^{\leftarrow} + 2\nabla \log q_{T-t}(X_t^{\leftarrow})] dt + \sqrt{2} dB_t, \quad X_0^{\leftarrow} \sim q_T \approx \gamma^d := \text{law of } N(0, I_d),$$

**Partition** : Partition the interval  $[0, T]$  to  $[kh, (k+1)h]$ ,  $k = 0, 1, \dots, K-1$  with  $h > 0$ ,  $K = T/h$ . Integrate the reverse process from  $[kh, t]$ ,  $t \in [kh, (k+1)h]$ ,

$$X_t^{\leftarrow} = X_{kh}^{\leftarrow} + \int_{kh}^t X_s^{\leftarrow} ds + \int_{kh}^t \underbrace{2 \nabla q_{T-s}(X_s^{\leftarrow})}_{\approx s_{T-kh}(X_{kh}^{\leftarrow})} ds + \sqrt{2}(B_t - B_{kh}).$$

## Score matching and integral approximation:

$$dX_t^{\leftarrow} = \{ X_t^{\leftarrow} + 2 s_{T-kh}(X_{kh}^{\leftarrow}) \} dt + \sqrt{2} dB_t, \quad t \in [kh, (k+1)h],$$

which is a linear SDE and can be integrated in closed form.



# Diffusion Model: Convergence Analysis

Let  $p_t := \text{Law}(X_t^{\leftarrow})$ , DDPM has mainly three types of errors.

1. The error made at initialization of reverse process,  $\gamma^d$  used instead of  $q_T$ .
2. The score matching error, which on the sample size  $N$ , the size of the function class  $\mathcal{F}$  and its closeness to the true score function.
3. The discretization of the reverse process, which depends on the step-size  $h$ .

## Assumption 1 (Lipschitz score)

For any  $t \geq 0$ , the score  $\nabla \log q_t$  is  $L$ -Lipschitz.

## Assumption 2 (Second moment bound)

Assume that  $M_2^2 := \mathbb{E}_{X \sim q_0} \|X\|_2^2 < \infty$ .

## Assumption 3 (Score estimation error bound)

For  $k = 1, \dots, K$ ,

$$\mathbb{E}_{q_{kh}} \|s_{kh} - \nabla \log q_{kh}\|_2^2 \leq \epsilon_{\text{score}}^2.$$

# Diffusion Model: Convergence Analysis

## Theorem ([Chen et al., 2023])

Under the three previous assumptions. Let  $p_T$  be the output of the DDPM algorithm at time  $T > 0$ , with  $h = T/K$  and  $K$  the number of steps, suppose  $h \lesssim 1/L$ , then

$$d_{\text{TV}}(p_T, q_0) \lesssim \underbrace{\sqrt{\text{KL}(q_0 \parallel \gamma^d)} e^{-T}}_{\text{convergence of forward process}} + \underbrace{(L\sqrt{d}h + LM_2h)\sqrt{T}}_{\text{discretization error}} + \underbrace{\epsilon_{\text{score}}\sqrt{T}}_{\text{score estimation error}}.$$

### Remark:

1. Unlike Langevin algorithms, this theorem does not assume any type of “bottleneck” condition such as log concave target distribution. **It means that DDPM can efficiently sample from multi-modal target measures as long as the score estimation is good.**
2. Even though the KL divergence term  $\text{KL}(q \parallel \gamma^d)$  might be large (even exponentially in dimension  $d$ ), the contraction of the forward process creates a  $\exp(-T)$  term which can make the first term small.

# References I

---



Chen, S., Chewi, S., Li, J., Li, Y., Salim, A., and Zhang, A. R. (2023).

Sampling is as easy as learning the score: theory for diffusion models with minimal data assumptions.



Dyer, M. E. and Frieze, A. M. (1988).

On the complexity of computing the volume of a polyhedron.

*SIAM Journal on Computing*, 17(5):967–974.



Jia, H., Laddha, A., Lee, Y. T., and Vempala, S. (2021).

Reducing isotropy and volume to kls: an  $\tilde{O}(n^3 \psi^2)$  volume algorithm.

In *Proceedings of the 53rd Annual ACM SIGACT Symposium on Theory of Computing*, pages 961–974.



Scott, D. W. (2015).

*Multivariate density estimation: theory, practice, and visualization*.

John Wiley & Sons.

# References II

---



Song, Y. and Ermon, S. (2019).

Generative modeling by estimating gradients of the data distribution.

*Advances in neural information processing systems*, 32.



Song, Y. and Ermon, S. (2020).

Generative modeling by estimating gradients of the data distribution.



Tsybakov, A. (2009).

Introduction to nonparametric estimation. springer series in statistics. springer, new york.

*Thanks!*