

I. INTRODUCTION

In this blogpost, we present a 'ground up' view of Diffusion Models, starting from its core element (i.e. the 'score'), building up to the modern form. While other models use a direct parametric function (with parameters θ) to transform noise into data distribution $q_{data}(x)$, i.e. $x = f_{\theta}(z)$, where $z \sim$ $\mathcal{N}(0, I)$, diffusion model do so iteratively that involves a parametric "periteration" function (estimate of the true 'score')

 $x = g_1(g_2(g_3(...z_{\dots}, s_{\theta}), s_{\theta}), s_{\theta}), \text{ where } z \sim \mathcal{I}$ The 'Score' of a distribution:

The 'score' of a distribution $q_{data}(x)$ is simply the gradient of the log-density, i.e. $\nabla_x \log q_{\text{data}}(x)$. This term was originally coined [1] long back in 1935 by Ronald Fisher in a slightly different context. But in machine learning, it is interpreted as a *guide* to go uphill in the log-density surface. An infinitesimal step in the direction of the score can get us to a state of higher likelihood $x'_t = x_t + \delta \cdot \nabla_x \log q_{\text{data}}(x)$

II. GENERATIVE MODELLING WITH SCORE FUNCTION

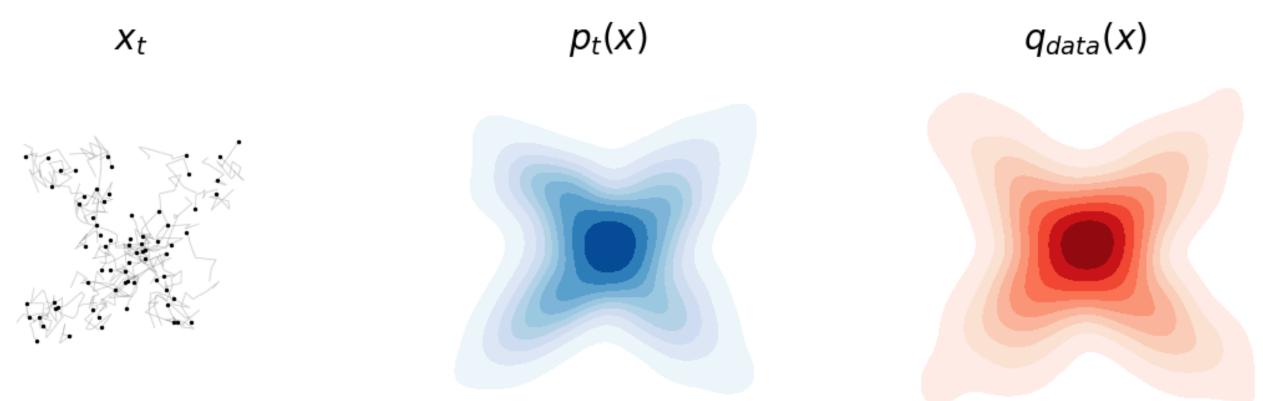
One can craft an iterative "sampling rule" solely based on the intutive interpretation of score provided in Equation 2

 $p_t(x)$

 $dx =
abla_x \log q_{ ext{data}}(x) \cdot dt$

However, this process is NOT guarenteed to converge to the true distribution $q_{\text{data}}(x)$. Turns out that this problem has been studied [2] in particle physics long ago by Paul Langevin, in order to explain movement of particles suspended in fluid. According to their theory, adding a little noise term fixes it.

 $dx = \nabla_x \log q_{\text{data}}(x) \cdot dt + \sqrt{2} \cdot dB_t$, where $dB_t = \mathcal{N}(0, dt)$ (4)



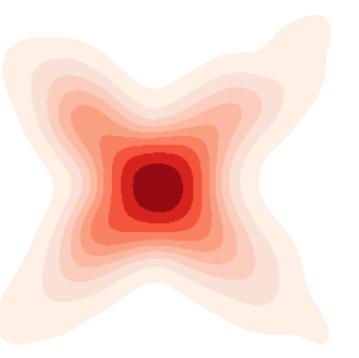
BUILDING DIFFUSION MODEL'S THEORY FROM GROUND UP

$$\mathcal{N}(0, I).$$
 (1)

(2)

(3)

 $q_{data}(x)$



Fokker-Planck Equation & a probability path:

The process in Equation 4 has a guarantee of convergence as $t \to \infty$. This can be validated by first noting $\mu_t(x) = \nabla_x \log q_{\text{data}}(x), \sigma_t(x) = \sqrt{2}$ and using the Fokker-Planck equation where we set $p_{\infty}(x) := q_{\text{data}}(x)$

$$\frac{\partial}{\partial t}p_{\infty}(x) = -\frac{\partial}{\partial x}p_{\infty}(x)\mu_{t}(x) + \frac{1}{2}\frac{\partial^{2}}{\partial x^{2}}p_{\infty}(x)\sigma_{t}^{2}(x)$$
(5)

$$p_t \mid p_0 = \mathcal{N}(0, I)$$

 $p_t \mid p_0 = \mathcal{N}(0, I)$

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The forward process & 'schedule':

It was argued [3] that learning an estimate of the true score everywhere on the data space x is extremely hard. The most popular solution to this problem is to learn a score specialized for t. To do so, one need samples from $p_t(x)$. The "forward process" is thus designed as an *ahead-of-time* description of the "path" taken by $p_t(x)$ on the probability space (refer to the above figure). One can revert the path by using the same Langevin Equation 4 but with end target being $\mathcal{N}(0, I)$ and starting from $q_{\text{data}}(x)$ $dx = \nabla_x \log \mathcal{N}(0, I) \cdot dt + \sqrt{2} \cdot dB$

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To see its similarity to the 'modern' form of forward process, one must redefine the interpretation of time to contain it within a finite interval (e.g. [0, 1]). This is required due to the fact that langevin equation only guarantees convergence with $t \to \infty$. One can do so by first discretizing the above equation and plugging a new time mapping $t' = \mathcal{T}(t) = 1 - \exp(-t)$ $x_{t'+dt'} = (1 - e^t dt') x_t$

This resembles DDPM's [4] forward process where $e^t dt'$ is analogous to β_t in DDPM (small and increating in time t). We can then sample x_t for any t by simulating Equation 7.

 $x_t \sim q_t(x)$

Now the score of a *t*-specialized density, i.e.

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 $p_t \mid p_0 = \mathcal{U}(-2, 2)$ $p_t \mid p_0 = \mathcal{U}(-2, 2)$

$$B_t = -x \cdot dt + \sqrt{2} \cdot dB_t \qquad (6)$$

$$+\sqrt{2e^t}dB_t \tag{7}$$

$$(8)$$

$$\nabla_x \log q_t(x) \text{ must be learned.}$$

The previous section deals with the generative modelling problem – given that we have access to the true score $\nabla_x \log q_{\text{data}}(x)$, which in reality, we don't. The very first credible solution to the score learning was proposed by [5], which turns the original Score Matching objective (not computable)

$$J(\theta) = \frac{1}{2} \mathbb{E}_{x \sim q_{\text{data}}(x)} \left[\parallel s_{\theta}(x) - \nabla_x \log q_{\text{data}}(x) \parallel^2 \right]$$
(9)

(ISM)" which does not require the true score (unavailable)

$$J_I(\theta) = \mathbb{E}_{x \sim q_{\text{data}}(x)} \left[\frac{1}{2} \| s_{\theta}(x) \|^2 + \operatorname{Tr}(\nabla_x s_{\theta}(x)) \right].$$
(10)

This objective was further upgraded by Vincent Pascal [6] as the "Denoising" Score Matching", the variant still used in modern Diffusion Models

$$J_{D}(\theta) = \mathbb{E}_{x \sim q_{\text{data}}(x), \epsilon \sim \mathcal{N}(0, I)} \left[\frac{1}{2} \| s_{\theta} \left(\underbrace{x + \sigma \epsilon}_{\tilde{x}} \right) - \left(-\frac{\epsilon}{\sigma} \right) \|^{2} \right]. \quad (11)$$

A slight modification (reparameterization) of the denoising score matching loss above leads to the widely used variant called "noise estimation" where instead of score, we learn the noise direction from a noisy sample

$$J_{\epsilon}(\theta) = \mathbb{E}_{x \sim q_{\text{data}}(x), \epsilon \sim \mathcal{N}(0, I)} \left[\frac{1}{2\sigma^2} \| \epsilon_{\theta}(\tilde{x}) - \epsilon \|^2 \right]$$
(12)

Yet another variant, named as "end point estimation" can be derived easily from the above equation, which predicts the clean sample from noisy one

$$J_{x}(\theta) = \mathbb{E}_{x \sim q_{\text{data}}(x), \epsilon \sim \mathcal{N}(0, I)} \left[\frac{1}{2\sigma^{4}} \| x_{\theta}(\tilde{x}) - x \|^{2} \right]$$
(13)

Connection to Tweedie's formula:

Equation 13 above has an interesting interpretation – as long as the noise is gaussian, it can be seen as learning posterior mean of clean quantity from noisy samples. In bayesian "inverse problem" literature, this is known as *Tweedie's formula* [7]

 $\mathbb{E}_{x \sim q(x \mid \tilde{x})}$

The similarity is obvious when

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III. ESTIMATING THE SCORE FUNCTION

.. to a practically computable version named "Implicit Score Matching

$$[x] = \tilde{x} + \sigma^2 \nabla_{\tilde{x}} \log p(\tilde{x}).$$
 (14)
In we observe the fact that $x_{\theta}(\tilde{x}) = \tilde{x} + \sigma^2 s_{\theta}(\tilde{x}).$

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[1] R. A. Fisher, "The detection of linkage with "dominant" abnormalities," *Annals of Eugenics*, vol. 6, 1935.

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