

Controllable Generative AI for Biological Sequences

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Outline

Mathematical Background

Continuous Diffusion Models

The Idealism-Realism Tradeoff

Guidance Methods

Discrete Diffusion

Guidance for Discrete Diffusion

Plus a bonus lecture on Applications to Biological Sequences.

Organisation



Ewa Szczurek (Lecture)



Jan Olszewski (Tutorial)

- **Website:** <https://phdopen.mimuw.edu.pl/index.php?page=126w4>
- **Slack:** https://join.slack.com/t/phdopen2026di-66v4471/shared_invite/zt-4180w6k1u-~JOA1Uryl95PL7XzREI6qA
- **Lecture:** Slides accompanied by lecture notes,
- **Tutorial:** python notebooks, **bring your laptops**
- **Plan for tomorrow:** room 5440, 14:15 - 15:45 Lecture, 16:15 - 17:30 Tutorial

Continuous random variables and densities

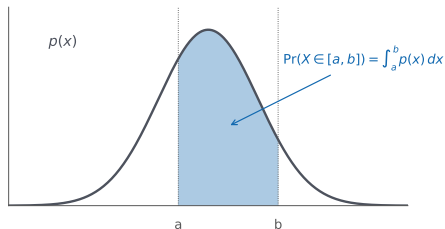
A continuous r.v. X in $\mathbb{R}^d$ has a density $p : \mathbb{R}^d \rightarrow \mathbb{R}_{\geq 0}$ with $\int p(x) dx = 1$.

$p(x)$ is not a probability

Densities can exceed one. Probabilities come from integration:

$$\Pr(X \in A) = \int_A p(x) dx.$$

Think of $p(x)$ telling **how likely** the sample x is.
Crucial for generative models.



Useful facts about continuous random variables and densities

For two r.v.s X, Y , we will think of

- x as a **sample**, with **likelihood** $p(x)$, learned by the **generative model**
- y as a label, with **conditional likelihood** $p(y|x)$, learned by **classifier**.

The following holds for the joint $p(x, y)$:

- **Marginal:** $p(x) = \int p(x, y) dy$
- **Conditional:** $p(x | y) = p(x, y)/p(y)$ where $p(y) > 0$
- **Product rule:** $p(x, y) = p(y|x)p(x)$.

Bayes' rule for densities

The identity from the discrete setting carries over to densities:

$$p(x | y) = \frac{p(y | x) p(x)}{p(y)}$$

Taking log on both sides and differentiating w.r.t. x :

$$\nabla_x \log p(x | y) = \nabla_x \log p(y | x) + \nabla_x \log p(x),$$

since $\nabla_x \log p(y) = 0$ (the marginal $p(y)$ does not depend on x).

Why this matters

The intractable normalizer $p(y) = \int p(y | x) p(x) dx$ **drops out** of the gradient.

Expectations

For any function f :

$$\mathbb{E}[f(X)] = \int f(x) p(x) dx, \quad \mathbb{E}[f(X) \mid Y = y] = \int f(x) p(x \mid y) dx.$$

Practical consequence

All training losses in this course will be expectations of squared errors or cross-entropies, approximated by Monte Carlo over training batches.

Gaussian algebra

The Gaussian density on $\mathbb{R}^d$ with mean μ and covariance Σ :

$$\mathcal{N}(x; \mu, \Sigma) = \frac{1}{(2\pi)^{d/2} |\Sigma|^{1/2}} \exp\left(-\frac{1}{2}(x - \mu)^\top \Sigma^{-1}(x - \mu)\right).$$

Fact 1: Linear transformation

If $X \sim \mathcal{N}(\mu, \Sigma)$ and $a \in \mathbb{R}$, $b \in \mathbb{R}^d$ are constants:

$$aX + b \sim \mathcal{N}(a\mu + b, a^2\Sigma).$$

Fact 2: Sum of independent Gaussians

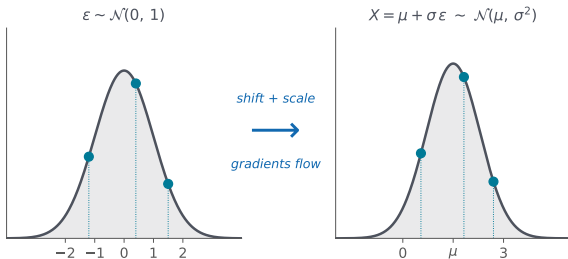
If $X \sim \mathcal{N}(\mu_1, \Sigma_1)$ and $Y \sim \mathcal{N}(\mu_2, \Sigma_2)$ are independent:

$$X + Y \sim \mathcal{N}(\mu_1 + \mu_2, \Sigma_1 + \Sigma_2).$$

Fact 3: The reparameterization trick

Reparameterization

$$X \sim \mathcal{N}(\mu, \sigma^2 I) \iff X = \mu + \sigma \epsilon, \quad \epsilon \sim \mathcal{N}(0, I).$$



Two payoffs:

- **Randomness outside:** X is a deterministic function of (μ, σ, ϵ) .
- **Gradients flow** through it, enables backprop.

KL divergence

To train a probabilistic model we need to measure closeness of distributions.

$$\text{KL}(q \parallel p) = \int q(x) \log \frac{q(x)}{p(x)} dx = \mathbb{E}_{X \sim q} \left[\log \frac{q(X)}{p(X)} \right].$$

- **Non-negative:** $\text{KL}(q \parallel p) \geq 0$, with equality iff $q = p$ a.e.
- **Asymmetric:** a divergence, not a metric.
- **Loss-like:** minimizing $\text{KL}(q \parallel p_\theta)$ in θ (with $q = \text{data}$) recovers **maximum likelihood**.

The Gaussian case (same covariance)

For $p = \mathcal{N}(\mu_1, \sigma^2 I)$ and $q = \mathcal{N}(\mu_2, \sigma^2 I)$:

$$\text{KL}(p \parallel q) = \frac{1}{2\sigma^2} \|\mu_1 - \mu_2\|^2 \longrightarrow \text{scaled squared error between means.}$$

The score function

For a density $p(x)$ on $\mathbb{R}^d$, the *score* is the gradient of the log-density:

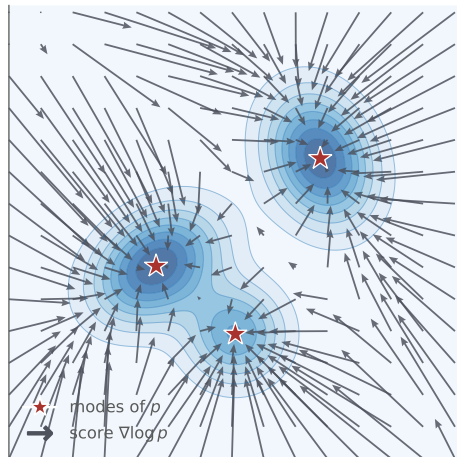
$$s(x) := \nabla_x \log p(x) \in \mathbb{R}^d.$$

Geometric meaning

$s(x)$ points in the direction of **steepest increase** of $\log p$.

- Following s moves samples *toward modes*.
- Score-based samplers follow this gradient, with added noise to avoid mode collapse.

$\nabla_x \log p(x)$ points toward the modes



The score: invariance to normalization

Suppose we only know p up to a constant: $p(x) = \tilde{p}(x)/Z$ with intractable $Z = \int \tilde{p}(x) dx$. Then

$$\nabla_x \log p(x) = \nabla_x (\log \tilde{p}(x) - \log Z) = \nabla_x \log \tilde{p}(x),$$

because Z is constant in x .

Huge practical advantage: we can work with unnormalized distributions and still compute scores exactly.

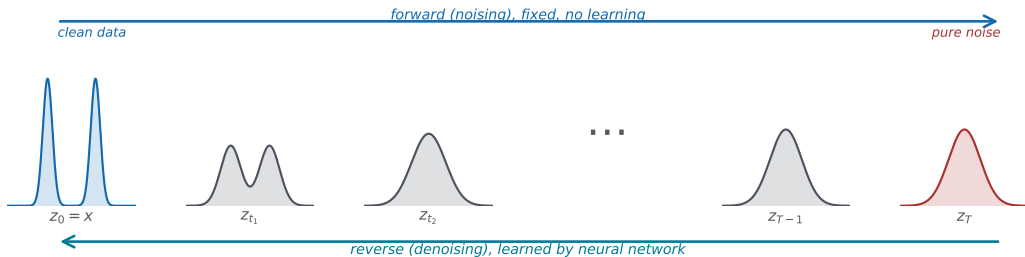
Why this matters for Bayesian guidance

Bayes gives $p(x | y) = p(y | x) p(x) / p(y)$ with intractable $p(y) = \int p(y | x) p(x) dx$. Taking the score wipes it out:

$$\nabla_x \log p(x | y) = \nabla_x \log p(y | x) + \nabla_x \log p(x).$$

Both right-hand-side terms are tractable: a classifier's gradient and the data score.

Diffusion in one picture



Forward: add Gaussian noise, T steps.

Fixed. No learning.

Reverse: undo the noise.

Learned by a neural net.

Forward process¹

A Markov chain that gradually corrupts x over T steps:

$$q(z_t | z_{t-1}) = \mathcal{N}\left(z_t; \sqrt{1 - \beta_t} z_{t-1}, \beta_t I\right), \quad z_0 := x.$$

- $\{\beta_t\}$: **variance schedule**, small per step (linear or cosine).
- Shrink by $\sqrt{1 - \beta_t}$, add fresh noise of variance β_t :

$$\text{Var}(z_t) = (1 - \beta_t) \text{Var}(z_{t-1}) + \beta_t \xrightarrow{\text{Var}(z_{t-1})=1} \text{Var}(z_t) = 1.$$

Coefficients sum to one, so variance stays 1 (does not explode).

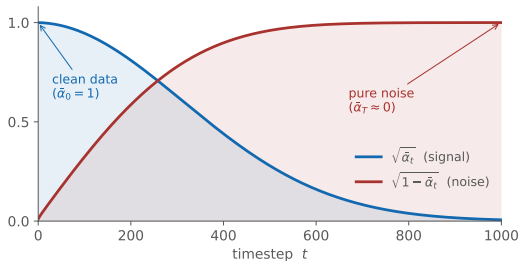
- Typical $T = 1000$: **many small steps** make each denoising decision easy.

¹Ho et al. (2020) *Denoising Diffusion Probabilistic Models* (DDPM)

Closed-form marginal: the workhorse equation

Define $\alpha_t := 1 - \beta_t$, $\bar{\alpha}_t := \prod_{s=1}^t \alpha_s$. Iterating the linear-Gaussian step gives:

$$z_t = \sqrt{\bar{\alpha}_t} x + \sqrt{1 - \bar{\alpha}_t} \epsilon, \quad \epsilon \sim \mathcal{N}(0, I).$$



- Jump to any noise level in one step, no chain simulation.
- $\sqrt{\bar{\alpha}_t}$: signal kept.
 $\sqrt{1 - \bar{\alpha}_t}$: noise injected.
- $\bar{\alpha}_0 = 1$, $\bar{\alpha}_T \approx 0$, which justifies starting reverse from $\mathcal{N}(0, I)$.

Reverse process is intractable

We want $q(z_{t-1} \mid z_t)$.

From the Bayes rule

$$q(z_{t-1} \mid z_t) = \frac{q(z_t \mid z_{t-1}) q(z_{t-1})}{q(z_t)}.$$

This is intractable

The marginals $q(z_{t-1})$ and $q(z_t)$ involve integrating over the entire data distribution.

Reverse process: the key trick

But if we additionally condition on $x \dots$

$q(z_{t-1})$ and $q(z_t)$ replaced by the **closed-form Gaussians** $q(z_{t-1} | x)$ and $q(z_t | x)$, and

$$q(z_{t-1} | z_t, x) = \mathcal{N}(z_{t-1}; \tilde{\mu}_t(z_t, x), \tilde{\beta}_t I)$$

is also a **closed-form Gaussian**, with mean

$$\tilde{\mu}_t(z_t, x) = \frac{\sqrt{\bar{\alpha}_{t-1}} \beta_t}{1 - \bar{\alpha}_t} x + \frac{\sqrt{\alpha_t} (1 - \bar{\alpha}_{t-1})}{1 - \bar{\alpha}_t} z_t.$$

and variance

$$\tilde{\beta}_t = \frac{1 - \bar{\alpha}_{t-1}}{1 - \bar{\alpha}_t} \beta_t \leq \beta_t \quad (\text{conditioning on } x \text{ shrinks the variance}).$$

The posterior mean: a convex combination

The coefficients are nonnegative and approx. sum to one (exactly so as $\beta_t \rightarrow 0$), making $\tilde{\mu}_t$ a convex combination of x and z_t .

$$\tilde{\mu}_t(z_t, x) = \frac{\sqrt{\bar{\alpha}_{t-1}} \beta_t}{1 - \bar{\alpha}_t} x + \frac{\sqrt{\alpha_t} (1 - \bar{\alpha}_{t-1})}{1 - \bar{\alpha}_t} z_t.$$

$\tilde{\mu}_t(z_t, x)$ is a convex combination of x and z_t

x (clean)

z_t (noisy)



low t
(close to x)

mid t

high t
(close to z_t)

teal squares = $\tilde{\mu}_t$ (the reverse posterior mean)

Training: predict x from z_t

At sampling time we don't have x . So: predict it.

Train a network $\hat{x}_\theta(z_t, t)$ to recover the clean data from the noisy state:

$$\mathcal{L}(\theta) = \mathbb{E}_{x, t, \epsilon} \left[\|\hat{x}_\theta(z_t, t) - x\|^2 \right], \quad z_t = \sqrt{\bar{\alpha}_t} x + \sqrt{1 - \bar{\alpha}_t} \epsilon.$$

$\mathcal{L}(\theta)$ follows from the variational bound (ELBO) on $\log p_\theta(x)$, after the KL terms collapse to squared errors.

Recipe per training example

Sample $(x, t, \epsilon) \rightarrow$ form $z_t \rightarrow$ predict $\hat{x}_\theta(z_t, t) \rightarrow$ MSE loss.

Sampling: DDPM ancestral algorithm

Algorithm

1. Sample $z_T \sim \mathcal{N}(0, I)$.
2. For $t = T, T-1, \dots, 1$:
 - ▶ Compute $\hat{x}_\theta(z_t, t)$ (one forward pass)
 - ▶ Compute posterior mean $\mu_\theta = \tilde{\mu}_t(z_t, \hat{x}_\theta(z_t, t))$
 - ▶ Sample $z_{t-1} \sim \mathcal{N}(\mu_\theta, \tilde{\beta}_t I)$
3. Return z_0 .

Cost: T forward passes per sample. For $T = 1000$, $\sim 1000\times$ slower than a VAE/GAN.

Each reverse step is a **small** correction. Many small denoising decisions beat one big leap from noise to x .

Three equivalent parameterizations

The trained network can be written in three forms, algebraically interchangeable:

Form	Predicts	Conversion
$\hat{x}_\theta(z_t, t)$	clean data x (Tweedie estimate)	–
$\hat{\epsilon}_\theta(z_t, t)$	the noise ϵ	$\hat{\epsilon}_\theta = \frac{z_t - \sqrt{\bar{\alpha}_t} \hat{x}_\theta}{\sqrt{1 - \bar{\alpha}_t}}$
$s_\theta(z_t, t)$	score $\nabla_{z_t} \log p_t$	$s_\theta = -\frac{\hat{\epsilon}_\theta}{\sqrt{1 - \bar{\alpha}_t}}$

- $\hat{x} \leftrightarrow \hat{\epsilon}$: pure algebra from the forward marginal $z_t = \sqrt{\bar{\alpha}_t} x + \sqrt{1 - \bar{\alpha}_t} \epsilon$.
- $\hat{\epsilon} \rightarrow s_\theta$: Tweedie's formula, the score of the noised marginal is the rescaled noise, $\nabla_{z_t} \log p_t = -\hat{\epsilon}_\theta / \sqrt{1 - \bar{\alpha}_t}$.

A remark: the continuous-time view

Everything so far was discrete time, $t = 1, \dots, T$. Letting the steps shrink ($\beta_t \rightarrow 0$) opens a second formulation. We only sketch it here.

- **Reverse SDE.** The small-step limit of the forward chain is a stochastic differential equation; **DDPM is one discretization of it.**
- **Probability-flow ODE.** A *deterministic* process (no injected noise) with the *same marginals* p_t , hence the same endpoint p_{data} . Closely related to **flow matching**.
- **Same trained network** drives both: no retraining, only the sampling changes.

Why it matters

- Faster sampling: **DDIM**² ~ 50 steps, **DPM-Solver**³ $\sim 10\text{--}20$ steps, vs. DDPM's ~ 1000 .
- The deterministic ODE enables **exact likelihoods**.

²Song et al. (2021)

³Lu et al. (2022)

The idealism–realism tradeoff

We have two competing needs:

Realism

Modeled by the **prior** $p_{\theta}(x) \approx p(x)$ gives plausible data: *any* peptide, molecule, image.

Idealism

Modeled by a **reward** $r(x) \geq 0$ scores how *desirable* an output is (binding affinity, antimicrobial activity, prompt match).

The question

How do we modify a generative model to favour high-reward outputs while keeping samples close to the data distribution?

KL-regularized optimization

Naive: maximize the reward.

This is degenerate

Returns a delta on the highest-reward point, which may not even be realistic (r is reliable only on the data manifold).

Fix: regularize π toward the prior with a KL penalty:

$$\pi^* = \arg \max_{\pi} \left[\mathbb{E}_{x \sim \pi} [r(x)] - \beta \text{KL}(\pi \parallel p_{\theta}) \right]$$

$\beta > 0$ trades off reward against proximity to the prior:

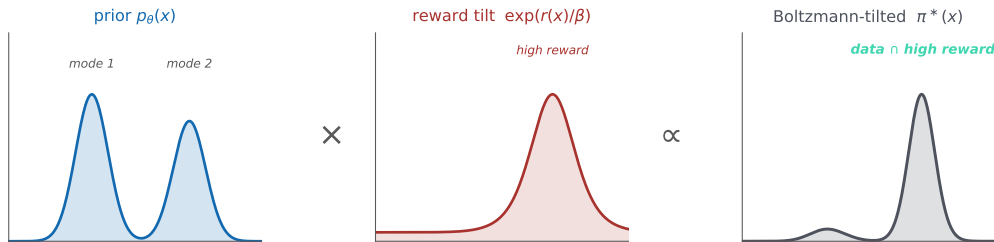
- **Small β :** aggressive; risks off-manifold samples, reward hacking.
- **Large β :** stay close to prior; ignore the reward.

Same form as KL-regularized RL. In physics: inverse temperature.

The Boltzmann-tilted target

The KL-regularized objective has a closed-form optimum

$$\pi^*(x) = \frac{1}{Z} p_\theta(x) \cdot \exp(r(x)/\beta), \quad Z = \int p_\theta(x) \exp(r(x)/\beta) dx.$$



Both have a vote: a sequence with high r but low p_θ is suppressed by the prior; a sequence with high p_θ but low r is suppressed by the reward.

Two special cases of the same framework

Bayesian posterior

Take $r(x) = \log p(y | x)$ and $\beta = 1$:

$$\pi^*(x) \propto p_\theta(x) p(y | x) = p_\theta(x | y).$$

The Boltzmann target *is* the Bayesian posterior.

Classifier guidance exemplifies this case.

Hard constraints

Take $r(x) = \log \mathbf{1}[x \in S]$ (0 on S , $-\infty$ off it), so $\exp(r(x)/\beta) = \mathbf{1}[x \in S]$:

$$\pi^* \propto p_\theta(x) \mathbf{1}[x \in S] \text{ concentrates on } S \cap \text{supp}(p_\theta).$$

Realistic outputs satisfying a hard constraint.

The obstacle and the score trick

The partition function tractability obstacle

The partition function $Z = \int p_{\theta}(x) \exp(r(x)/\beta) dx$ is **intractable** in high dimensions. We know π^* only up to the unknown Z . Cannot evaluate or directly sample.

The score saves us

Take $\nabla_x \log$ on both sides. **Z disappears:**

$$\nabla_x \log \pi^*(x) = \nabla_x \log p_{\theta}(x) + \frac{1}{\beta} \nabla_x r(x)$$

Additive decomposition: prior score + scaled reward gradient.
Both pieces are tractable. **Used by some guidance methods.**

Time-dependent extension: the value function u_t

Diffusion gives a *family* p_t for $t \in [0, T]$.

We need a family of tilted targets:

$$q_t^*(z_t) \propto p_t(z_t) u_t(z_t), \quad u_t(z_t) := \mathbb{E}_P[\exp(r(Z_0)/\beta) \mid Z_t = z_t].$$

u_t asks: *starting from z_t now, what is the expected exponentiated reward if the prior dynamics run to completion?*

- P is the **prior path measure**: the joint distribution of **whole trajectories** $(Z_t)_{t \in [0, T]}$ induced by running the diffusion model. The expectation in u_t is taken over these trajectories, those that pass through z_t at time t .
- Boundary: $u_0(z_0) = \exp(r(z_0)/\beta)$, which recovers the static tilt.
- Known by many names: **value function** (RL), *desirability function* (stochastic optimal control), *Schrödinger potential* (Schrödinger bridges).

The time-dependent score decomposition

$$\nabla_{z_t} \log q_t^*(z_t) = \nabla_{z_t} \log p_t(z_t) + \nabla_{z_t} \log u_t(z_t).$$

At $t = 0$, $\log u_0 = r/\beta$, recovering the static decomposition.

u_t is not directly computable

→ **Guidance methods** are different approximations to u_t or $\nabla \log u_t$.

Refresher of key takeaways

The machinery (continuous diffusion)

- **Forward marginal:** $q(z_t | x) = \mathcal{N}(\sqrt{\bar{\alpha}_t} x, (1 - \bar{\alpha}_t)I)$.
- One network, **three views:** $\hat{x}_\theta$ (posterior mean, Tweedie), noise $\hat{\epsilon}_\theta = (z_t - \sqrt{\bar{\alpha}_t} \hat{x}_\theta) / \sqrt{1 - \bar{\alpha}_t}$, score $s_\theta = -\hat{\epsilon}_\theta / \sqrt{1 - \bar{\alpha}_t}$.
- **Reverse step:** substitute $\hat{x}_\theta$ into the Gaussian posterior $q(z_{t-1} | z_t, x)$ & sample.

The goal (idealism-realism)

- **Tilted target:** $q_0^*(x) \propto p_0(x) \exp(r(x)/\beta)$, family $q_t^* \propto p_t u_t$.
- **Score decomposition:** $\nabla_{z_t} \log q_t^* = \nabla_{z_t} \log p_t + \nabla_{z_t} \log u_t$.

Next: controllability mechanisms

Ways to estimate the guidance term $\nabla_{z_t} \log u_t$ (or to avoid it).

Plain conditioning, the simplest mechanism

Enlarge the denoiser's input with a conditioning vector c :

$$\hat{x}_\theta(z_t, t) \longrightarrow \hat{x}_\theta(z_t, t, c).$$

Training and sampling

- Loss: same MSE, threaded with c : $\mathcal{L}(\theta) = \mathbb{E}_{(x,c),t,\epsilon} [\|\hat{x}_\theta(z_t, t, c) - x\|^2]$
- Training data: (x, c) pairs.
- Sampling: identical algorithm, with desired c fed in at every step.

How conditioning enters the network

Three standard mechanisms. The choice does not change the loss or sampler, only how c is presented.

Concatenation

Append c (or an embedding) to the input or to intermediate features. *Simple, robust; works when c is a low-dimensional vector.*

Cross-attention

Treat c as a sequence of tokens. Hidden states query the tokens of c . *Essential when c is itself a sequence (text prompt).* [Stable Diffusion](#), [Imagen](#).

Feature-wise Linear Modulation(FiLM) / adaptive normalization

Produce per-channel scale and shift from c ; modulate normalization layers. *Compact, expressive.* [Ablated Diffusion Model \(ADM\)](#)⁴

⁴Dhariwal & Nichol (2021)

The score-level Bayes identity

Specialized to diffusion latents at noise level t :

$$\nabla_{z_t} \log p_t(z_t | y) = \nabla_{z_t} \log p_t(z_t) + \nabla_{z_t} \log p_t(y | z_t)$$

Two tractable pieces:

- $\nabla_{z_t} \log p_t(z_t)$: the *prior score*, from the diffusion model.
- $\nabla_{z_t} \log p_t(y | z_t)$: the gradient of a separately trained classifier.

Consequence

An **unconditional** diffusion model + a classifier is enough to sample **conditionally**, with no conditional generative training.

This is **classifier guidance**. The classifier gradient plays the role of $\nabla \log u_t$.

Classifier guidance and the guidance scale γ

Dhariwal & Nichol (2021): amplify the classifier term with a scale $\gamma \geq 0$:

$$\nabla_{z_t} \log p_t^\gamma(z_t | y) = \nabla_{z_t} \log p_t(z_t) + \gamma \nabla_{z_t} \log p_t(y | z_t).$$

Equivalent to sampling from $p_t^\gamma(z_t | y) \propto p_t(z_t) \cdot p_t(y | z_t)^\gamma$.

Three regimes

- $\gamma = 0$: **unconditional** sampling. Classifier ignored.
- $\gamma = 1$: the exact **Bayesian posterior** $p_t(z_t | y)$.
- $\gamma > 1$: **sharpened** toward modes where the classifier is most confident.

Typical $\gamma \in [1, 10]$. Higher γ : better alignment, less diversity, eventually artifacts. Setting $\gamma > 1$ is a useful *heuristic*, not exact posterior sampling.

Training the classifier

The classifier must operate on *noisy* z_t at every level: train it **noise-conditionally**.

Off-the-shelf clean-data classifier fails

On high-noise z_t it produces arbitrary outputs; its gradient is **adversarial noise**. The classifier must see the noised distribution itself.

Recipe: noise-conditional classifier $p_\phi(y \mid z_t, t)$

Sample (x, y) , t , ϵ . Form z_t . Train p_ϕ by cross-entropy on (z_t, t, y) .

Classifier-guided sampling

Shift the clean-data predictor by the classifier gradient

$$\tilde{x}_\theta(z_t, t, y) = \hat{x}_\theta(z_t, t) + \frac{1 - \bar{\alpha}_t}{\sqrt{\bar{\alpha}_t}} \gamma \nabla_{z_t} \log p_\phi(y | z_t, t),$$

then take the standard DDPM reverse step using $\tilde{x}_\theta$.

Cost: one diffusion forward + one classifier forward+backward per step.

Reconstruction guidance, for clean-data classifiers

The practical issue

Most scientific property predictors take *clean* inputs, e.g. binding affinity for molecules. Retraining a noise-conditional version is expensive or impossible.

The Diffusion Posterior Sampling (DPS) trick⁵

Evaluate a clean-data classifier $p_\phi(y \mid x)$ on the Tweedie estimate $\hat{x}_\theta(z_t, t)$, then backprop:

$$\nabla_{z_t} \log p_\phi(y \mid \hat{x}_\theta(z_t, t)).$$

- **Cost:** backprop through *both* classifier and diffusion network.
- **Approximation:** biased at high noise (poor $\hat{x}_\theta$), *usually mild in practice*.
- Enables off-the-shelf property predictors with pretrained diffusion models.

⁵Chung et al., (2023)

The reverse Bayes trick

Apply Bayes the *other* way. Starting from $p(y | x) = p(x | y) p(y) / p(x)$:

$$\nabla_{z_t} \log p_t(y | z_t) = \nabla_{z_t} \log p_t(z_t | y) - \nabla_{z_t} \log p_t(z_t)$$

The classifier gradient as a **difference of two scores**:

- **Conditional score** (diffusion net in conditional mode).
- **Unconditional score** (same net in unconditional mode).

Consequence

If a **single diffusion network** can act in both modes, the classifier gradient comes for free.

No separate classifier to train. → **Classifier-free guidance**.⁶

⁶Ho & Salimans, (2022)

Classifier-free guidance: training

Train one diffusion network to act in *both* conditional and unconditional modes via **conditioning dropout**.

Training recipe

During training, with probability p_{drop} (typically 0.1–0.2), replace c with a special null token $\emptyset$ before passing it to the network. Otherwise pass the real c . The loss is unchanged.

After training, the same network functions in two modes:

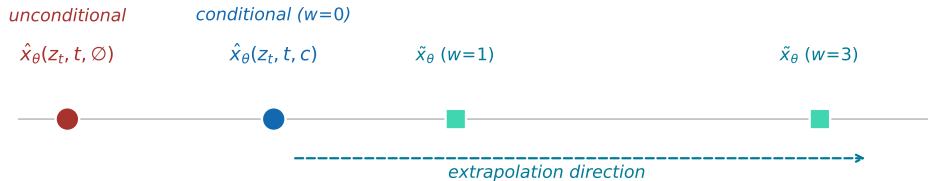
$$\begin{aligned}\hat{x}_{\theta}(z_t, t, c) & \text{ approximates the conditional score} \\ \hat{x}_{\theta}(z_t, t, \emptyset) & \text{ approximates the unconditional score}\end{aligned}$$

Training cost: identical to plain conditional. One extra line of code.

CFG: sampling formula and geometric picture

In the clean-data parameterization, the guided predictor is a linear combination:

$$\tilde{x}_\theta(z_t, t, c) = (1 + w) \hat{x}_\theta(z_t, t, c) - w \hat{x}_\theta(z_t, t, \emptyset).$$



- **Extrapolation**, not interpolation.
- $w = 0$: standard conditional.
- $w \in [1, 7]$: typical for text-to-image.

Cost: 2 forward passes per step (one with c , one with $\emptyset$). No backward passes.

Sequential Monte Carlo steering: the idea

A *different paradigm*: not score modification within one trajectory, but **many trajectories reweighted by reward**.

Target

$$q_0^*(x) \propto p_0(x) \exp(r(x)/\beta), \quad q_t^*(z_t) \propto p_t(z_t) u_t(z_t)$$

u_t : value function, expected $\exp(r/\beta)$ over completions from z_t . Boundary $u_0 = \exp(r/\beta)$. **Reweight** intermediate samples by *potential function* g_t .

Optimal potential $g_t^* = u_t$ is intractable.

But the intermediate g_t **cancel across the trajectory**, so any g_t with $g_0 \propto u_0$ reaches q_0^* . Freedom to pick a cheap, heuristic g_t .

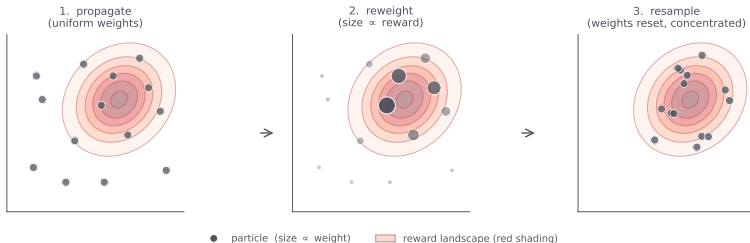
SMC algorithm

Algorithm (weight steering version)

Init $z_T^{(i)} \sim \mathcal{N}(0, I)$, $w^{(i)} = 1/N$. For $t = T, \dots, 1$:

1. **Propagate:** $z_{t-1}^{(i)} \sim p_\theta(z_{t-1} | z_t^{(i)})$.
2. **Reweight:** $w^{(i)} \leftarrow w^{(i)} \cdot g_{t-1}(z_{t-1}^{(i)}) / g_t(z_t^{(i)})$.
3. **Resample:** if $\text{ESS} < N/2$, resample by weight, reset to $1/N$.

Return $\{(z_0^{(i)}, w^{(i)})\}$. Sample: draw one particle $\propto w^{(i)}$.



SMC: resampling

Effective sample size

Each reweight multiplies by a random factor, so weights spread with the number of steps: a few particles capture all the weight.

$$\text{ESS} = \frac{(\sum_i w^{(i)})^2}{\sum_i (w^{(i)})^2} \in [1, N].$$

N : uniform. 1: one particle dominates. Resample when $\text{ESS} < N/2$.

Resampling restores effective sample size

Duplicates high-weight particles, culls the rest. Identical copies **re-separate** at the next step (stochastic kernel). Weight becomes multiplicity, target unchanged.

SMC: potential

Choosing g_t (bias-variance, all valid as $N \rightarrow \infty$)

- **Terminal** ($g_t \equiv 1, t > 0$): no bias via $\hat{x}_\theta$, **highest variance**.
- **Rollout** ($g_t = \exp(r(\hat{x}_\theta)/\beta)$): **lower variance**, biased at high t .
- **Learned twist** (Twisted Diffusion Sampler, TDS⁷), $g_t \approx u_t$: **lowest variance**, training cost. Also steers the proposal.

The consequence

- **No gradients**. Reward enters as a weight, never a derivative.
- Works for **non-differentiable** rewards: simulators, oracles, energy functions.
- Asymptotically exact as $N \rightarrow \infty$.

⁷Wu et al. 2023

Comparing the four methods

	Plain cond.	Classifier g.	CFG	SMC
Acts at	training	inference	training + inf.	inference
Mechanism	conditional NN	score modify	score modify	particle reweight
Approx. of u_t	implicit	explicit (clf. grad.)	implicit	none (asyp. exact)
Training data	(x, c) pairs	unlab. x + (x, y) clf.	(x, c) + dropout	unlab. x + scorer
Reward diff.?	N/A	required	N/A	not required
Grad. at inference?	no	yes	no	no
Cost / step	$1\times$	1 fwd + clf fwd/bwd	$2\times$	$N\times$
Inference knob	none	γ	w	N , potential g_t
Exact at	N/A	$\gamma = 1$	$w = 0$	$N \rightarrow \infty$
Failure mode	weak control	adversarial gradient	sharpened, not posterior	mode collapse

Why discrete data is different

Biological sequences live in *finite alphabets*: 20 amino acids, 4 nucleotides. Continuous diffusion assumes data in $\mathbb{R}^d$.

The central obstacle

$\nabla_x \log p(x)$ is **undefined** on a discrete state space.

The whole score-based machinery does not transfer directly.

What carries over vs. what changes

- **Carries over**: forward corruption, learned reverse denoising, many small steps, *predict-clean-data* training trick.
- **Changes**: forward kernel (no Gaussian), training objective (no score matching), some guidance methods (no classifier guidance via gradients).

Setup and the two main families (D3PM)

Let V = finite alphabet. Extend with a mask token m , giving $|V| = 21$ for proteins. A sequence of length L is $x = (x_1, \dots, x_L) \in V^L$.

Discrete Denoising Diffusion Probabilistic Models (D3PM⁸) forward chain

Position-wise Markov chain over V via row-stochastic transition matrices Q_t :

$$q\left(x_t^{(i)} = j \mid x_{t-1}^{(i)} = k\right) = [Q_t]_{kj}, \quad [Q_t]_{kj} \geq 0, \quad \sum_j [Q_t]_{kj} = 1.$$

Two corruption types dominate:

- **Uniform**: each token replaced by a uniformly random token with prob. β_t .
- **Absorbing (masked)**: each token replaced by the mask token m ; stays masked.

⁸Austin et al., (2021)

Uniform vs absorbing: which to use

Different denoising tasks, different trade-offs.

Uniform corruption

- Network must **detect** corruption *and* predict the original.
- Even far in the chain, each position retains some info.
- Reverse posterior is more complex.

Absorbing (masked)

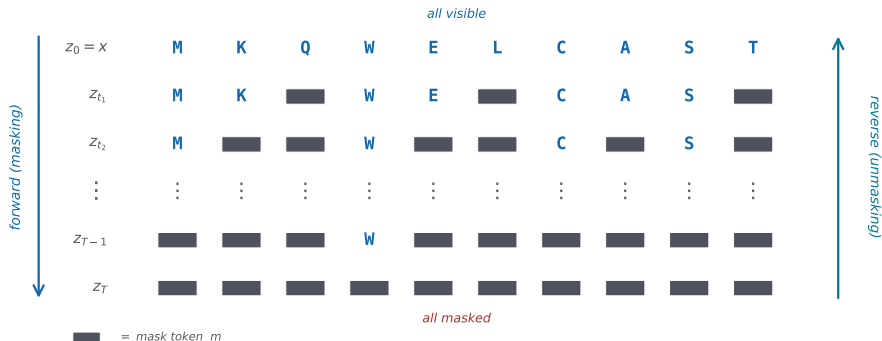
- Corrupted positions are **explicit** (the mask token).
- Each position is either fully known or fully unknown.
- Reverse posterior is simpler: stay masked vs. unmask to x_0 .

Rest of this lecture: absorbing diffusion.

Absorbing forward process

Let $\beta_t \in [0, 1]$ be the per-step mask probability. Position-wise:

$$q\left(x_t^{(i)} \mid x_{t-1}^{(i)}\right) = \begin{cases} (1 - \beta_t) \delta_{x_t^{(i)}=x_{t-1}^{(i)}} + \beta_t \delta_{x_t^{(i)}=m} & \text{if } x_{t-1}^{(i)} \neq m, \\ \delta_{x_t^{(i)}=m} & \text{if } x_{t-1}^{(i)} = m. \end{cases}$$



Absorbing: once a position is masked, it stays masked.

Closed-form marginal: the workhorse equation

Define the cumulative mask probability $\bar{\beta}_t := 1 - \prod_{s=1}^t (1 - \beta_s)$. Then for each position,

$$q\left(x_t^{(i)} \mid x_0^{(i)}\right) = (1 - \bar{\beta}_t) \delta_{x_t^{(i)}=x_0^{(i)}} + \bar{\beta}_t \delta_{x_t^{(i)}=m}.$$

- Each position independently: **kept** with prob. $1 - \bar{\beta}_t$, **masked** with prob. $\bar{\beta}_t$.
- Common choice: set β_t so that $\bar{\beta}_t$ linear from 0 at $t = 0$ to 1 at $t = T$.
- **Jump to any noise level** in one step: sample one Bernoulli per position.
- Compare with continuous: closed-form *Bernoulli mixture* vs. closed-form *Gaussian*, both indexed by a single cumulative quantity ($\bar{\beta}_t$ vs. $\bar{\alpha}_t$).

Reverse process: unmasking and the closed-form posterior

Strong structural constraint from the forward chain:

- If position i is **unmasked** at time t , it must keep the same value at $t - 1$.
- If position i is **masked** at time t , it either stays masked or unmask to some token.

Conditional reverse posterior on a masked position

Conditioning on x_0 makes it tractable (parallels the continuous case):

$$q\left(x_{t-1}^{(i)} \mid x_t^{(i)} = m, x_0^{(i)}\right) = \frac{\bar{\beta}_{t-1}}{\bar{\beta}_t} \delta_{x_{t-1}^{(i)}=m} + \left(1 - \frac{\bar{\beta}_{t-1}}{\bar{\beta}_t}\right) \delta_{x_{t-1}^{(i)}=x_0^{(i)}}.$$

A Bernoulli over {stay masked, unmask to x_0 }.

Training trick (same as continuous): replace x_0 by the network's prediction $\hat{x}_\theta(\cdot \mid x_t, t)_i$, a distribution over V .

Training loss and the BERT connection

After expanding the variational lower bound (Sahoo et al. 2024, Shi et al. 2024):

$$\mathcal{L}(\theta) = \mathbb{E}_{x_0, t, x_t} \left[-\frac{\beta_t}{\bar{\beta}_t} \sum_{i: x_t^{(i)}=m} \log \hat{x}_\theta \left(x_0^{(i)} \mid x_t, t \right)_i \right].$$

Time-weighted cross-entropy on *masked positions only*.

Connection to BERT-style masked language modelling

Same structure: mask tokens, predict them, sum cross-entropy. Differences:

- **Mask ratio:** BERT fixes e.g. 15%; masked diffusion samples it per example, from 0% to 100%.
- **Time weight:** BERT has none; diffusion uses $\beta_t/\bar{\beta}_t$ from the ELBO.

Discrete diffusion is a principled generalization of BERT to a continuum of mask ratios. Models can be *warm-started* from BERT weights.

Sampling: algorithm and order variants

Ancestral sampling

1. Initialize $x_T = (m, m, \dots, m)$.
2. For $t = T, T - 1, \dots, 1$:
 - ▶ Compute $\hat{x}_\theta(\cdot \mid x_t, t)_i$ at every position (one forward pass).
 - ▶ Sample x_{t-1} from the reverse posterior; copy unmasked positions through.
3. Return x_0 .

Sampling order: parallel vs. one-position-per-step

- **Parallel** (default): many positions can be unmasked per step.
- **One-per-step** ($T = L$): exactly one masked position is unmasked at a time, in some order (random, confidence-based, entropy-based).
- Random order $\Rightarrow$ **order-agnostic autoregressive** (OADM, Hoogetboom et al. 2022). **EvoDiff** uses this.
- Same network, **both** regimes: choice is a sampling-time hyperparameter.

Continuous vs. discrete diffusion

	Continuous diffusion	Discrete (masked) diffusion
Data type	real-valued $x \in \mathbb{R}^d$	tokens $x \in V^L$
Forward process	Gaussian noising	probabilistic masking
Per-step kernel	$z_t = \sqrt{1 - \beta_t} z_{t-1} + \sqrt{\beta_t} \epsilon$	each token masked w.p. β_t
Closed-form marginal	$z_t = \sqrt{\bar{\alpha}_t} x + \sqrt{1 - \bar{\alpha}_t} \epsilon$	each position kept w.p. $1 - \bar{\beta}_t$
Reverse step	Gaussian posterior in $\mathbb{R}^d$	categorical posterior on V
Network output	clean $\hat{x}_\theta$ or noise $\hat{\epsilon}_\theta$	categorical $\hat{x}_\theta(\cdot \mid x_t, t)_i$
Training loss	MSE on clean data or noise	weighted cross-entropy on masked positions
$\nabla_x \log p$	defined; learnable as a score	undefined (discrete state)
Bio-seq examples	RFdiffusion, AlphaFold3, OmegAMP	EvoDiff, MDLM, SEDD

The genuine asymmetry is the $\nabla_x \log p$ row. **Consequence:** guidance methods that rely on classifier gradients do not transfer directly.

Guidance for discrete diffusion: what changes

All four methods extend to absorbing diffusion, with different effort.

Summary of changes

- **Plain conditioning:** *trivial port*, identical to continuous.
- **Classifier guidance:** *requires workarounds*, since $\nabla_z \log p$ is undefined on a discrete state space.
- **Classifier-free guidance:** same training trick; sampling formula now mixes *logits*.
- **Sequential Monte Carlo:** structurally identical, and gains a useful new tool: *constrained proposals*.

Practical takeaway: in discrete settings, CFG and SMC are the *workhorses*; classifier guidance is rarely the cleanest choice.

Plain conditioning: a trivial port

Add the conditioning vector c to the denoiser input, as in the continuous case:

$$\hat{x}_\theta(\cdot \mid x_t, t) \longrightarrow \hat{x}_\theta(\cdot \mid x_t, t, c).$$

The network now outputs a *conditional* categorical distribution at every position.

Everything else is unchanged

- Loss: time-weighted cross-entropy on masked positions, conditioned on c .
- Sampling: identical ancestral algorithm, with desired c fed in at every step.
- Training data: (x, c) pairs.

Architectural choices (concatenation, cross-attention, FiLM) carry over identically.

Classifier guidance: the gradient problem and workarounds

The obstacle

Classifier guidance needs $\nabla_{z_t} \log p_\phi(y \mid z_t, t)$. On a discrete state space, **this gradient is undefined**: z_t takes values in a finite set, not in $\mathbb{R}^d$.

Three families of workarounds

- **Hidden-state guidance** (NOS, Nisonoff et al. 2024): backprop the classifier into the network's *continuous* hidden activations $h(z_t)$ rather than into z_t itself.
- **Reconstruction-style**: evaluate a clean-data classifier on the predicted distribution $\hat{x}_\theta(\cdot \mid z_t, t)$ and backprop through it (DPS analog).
- **Logit-level modification**: shift the network's output logits by a learned or heuristic term derived from the classifier.

All are **approximations** with no clean analog to continuous classifier guidance. CFG and SMC are usually preferred for discrete data.

CFG for discrete diffusion: training and logit-space sampling

Training: identical to continuous case. During training, with probability p_{drop} , replace c by the null token $\emptyset$ before passing to the network. After training, the same network operates in conditional and unconditional modes.

Sampling: the network outputs logits $\ell_{\theta}(z_t, t, \cdot)_i \in \mathbb{R}^{|V|}$ at every position. Mix them at the logit level:

$$\tilde{\ell}_{\theta}(z_t, t, c)_i = (1 + w) \ell_{\theta}(z_t, t, c)_i - w \ell_{\theta}(z_t, t, \emptyset)_i, \quad \tilde{p}_i = \text{softmax}(\tilde{\ell}_{\theta}(z_t, t, c)_i).$$

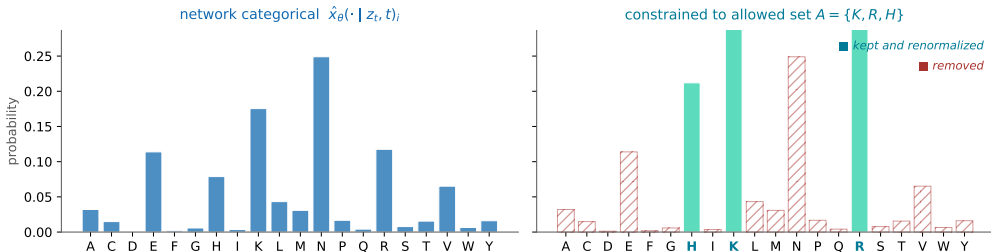
Connection to continuous CFG

Equivalent to sampling from $\tilde{p}_i \propto p_{\theta}(\cdot \mid z_t, t, c)^{1+w} \cdot p_{\theta}(\cdot \mid z_t, t, \emptyset)^{-w}$. Same extrapolation structure as the continuous case, now in logit space.

Cost: $2 \times$ forward passes per step. Workhorse for masked-diffusion text and protein generation.

SMC for discrete diffusion: constrained proposals

The SMC algorithm (propagate, reweight, resample) is **structurally identical**. Discrete data unlocks a useful extra tool: **constrained proposals**.



Why this is natural for discrete

At each masked position the network outputs a full categorical over V . **Restrict** it to an allowed set $A \subset V$, sample from the renormalized restricted distribution, correct with an importance weight. Works without retraining.

Choosing a method in practice

Same trade-offs as the continuous case, with a few discrete-specific notes.

Decision criteria

- Quick baseline with paired data? → **plain conditioning**.
- Differentiable property predictor at hand? → **classifier guidance** *workaround* (NOS / DPS-style).
- Want clean control with a single knob, no extra model? → **classifier-free guidance**.
- Reward is non-differentiable (oracle, simulator), or you have *hard constraints*? → **SMC** (with constrained proposals).

In practice

Production pipelines often combine these: a CFG-trained model as conditional base, with SMC for non-differentiable scorers and hard constraints at inference.

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Guidance

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