TEST-TIME ANCHORING FOR DISCRETE DIFFUSION POSTERIOR SAMPLING

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ABSTRACT

We study the problem of posterior sampling using pretrained discrete diffusion foundation models, aiming to recover images from noisy measurements without retraining task-specific models. While diffusion models have achieved remarkable success in generative modeling, most advances rely on continuous Gaussian diffusion. In contrast, discrete diffusion offers a unified framework for jointly modeling categorical data such as text and images. Beyond unification, discrete diffusion provides faster inference, finer control, and principled training-free Bayesian inference, making it particularly well-suited for posterior sampling. However, existing approaches to discrete diffusion posterior sampling face severe challenges: derivative-free guidance yields sparse signals, continuous relaxations limit applicability, and split Gibbs samplers suffer from the curse of dimensionality. To overcome these limitations, we introduce Anchored Posterior Sampling (APS) for *masked diffusion* foundation models, built on two key innovations—*quantized expectation* for gradient-like guidance in discrete embedding space, and *anchored remasking* for adaptive decoding. Our approach achieves state-of-the-art performance among discrete diffusion samplers across linear and nonlinear inverse problems on the standard benchmarks. We further demonstrate training-free stylization and text-guided editing using our sampler.

1 INTRODUCTION

Diffusion models have become the state-of-the-art across a wide range of generative tasks, including images (Ramesh et al., 2021; Rombach et al., 2022; Baldridge et al., 2024; Esser et al., 2024; Black Forest Labs, 2024), audio (Huang et al., 2023; Veo, 2025), and video (Singer et al., 2023; OpenAI, 2024; Veo, 2025). Most of this progress has been driven by *continuous* diffusion models, where Gaussian noise is gradually added in pixel or latent space and then reversed by a learned denoiser (Sohl-Dickstein et al., 2015; Ho et al., 2020). Recently, however, *discrete* diffusion has emerged as a powerful alternative, showing superior performance in modeling categorical distributions such as text (Lou et al., 2024; Sahoo et al., 2024; Shi et al., 2024; Nie et al., 2025; Rout et al., 2025a) and images (Yang et al., 2025). Discrete diffusion further enables a unified framework for both image and text generation, supporting multimodal generation and editing.

Beyond unification, discrete diffusion offers several advantages over continuous diffusion that are particularly relevant for posterior sampling. First, it achieves *faster inference*, often generating high-quality samples in significantly fewer reverse steps (Shi et al., 2024; Schiff et al., 2025; Ma et al., 2025). Second, it provides *finer control*: the model predicts a normalized categorical distribution per token (e.g., a pixel or a patch), which decouples different parts of the image, unlike Gaussian diffusion where the entire image is coupled. Third, it enables *training-free posterior sampling*: since the model outputs full conditional distributions at each step, these can be reweighted by the likelihood to yield a better posterior estimate posterior (Murata et al., 2024; Chu et al., 2025). This property unlocks precise image editing and solving inverse problems without additional training (§4), motivating our approach to use discrete diffusion (Yang et al., 2025) as a prior.

State-of-the-art posterior samplers (Rout et al., 2023; 2024; Chung et al., 2024; Zhang et al., 2025) use continuous diffusion as a prior. These approaches rely on guiding the reverse diffusion process using likelihood gradients in continuous latent spaces (Rout et al., 2023; 2024; Chung et al., 2024; Zhang et al., 2025). This is infeasible for *discrete* diffusion due to non-differentiability in token space. Derivative-free discrete methods (Li et al., 2024) inspired by reinforcement learning provide weak signals. G2D2 (Murata et al., 2024) uses Gumbel-softmax relaxation but it is limited to discrete tokens with continuous embeddings. SGDD (Chu et al., 2025) introduces a split Gibbs sampler, but
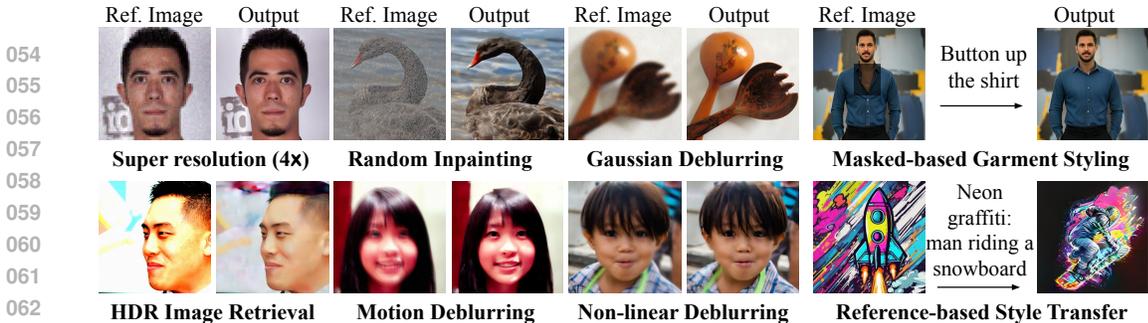
Figure 1: We introduce Anchored Posterior Sampling (APS) for *masked diffusion* foundation models, built on two key innovations: (i) *quantized expectation*, which provides gradient-like guidance in discrete embedding space, and (ii) *anchored remasking*, which enables adaptive decoding. Our method supports a variety of linear and nonlinear image restoration tasks (left three columns), as well as mask-based garment styling and reference-guided style transfer (last column).

suffers from exponential complexity in sequence length (Chewi, 2023). Moreover, these methods unmask tokens in *random order*, which is suboptimal compared to adaptive decoding strategies in language modeling (Yang et al., 2025; Rout et al., 2025a). These limitations underscore the need for a discrete diffusion posterior sampler with adaptive decoding, leveraging next-generation multimodal models (Yang et al., 2025; Gemini Team, 2024).

Existing discrete diffusion methods explore posterior sampling under uniform (Chu et al., 2025) or mixed-noise processes (Murata et al., 2024), but these methods are tied to specific noising schemes and lack generalization (§4). Meanwhile, recent advances show that *masked diffusion* achieves state-of-the-art performance in image generation (Ma et al., 2025; Yang et al., 2025), yet its potential for posterior sampling remains underexplored. In this work, we take the first step towards leveraging multimodal masked diffusion models such as MMaDA (Yang et al., 2025) for posterior sampling. We introduce two key algorithmic innovations: (i) *quantized expectation*, which provides gradient-like guidance in purely discrete embedding space by updating the full conditional probability table (§3.2.1), and (ii) *anchored remasking*, an adaptive inference strategy that decodes important "anchor" tokens early in the reverse process (§3.2.2). Together, these techniques overcome the non-differentiability challenge in discrete diffusion and yield scalable posterior samplers. Extensive experiments on linear and nonlinear inverse problems demonstrate improvements of up to 31.36% in LPIPS and 7.05% in PSNR (§4), and showcase training-free stylization as illustrated in Figure 1.

Our contributions are summarized below.

- **Theoretical results:** we derive (i) a *training* upper bound, $\mathcal{L}_{\text{DDPS}}$ (**Theorem 3.1**), that integrates measurements into the reverse diffusion process, and (ii) a *test-time* bound, $\mathcal{L}_{\text{APS}}$ (**Theorem 3.2**), that reuses a pretrained denoiser without expensive retraining per downstream task (§3).

- **Quantized expectation:** a novel strategy to update *all* entries of the conditional probability table, enabling strong *gradient-like* guidance in the discrete embedding space (§3.2.1).

- **Anchored remasking:** an adaptive decoding strategy that unmasks "anchor" tokens early in the reverse process, better utilizing model's capacity to decode remaining tokens (§3.2.2).

- **Extensive experiments:** comprehensive evaluation on linear (super resolution, Gaussian deblurring, inpainting, motion deblur) and nonlinear (HDR, nonlinear deblurring) inverse problems on FFHQ and ImageNet, where our method achieves up to 35.82% LPIPS and 10.94% PSNR gains on ImageNet super resolution ($4\times$), and 31.36% LPIPS and 7.05% PSNR gains on FFHQ, over the prior state-of-the-art discrete sampler. We further demonstrate *training-free stylization* enabled by our discrete posterior sampler, highlighting flexibility beyond inverse problems (§4).

2 RELATED WORKS

Gaussian pixel-space diffusion methods (Chung et al., 2023; Kawar et al., 2022; Zhu et al., 2023) address noisy inverse problems by guiding the reverse process of diffusion models trained directly in pixel space. These priors are domain-specific—for instance, a model trained on ImageNet must be used for ImageNet tasks, and one trained on FFHQ for FFHQ tasks—yielding highly informative but narrow priors. Mixing domains during training dilutes this specificity, and at the extreme of

internet-scale training, the prior remains valid for generation (Ramesh et al., 2021; Baldridge et al., 2024; Esser et al., 2024; Black Forest Labs, 2024) but is less informative for domain-specific inverse problems. This motivates the challenge of extracting domain-relevant information from general-purpose priors.

Gaussian latent diffusion. PSLD (Rout et al., 2023) introduced posterior sampling with latent diffusion, showing how domain-specific priors can be extracted from large-scale foundation models. This line of work (Chung et al., 2024; Song et al., 2024; Rout et al., 2024; 2025b; Noroozi et al., 2024; Zhang et al., 2025; Chung et al., 2025) leverages the advantages of latent diffusion: a single pretrained model can handle multiple domains, enabling inverse problems and semantic edits without retraining, while also being faster and more scalable to high-resolution synthesis. A drawback, however, is that posterior sampling often requires backpropagation through large denoisers (e.g., Flux (Black Forest Labs, 2024), SD3.5 (Esser et al., 2024)), which is prohibitively slow. RB-Modulation (Rout et al., 2025c) alleviates this by framing the problem as stochastic optimal control, directly optimizing the terminal latent state and reducing runtime from minutes (PSLD: ~12 min, P2L: ~30 min, STSL: ~3 min) to under 40 seconds with minimal overhead. This efficiency relies on continuous, differentiable latent embeddings, a property that does not extend to discrete diffusion. Addressing this gap motivates the need for new posterior sampling approaches in discrete settings.

Uniform discrete diffusion. Recent works have explored posterior sampling with discrete diffusion. G2D2 (Murata et al., 2024) extends the proximal sampler of RB-Modulation to VQ-diffusion (Gu et al., 2022) using a star-shaped noising process and Gumbel-Softmax dequantization (Gumbel, 1954; Jang et al., 2017; Maddison et al., 2017). While it enables gradient guidance, G2D2 depends on continuous relaxations, requires storing log-probabilities from previous step, and struggles to generalize to purely discrete token embeddings (§4). SGDD (Chu et al., 2025) instead proposes a split Gibbs sampler with Hamming-distance reweighting and rejection sampling via Metropolis–Hastings, but its exponential rejection rate restricts results to low-resolution tasks.

Masked (absorbing) discrete diffusion. While G2D2 and SGDD can, in principle, be adapted to masked diffusion, they perform poorly with purely discrete token embeddings. In contrast, our method leverages a unified masked discrete diffusion model and introduces two key components: *quantized expectation* (§3.2.1) and *anchored remasking* (§3.2.2). Together, these yield an efficient and scalable posterior sampler for high-resolution inverse problems. To our knowledge, this is the first inverse problem solver tailored for masked discrete diffusion with purely discrete embeddings, outperforming prior discrete samplers and remaining competitive—often superior—to continuous diffusion methods at substantially lower inference cost (§4).

3 METHOD

3.1 PRELIMINARIES

Visual Tokenizer. A cornerstone of modern image tokenization is the VQ-VAE (Van Den Oord et al., 2017), which maps images into discrete codebook indices. It consists of an encoder $\mathcal{E} : \mathbb{R}^{H \times W \times 3} \to \mathbb{R}^{h \times w \times d}$ that projects an image into a latent embedding of size $h \times w \times d$. The embeddings are reshaped into a sequence $\mathbf{e} \in \mathbb{R}^{L \times d}$ of length $L = h \times w$. We denote images by $\hat{\mathbf{x}}$. The encoder produces $\mathbf{e} = \mathcal{E}(\hat{\mathbf{x}})$ where each embedding $\mathbf{e}^l$ for $l = 1, \ldots, L$ is quantized to the nearest codebook entry $\mathbf{c}_j \in \mathcal{C}$, where $\mathcal{C} \in \mathbb{R}^{K \times d}$ is a learned codebook:

$$\mathcal{Q}_{\text{vq}}(\mathbf{e}^l) \coloneqq \mathbf{c}_j, \quad j = \underset{k \in \{1, \ldots, K\}}{\arg\min} \left\| \mathbf{e}^l - \mathbf{c}_k \right\|_2. \tag{1}$$

Codebook entries may be continuous vectors $\mathbf{c}_j \in \mathbb{R}^d$ or purely discrete binary embeddings $\mathbf{c}_j \in \{-1, +1\}^d$ used in this paper. Binary embeddings are particularly appealing because prior studies (Yu et al., 2024) show that masked diffusion models degrade in generation quality as continuous vocabulary size grows, yet lookup-free quantization (LFQ) with binary embeddings achieves both strong generation and reconstruction quality. In LFQ, the codebook is not learned but obtained by thresholding the encoder output:

$$\mathcal{Q}_{\text{lfq}}(\mathbf{e}^l) \coloneqq \text{sign}(\mathbf{e}^l), \text{ where } [\mathcal{Q}_{\text{lfq}}(\mathbf{e}^l)]_i = \begin{cases} +1 & \text{if } \mathbf{e}^l[i] > 0, \\ -1 & \text{otherwise.} \end{cases} \tag{2}$$

Finally, each image is represented as a sequence of tokens $x = (x^1, \ldots, x^L)$ corresponding to the selected indices from (1) as $x^l = j$, or equivalently for LFQ the token index is obtained by

$x^l = j = \sum_{i=1}^{d} 2^{i-1} \mathbf{1}_{\{\mathbf{e}_l[i]>0\}}$. Equivalently, we represent each sequence as a mixture of one-hot vectors $\mathbf{x} = (\mathbf{x}^1, \ldots, \mathbf{x}^L)$, where $\sum_{k=1}^{K} \mathbf{x}^l[k] = 1, \mathbf{x}^l[k] \geq 0$ and $\mathbf{x}^l[x^l] = 1$. This discrete token representation forms the basis of our masked diffusion posterior sampler.

Notation: We use ':=' to indicate architectural and parameterization choices in our model, to distinguish from '=' which are identities that follow from mathematical derivations.

Masked Diffusion. We now describe masked discrete diffusion, which defines a generative model over the discrete state space $\mathcal{S} = \mathcal{V}^L$, where $\mathcal{V} = \{1, \ldots, K, K+1\}$ consisting of K codebook indices and a special [MASK] token $\mathbf{m}$ corresponding to index $K+1$. Let $X = \{x^l\}_{l=1}^{L} \in \mathcal{S}$ denote a sequence of tokens (or equivalently its one-hot representation $\mathbf{x} = \{\mathbf{x}^l\}_{l=1}^{L}$). The data distribution is denoted by $q(\cdot)$ over $\mathcal{S}$. The goal of masked diffusion is to learn a generative model that samples from $q(\cdot)$. Masked diffusion models (MDMs) (Austin et al., 2021; Lou et al., 2024; Sahoo et al., 2024; Shi et al., 2024) construct a discrete-time Markov chain with T steps, parameterized by α_t with $t \in [0, 1]$. The forward process gradually replaces each token with the [MASK] token:

$$q(\mathbf{z}_t|\mathbf{x}) = \prod_{l=1}^{L} q(\mathbf{z}_t^l|\mathbf{x}), \quad q(\mathbf{z}_t^l|\mathbf{x}) = \mathrm{Cat}\left(\mathbf{z}_t^l; \alpha_t \mathbf{x}^l + (1-\alpha_t)\mathbf{m}\right), \quad (3)$$

where $\mathbf{z}_t^l$ is either preserved from $\mathbf{x}^l$ with probability α_t or replaced with $\mathbf{m}$ otherwise. The corresponding reverse process is parameterized by a neural network p_θ, which predicts categorical distributions over tokens. For each token position l, the transition distribution is

$$p_\theta(\mathbf{z}_s^l|\mathbf{z}_t) := q(\mathbf{z}_s^l|\mathbf{z}_t^l, \mathbf{x}_\theta(\mathbf{z}_t)) = \begin{cases} \mathrm{Cat}(\mathbf{z}_s^l; \mathbf{z}_t^l), & \mathbf{z}_t^l \neq \mathbf{m}, \\ \mathrm{Cat}\left(\mathbf{z}_s^l; \frac{\alpha_s - \alpha_t}{1-\alpha_t}\mathbf{x}_\theta(\mathbf{z}_t) + \frac{1-\alpha_s}{1-\alpha_t}\mathbf{m}\right), & \mathbf{z}_t^l = \mathbf{m}, \end{cases} \quad (4)$$

where $\mathbf{x}_\theta(\mathbf{z}_t)$ denotes the network prediction. Training minimizes the negative evidence lower bound (NELBO) by aligning the reverse transition $p_\theta(\mathbf{z}_s^l|\mathbf{z}_t)$ with the inference posterior $q(\mathbf{z}_s^l|\mathbf{z}_t^l, \mathbf{x})$ derived from the forward process (3). Concretely, the training objective $\mathcal{L}_{\mathrm{NELBO}}(\mathbf{x}; \theta) :=$

$$\mathbb{E}_{Z_0 \sim q(\cdot|\mathbf{x})}\left[-\log p_\theta(\mathbf{x}|Z_0)\right] + \sum_{i=1}^{T} \mathbb{E}_{Z_t \sim q(\cdot|\mathbf{x})}\left[\frac{\alpha_t - \alpha_s}{1-\alpha_t}\sum_{l=1}^{L}\log\langle\mathbf{x}_\theta^l(Z_t), \mathbf{x}^l\rangle\mathbf{1}_{\{Z_t^l=\mathbf{m}\}}\right], \quad (5)$$

where, for brevity, we drop i from $t(i) = i/T$ and $s(i) = (i-1)/T$.

3.2 Test-Time Anchored Posterior Sampling

This section develops our theoretical framework for posterior sampling with discrete diffusion. We first derive a training-based objective (**Theorem 3.1**) that incorporates measurements into the reverse diffusion process, and then present a lightweight test-time training bound (**Theorem 3.2**) that reuses a pretrained denoiser without retraining. Building on this, we introduce two key mechanisms—*Quantized Expectation* (§3.2.1) for differentiable likelihood evaluation and *Anchored Remasking* (§3.2.2) for adaptive unmasking—that enable scalable and accurate posterior sampling.

In posterior sampling, our goal is to construct a Markov chain whose stationary distribution coincides with the Bayesian posterior: $q(\mathbf{x}|\mathbf{y}) \propto q(\mathbf{y}|\mathbf{x})\,q(\mathbf{x})$, where $\mathbf{y} = \mathcal{A}(\mathcal{D}(\mathbf{x})) + \sigma\varepsilon$ with measurement operator $\mathcal{A}(\cdot)$, image decoder $\mathcal{D}(\cdot)$, Gaussian noise $\varepsilon \sim \mathcal{N}(0, I)$, and standard deviation σ. When $\mathcal{A}$ is linear the task reduces to a *linear inverse problem*; otherwise a *nonlinear inverse problem*. We approximate $q(\mathbf{x}|\mathbf{y})$ with a tractable sampler $p_\varphi(\mathbf{x}|\mathbf{y})$ using only a masked diffusion model $p_\theta(\mathbf{x})$ previously trained to approximate $q(\mathbf{x})$.

To sample from the posterior $q(\cdot|\mathbf{y})$, we construct a Markov chain with the joint distribution defined as: $p_\varphi(\mathbf{x}, \mathbf{z}_{0:1}|\mathbf{y}) = p_\varphi(\mathbf{z}_1|\mathbf{y})\,p_\varphi(\mathbf{x}|\mathbf{z}_0, \mathbf{y})\prod_{i=1}^{T} p_\varphi\left(\mathbf{z}_{s(i)}|\mathbf{z}_{t(i)}, \mathbf{y}\right)$, where $\mathbf{z}_{0:1} = \mathbf{z}_0, \mathbf{z}_{1/T}, \ldots, \mathbf{z}_1$. We parameterize measurement conditional transitions by tilting the unconditional transition (4) with the likelihood of measurements given the current estimate:

$$p_\varphi(\mathbf{z}_s|\mathbf{z}_t, \mathbf{y}) := \prod_{l=1}^{L} p_\varphi(\mathbf{z}_s^l|\mathbf{z}_t, \mathbf{y}), \text{ where } p_\varphi(\mathbf{z}_s^l|\mathbf{z}_t, \mathbf{y}) \propto q\left(\mathbf{z}_s^l|\mathbf{z}_t^l, \mathbf{x}_\varphi(\mathbf{z}_t)\right)q\left(\mathbf{y}|\mathbf{x}_\varphi(\mathbf{z}_t; \mathbf{z}_s^l)\right). \quad (6)$$

We can compute the measurement likelihood $q(\mathbf{y}|\mathbf{x})$ given a sequence of tokens $\mathbf{x} = (\mathbf{x}^1, \mathbf{x}^2, \ldots, \mathbf{x}^L)$, where each $\mathbf{x}^l$ is a one-hot vector. During inference, we have access to the model output $\mathbf{x}_\varphi(\mathbf{z}_t)$ rather than the ground-truth sequence $\mathbf{x}$. With slight abuse of notation, we define the

log likelihood given the model's output as $\log q(\mathbf{y}|\mathbf{x}_\varphi(\mathbf{z}_t)) := \mathbb{E}_{\mathbf{x} \sim \mathbf{x}_\varphi(\mathbf{z}_t)}[\log q(\mathbf{y}|\mathbf{x})]$. Similarly, $\log q(\mathbf{y}|\mathbf{x}_\varphi(\mathbf{z}_t; \mathbf{z}_s^l))$ denotes the same log likelihood, but with $\mathbf{x}^l$ replaced by $\mathbf{z}_s^l$ while keeping all other positions drawn from $\mathbf{x}_\varphi(\mathbf{z}_t)$. We denote the normalizing factor in (6) by $\mathcal{Z}_\varphi(\mathbf{z}_t, \mathbf{y})$.

Theorem 3.1 (Discrete Diffusion Posterior Sampling (DDPS)). *Given a sample* $\mathbf{x} \sim q$, *let* $q(Z_{0:1}|\mathbf{x})$ *denote the forward noising law of (3). Then, for any measurement* $\mathbf{y} \sim q(\cdot|\mathbf{x})$, $-\log p_\varphi(\mathbf{x}|\mathbf{y}) \leq$

$$\mathcal{L}_{\text{DDPS}}(\mathbf{x}, \mathbf{y}; \varphi) := \mathbb{E}_{q(Z_0|\mathbf{x})}[-\log p_\varphi(\mathbf{x}|Z_0)] + \sum_{i=1}^T \mathbb{E}_{q(Z_t|\mathbf{x})}\big[\log \mathcal{Z}_\varphi(Z_t, \mathbf{y})\big] +$$

$$\sum_{i=1}^T \mathbb{E}_{q(Z_t|\mathbf{x})}\left[\frac{\alpha_t - \alpha_s}{1 - \alpha_t} \sum_{l=1}^L \log \langle \mathbf{x}_\varphi^l(Z_t), \mathbf{x}^l \rangle \mathbf{1}_{\{Z_t^l = \mathbf{m}\}} - \mathbb{E}_{q(Z_s|Z_t, \mathbf{x})}\left[\sum_{l=1}^L \log q(\mathbf{y}|\mathbf{x}_\varphi(Z_t; Z_s^l))\right]\right].$$

Implications. Theorem 3.1 shows that $\mathcal{L}_{\text{DDPS}}(\mathbf{x}, \mathbf{y}; \varphi)$ is a principled training criterion for discrete posterior samplers. The likelihood-based tilt $\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_s))$ enforces measurement consistency When $\mathbf{y}$ is absent, the tilting terms vanish and the objective reduces to the standard masked diffusion NELBO (5). Following the widely adopted approximation from diffusion guidance, we can treat $\mathcal{Z}_\varphi$ as a *stop-gradient term*—used only for normalization in the forward computation but excluded from gradient updates (Sohl-Dickstein et al., 2015; Dhariwal & Nichol, 2021; Ho & Salimans, 2022; Murata et al., 2024). Although expensive, a more formal treatment for the gradient of the log partition function can be obtained through importance sampling (Kahn, 1950), Hamiltonian Monte Carlo with leapfrog steps (MacKay, 2003), or noise contrastive estimation (Gutmann & Hyvärinen, 2010). The cross-entropy term $\log \langle \mathbf{x}_\varphi^l(Z_t), \mathbf{x}^l \rangle$ is zero for clean tokens and gets supervision for masked tokens, with weights determined by the noise schedule.

For retraining, one can minimize $\mathcal{L}_{\text{DDPS}}(\mathbf{x}, \mathbf{y}; \varphi)$ with respect to φ to obtain a discrete posterior sampler. In practice, however, retraining a large-scale foundation model per task is often expensive due to excessive compute and lack of training data. We therefore focus on the training-free case.

Theorem 3.2 (Test-time Anchored Poster Sampling (APS)). *Given a sample* $\mathbf{x} \sim q$, *let* $q(Z_{0:1}|\mathbf{x})$ *denote the forward noising law of (3). Suppose the pretrained network* $p_\theta(\mathbf{x})$ *closely approximates the unconditional prior* $q(\mathbf{x})$. *Then, for any measurement* $\mathbf{y} \sim q(\cdot|\mathbf{x})$, *the negative log-posterior*

$$-\log p_\varphi(\mathbf{x}|\mathbf{y}) \leq \mathcal{L}_{\text{APS}}(\mathbf{x}, \mathbf{y}; \varphi) := \mathcal{L}_{\text{NELBO}}(\mathbf{x}; \theta) + \sum_{i=1}^T \mathbb{E}_{q(Z_t|\mathbf{x})}\left[\log \mathcal{Z}_\varphi(Z_t, \mathbf{y})\right] +$$

$$\sum_{i=1}^T \mathbb{E}_{q(Z_t|\mathbf{x})}\left[\frac{\alpha_s - \alpha_t}{1 - \alpha_t} \sum_{l=1}^L \log \frac{\langle \mathbf{x}_\theta^l(Z_t), \mathbf{x}^l \rangle}{\langle \mathbf{x}_\varphi^l(Z_t), \mathbf{x}^l \rangle} \mathbf{1}_{\{Z_t^l = \mathbf{m}\}} - \mathbb{E}_{q(Z_s|Z_t, \mathbf{x})}\left[\sum_{l=1}^L \log q(\mathbf{y}|\mathbf{x}_\varphi(Z_t; Z_s^l))\right]\right].$$

Implications. Theorem 3.2 shows that posterior sampling can be performed *without retraining* by reusing a pretrained masked diffusion model.

- *Efficient test-time training.* The bound $\mathcal{L}_{\text{APS}}(\mathbf{x}, \mathbf{y}; \varphi)$ is expressed in terms of the pretrained NELBO from (5). Since this term is constant with respect to the new parameters φ, it can be ignored during optimization. As a result, test-time training only needs to update the lightweight adaptation and measurement-consistency terms, while reusing the fixed pretrained network $\mathbf{x}_\theta(\cdot)$. This avoids backpropagation through the large-scale denoiser (e.g., billions of parameters), making posterior sampling feasible and efficient at test time.

- *Training-free inference.* Although efficient test-time training requires paired $(\mathbf{x}, \mathbf{y})$ data, in posterior sampling we only observe $\mathbf{y}$. Interestingly, the bound $\mathcal{L}_{\text{APS}}$ points to training-free inference by substituting the pretrained model prediction $\mathbf{x}_\theta^l(Z_{t(i)})$ in place of $\mathbf{x}^l$ (see §3.2.1). This makes posterior sampling feasible directly from measurements, without retraining or labeled pairs.

- *Adaptation gap.* The log-ratio terms capture the mismatch between unconditional predictions $\mathbf{x}_\theta(Z_{t(i)})$ and adapted posterior predictions $\mathbf{x}_\varphi(Z_{t(i)})$, active only at masked positions.

- *Measurement consistency.* The final summation enforces agreement with the measurement likelihood, ensuring the sampler produces samples consistent with observed measurements $\mathbf{y}$.

Discussion. In summary, retraining a new network φ for every task would require backpropagation through massive denoisers (e.g., 8B parameters in MMaDA), which is computationally prohibitive. Our theoretical results show that posterior sampling can be done efficiently by reusing the pretrained model and optimizing only lightweight parameters at test time. Next, we introduce two key ingredients—*Quantized Expectation* (§3.2.1) and *Anchored Remasking* (§3.2.2)—that make this training-free posterior sampling practically implementable. These two ideas together form our **Algorithm 1: Anchored Posterior Sampling (APS)**; please see Appendix B.2 for a detailed discussion.

Table 1: **Quantitative results on Super Resolution (4×) and Gaussian Deblurring.** APS consistently outperforms prior discrete samplers (G2D2, SGDD, and SVDD-PM) and remains competitive with strong continuous diffusion baselines (shaded gray).

		(a) FFHQ				(b) ImageNet			
		SR (4×)		Deblur		SR (4×)		Deblur	
Type	Method	LPIPS ↓	PSNR ↑	LPIPS ↓	PSNR ↑	LPIPS ↓	PSNR ↑	LPIPS ↓	PSNR ↑
Pixel	DPS	0.269	25.86	0.219	25.87	0.367	22.61	0.443	19.04
	DDRM	0.282	26.58	0.239	24.93	0.352	24.00	0.246	27.30
	DiffPIR	0.260	26.64	0.236	27.36	0.371	23.18	0.355	22.80
	DAPS	0.177	29.07	0.165	29.19	0.276	25.89	0.253	26.15
Latent	PSLD	0.276	27.62	0.304	27.37	0.332	24.43	0.365	24.04
	ReSample	0.507	22.98	0.329	25.69	0.382	22.63	0.438	22.32
	LatentDAPS	0.182	27.48	0.234	27.93	0.276	25.06	0.345	25.05
Uniform (Mask)	SVDD-PM	0.594	12.08	–	–	–	–	–	–
	G2D2	0.271	26.93	0.287	26.35	0.349	23.20	0.375	22.71
	SGDD	0.288	25.85	–	–	–	–	–	–
Mask	**APS**	**0.234**	**27.50**	**0.276**	**27.90**	**0.324**	**24.30**	**0.375**	**24.71**
	APS-L	**0.186**	**28.83**	**0.241**	**29.50**	**0.224**	**25.74**	**0.282**	**26.35**

3.2.1 QUANTIZED EXPECTATION

Following the standard practice in test-time optimization (Chung et al., 2023; Rout et al., 2023; 2024; Chung et al., 2024; Rout et al., 2025c; Zhang et al., 2025; Murata et al., 2024), the upper bound in **Theorem 3.2** is optimized at every time step t. Ignoring the normalizing factor via the stop-gradient trick discussed in §3.2, the optimization objective per time step is defined as

$$\mathcal{L}_t(\varphi) := \frac{\alpha_s - \alpha_t}{1 - \alpha_t} \sum_{l=1}^{L} \left[\log \frac{\langle \mathbf{x}_\theta^l(\mathbf{z}_t), \mathbf{x}^l \rangle}{\langle \mathbf{x}_\varphi^l(\mathbf{z}_t), \mathbf{x}^l \rangle} \mathbf{1}_{\{\mathbf{z}_t^l = \mathbf{m}\}} \right] - \mathbb{E}_{q(Z_s|\mathbf{z}_t, \mathbf{x})} \left[\sum_{l=1}^{L} \log q(\mathbf{y}|\mathbf{x}_\varphi(\mathbf{z}_t; Z_s^l)) \right].$$

This form reuses the pretrained base model $\mathbf{x}_\theta(\cdot)$ but exactly evaluating it remains intractable. Since we have access to the latent variable $\mathbf{z}_t$ during inference, we can easily compute $\mathbf{x}_\theta(\mathbf{z}_t)$ and $\mathbf{x}_\varphi(\mathbf{z}_t)$. However, the measurement consistency term (the second term in $\mathcal{L}_t(\varphi)$) requires $\mathcal{O}(K^L)$ compute[1].

Since $q(\mathbf{y}|\mathbf{x}_\varphi(\mathbf{z}_t; Z_s^l))$ involves an expectation over a parameterized distribution (as defined in (6)), one could in principle, optimize $\mathcal{L}_t(\varphi)$ using policy-gradient methods such as REINFORCE (Williams, 1992), PPO (Schulman et al., 2017), or GRPO (Shao et al., 2024), as adopted in SVDD (Li et al., 2024). However, this leads to high-variance gradients and sparse rewards, resulting in unstable optimization and poor sample quality, as reflected in Table 1.

To overcome this, we introduce *quantized expectation*, a differentiable approximation to $\mathcal{L}_t(\varphi)$. Building on the *training-free inference* implication of **Theorem 3.2**, we note that only the measurements $\mathbf{y}$ are observed at test time, so we replace the ground-truth token $\mathbf{x}^l$ with its predicted mixture $\mathbf{x}_\theta^l(\mathbf{z}_t)$. Recall that the measurement likelihood is modeled as $q(\mathbf{y}|\mathbf{x}) \propto \exp(-\frac{\|\mathbf{y} - \mathcal{A}(\mathcal{D}(\mathbf{x}))\|_2^2}{2\sigma^2})$, where $\mathcal{D}(\cdot)$ is the decoder. A naive sampling approach $\mathbf{x} \sim \mathbf{x}_\varphi(\mathbf{z}_t)$ makes $\mathcal{L}_t(\varphi)$ non-differentiable and noisy. Instead, our quantized expectation formulation approximates it as

$$\hat{\mathcal{L}}_t(\varphi) := \frac{\alpha_t - \alpha_s}{1 - \alpha_t} \sum_{l=1}^{L} \left[\log \langle \mathbf{x}_\varphi^l(\mathbf{z}_t), \mathbf{x}_\theta^l(\mathbf{z}_t) \rangle \mathbf{1}_{\{\mathbf{z}_t^l = \mathbf{m}\}} \right] - L \cdot \log q(\mathbf{y}|\tilde{\mathbf{x}}_\varphi(\mathbf{z}_t)), \tag{7}$$

where $\tilde{\mathbf{x}}_\varphi(\mathbf{z}_t)$ is defined using a quantized expectation as discussed below.

Differentiability has propelled posterior sampling to achieve state-of-the-art results using continuous diffusion (Chung et al., 2023; Rout et al., 2023; 2024; Zhang et al., 2025). To restore differentiability in discrete diffusion, we parameterize $\mathbf{x}_\varphi(\mathbf{z}_{t(i)}) = \text{Softmax}(\varphi_{t(i)})$ with $\varphi_{t(i)} = \{\varphi_{t(i)}^l\}_{l=1}^L \in \mathbb{R}^{K \times L}$ containing logits $\varphi_{t(i)}^l$ over the codebook $\{\mathbf{c}_k \in \mathcal{C}\}$ for each position l. We compute the expected embedding $\bar{\mathbf{x}}^l = \sum_{k=1}^K \mathbf{c}_k \mathbf{x}_\varphi^l(\mathbf{z}_{t(i)}) \in \mathbb{R}^d$, and then *quantize* it using LFQ (2) (Yu et al., 2024): $\mathbf{x}^l = \mathcal{Q}_{\text{lfq}}(\bar{\mathbf{x}}^l) \in \{-1, +1\}^d$. We then apply the straight-through estimator (Van Den Oord

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[1] Recall that the $\log(\cdot|\cdot)$ term within the outer expectation is defined by an expectation in §3.2. The marginalization over Z_s in the outer expectation costs $\mathcal{O}(2^L)$ (given Z_t and $\mathbf{x}$, observe that $Z_s \in \{\mathbf{x}, \mathbf{m}\}$). For each Z_s, the computation of the inner expectation inside the summation over other random variables except Z_s^l costs $\mathcal{O}(LK^{L-1})$. Thus, the total cost becomes $\mathcal{O}(K^L)$.
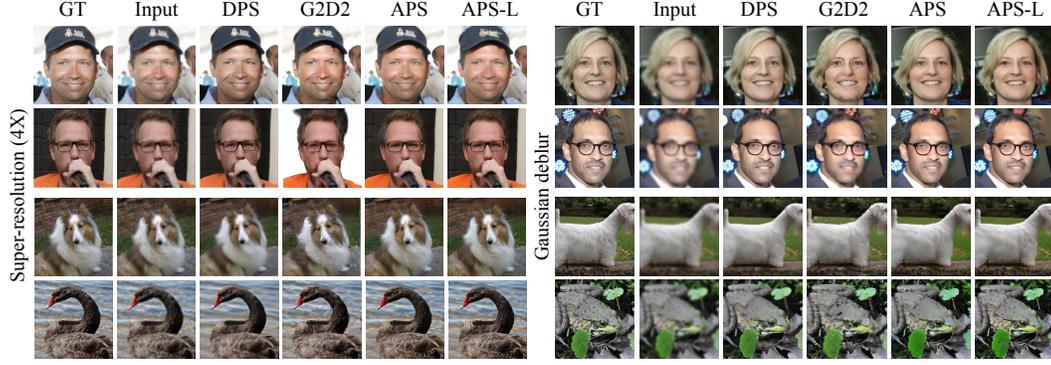
Figure 2: **Qualitative results on FFHQ and ImageNet for SR** ($4\times$) **and Gaussian deblur.** Compared to DPS and G2D2, APS yields better results with sharper texture and refined facial features. For instance, in the third row, APS reconstructs fine strands of the white and brown dog's fur.
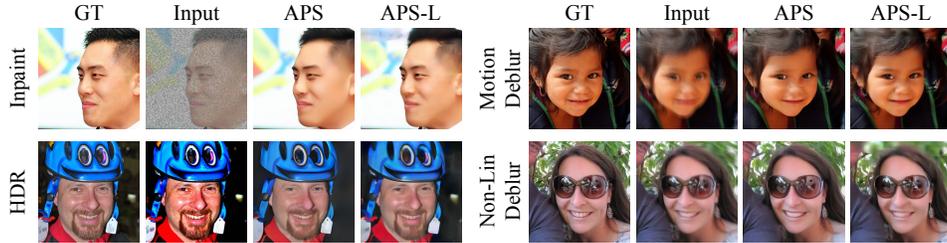

Figure 3: **Qualitative results on FFHQ** for linear (top row) and nonlinear (bottom row) inverse problems. APS and APS-L recover high-fidelity images from severely degraded inputs.

et al., 2017) to obtain an image $\hat{\mathbf{x}} = \mathcal{D}(\tilde{\mathbf{x}})$, $\tilde{\mathbf{x}} = \bar{\mathbf{x}} + \left[\mathbf{x} - \bar{\mathbf{x}}\right]_{\text{sg}}$, where sg denotes the stop-gradient operator. Finally, we compute the differentiable likelihood as $q(\mathbf{y}|\tilde{\mathbf{x}}_\varphi(\mathbf{z}_t)) \propto \exp(-\frac{\|\mathbf{y} - \mathcal{A}(\hat{\mathbf{x}})\|_2^2}{2\sigma^2})$.

Tokenwise expectation to differentiable surrogate. Define the measurement term in $\mathcal{L}_t(\varphi)$ as

$$\mathcal{L}_t^{\text{meas}} := \mathbb{E}_{q(Z_s|\mathbf{z}_t, \mathbf{x})}\Big[\sum_{l=1}^{L} -\log q\big(\mathbf{y} \mid \mathbf{x}_\varphi(\mathbf{z}_t; Z_s^l)\big)\Big] = \sum_{l=1}^{L} \mathbb{E}_{q(Z_s^l|\mathbf{z}_t^l, \mathbf{x})}\Big[-\log q\big(\mathbf{y} \mid \mathbf{x}_\varphi(\mathbf{z}_t; Z_s^l)\big)\Big],$$

where the last equality is due to $q(Z_s|\mathbf{z}_t, \mathbf{x})$ factorizing across sites given $\mathbf{z}_t$ (the forward process is conditionally independent tokenwise). Each inner expectation depends on the *local* categorical $q(Z_s^l|\mathbf{z}_t^l, \mathbf{x})$, which—under the mask process—has support on only two outcomes (e.g., "keep" vs. "mask"). Evaluating $\mathcal{L}_t^{\text{meas}}$ exactly would still require propagating each choice through $\mathcal{D}$ and $\mathcal{A}$, which is expensive when done for all l. We adopt a plug-in approximation that replaces the token-wise expectation by a single differentiable surrogate. Let $h(\mathbf{x}) := -\log q(\mathbf{y}|\mathbf{x})$ and $h(\mathbf{x}_\varphi(\mathbf{z}_t; \mathbf{z}_s^l))$ be defined as in §3.2. Then, a first-order Taylor's approximation yields

$$\mathbb{E}_{Z_s^l \sim q(\cdot|\mathbf{z}_t^l, \mathbf{x})}\big[h\big(\mathbf{x}_\varphi(\mathbf{z}_t; Z_s^l)\big)\big] \approx h\Big(\mathbb{E}_{Z_s^l \sim q(\cdot|\mathbf{z}_t^l, \mathbf{x})}\mathbb{E}_{\substack{X \sim \mathbf{x}_\varphi(\mathbf{z}_t), \\ X^l \leftarrow Z_s^l}}\big[X\big]\Big).$$

The double expectation inside $h(\cdot)$ scales linearly in sequence length (L) and vocabulary size (K). With the outer sum, the total complexity of $\mathcal{L}_t^{\text{meas}}$ becomes $\mathcal{O}(KL^2)$.

Next, we discuss how to reduce the complexity from $\mathcal{O}(KL^2)$ to $\mathcal{O}(KL)$. Note that $\mathbf{x}_\varphi(\mathbf{z}_t; Z_s^l)$ differs from $\mathbf{x}_\varphi(\mathbf{z}_t)$ only at position l. Assuming local smoothness of $\mathcal{A}\circ\mathcal{D}$, we approximate $q(\cdot|\mathbf{z}_t^l, \mathbf{x})$ by $\mathbf{x}_\varphi^l(\mathbf{z}_t)$ resulting in the *quantized expectation*: for each site, form the expected codebook embedding $\bar{\mathbf{x}}^l = \sum_k \mathbf{c}_k \mathbf{x}_\varphi^l(\mathbf{z}_t)$, quantize via LFQ as $\mathbf{x}^l = \mathcal{Q}_{\text{lfq}}(\bar{\mathbf{x}}^l)$, and apply the straight-through estimator to obtain a single surrogate sequence $\tilde{\mathbf{x}}_\varphi(\mathbf{z}_t) = \bar{\mathbf{x}} + [\mathbf{x} - \bar{\mathbf{x}}]_{\text{sg}}$. Substituting the same

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surrogate for all l yields the following[2]:

$$\mathcal{L}_t^{\text{meas}} \approx \sum_{l=1}^{L} h(\tilde{\mathbf{x}}_\varphi(\mathbf{z}_t)) = L \cdot \left(-\log q(\mathbf{y}|\tilde{\mathbf{x}}_\varphi(\mathbf{z}_t)) \right).$$

Discussion. We minimize (7) at each step t to obtain the optimal logits φ_t^*. This ensures that the optimized probabilities $\mathbf{x}_{\varphi^*}(\mathbf{z}_t)$ remain close to the prior predictions $\mathbf{x}_\theta(\mathbf{z}_t)$ while aligning with measurements $\mathbf{y}$. To our knowledge, this is the first use of *quantized expected codebook embeddings* for posterior sampling in discrete diffusion, allowing measurement gradients to backpropagate through $\bar{\mathbf{x}}$ and update all entries of $\mathbf{x}_\varphi(\mathbf{z}_t)$. Our quantized expectation reduces the cost for the measurement term from $\mathcal{O}(K^L)$ to $\mathcal{O}(KL)$ because it requires only *one* forward pass through $\mathcal{A} \circ \mathcal{D}$ using $\tilde{\mathbf{x}}_\varphi(\mathbf{z}_t)$.

3.2.2 ANCHORED REMASKING

In masked diffusion, the reverse process progressively unmasks tokens. Most discrete samplers (Austin et al., 2021; Chang et al., 2022; Lou et al., 2024; Sahoo et al., 2024; Shi et al., 2024) choose to unmask tokens in random order. ADLM (Rout et al., 2025a) shows that prioritizing "anchor" tokens (e.g., nouns or verbs in language, rather than articles or conjunctions) early in the reverse process reduces conditional entropy of the remaining sequence and improves generation quality. To decode anchor tokens, ADLM jointly trains an anchor network in addition to the standard denoising network using an anchored NELBO objective. Here we propose a *training-free* variant of anchored denoising, enabling posterior sampling with pretrained masked diffusion models.

Let φ be a minimizer of (7), resulting in $\mathbf{x}_\varphi^*(\mathbf{z}_t) = \{(\mathbf{x}_\varphi^*(\mathbf{z}_t))^l\}_{l=1}^{L}$ that represents categorical distributions over tokens at each position $l \in \{1, \dots, L\}$ at time step t. Anchored remasking selects a subset of positions $\mathcal{P}_t \subseteq \{1, \dots, L\}$ to decode early. The selection is based on the confidence of quantized tokens $\mathbf{x}$ (as defined in §3.2.1) under the posterior estimate $\mathbf{x}_\varphi^*(\mathbf{z}_t)$. Importantly, the posterior estimate is a function of the L-length sequence $\mathbf{z}_t$, and hence encodes the *joint* relation across all tokens; thus, anchored remasking is a function of *all* tokens, as compared to standard remasking (Chang et al., 2022; Yang et al., 2025) that depend only on per-token confidence. Formally, we compute the confidence of $\mathbf{x}^l$ as $\kappa_t^l = \langle (\mathbf{x}_\varphi^*(\mathbf{z}_t))^l, \mathbf{x}^l \rangle$, and choose anchor positions as $\mathcal{P}_t = \{l : \kappa_t^l \geq \tau_t\}$, where τ_t is an adaptive threshold that follows the cosine schedule from MMaDA. We then update the state by fixing anchor tokens and remasking the rest: $\mathbf{z}_s^l = \mathbf{x}$ with probability $\frac{\alpha_s - \alpha_t}{1 - \alpha_t}$ if $l \in \mathcal{P}_t$ else $\mathbf{m}$. Once a token is unmasked, it remains fixed in subsequent steps.

Discussion. Diffusion language models tend to be overconfident on low-information tokens such as articles ("a", "an" or "the") or conjunctions (Rout et al., 2025a); similarly, discrete image samplers using *independent per-token* confidence often unmask background pixels first. In contrast, our method leverages the *joint posterior* $\mathbf{x}_\varphi(\mathbf{z}_t)$ to identify anchor tokens consistent with the measurements. This leads to earlier decoding of informative tokens (e.g., a bird against a flat background), enabling faster reconstruction and improved likelihood of generated samples; refer §B.3 for details.

4 EXPERIMENTS

Baselines. Since our focus is on discrete diffusion, we first compare against existing discrete posterior samplers: G2D2 (Murata et al., 2024) and SGDD (Chu et al., 2025) (§2). To provide a comprehensive evaluation, we also include established continuous baselines, both in pixel space (DPS (Chung et al., 2023), DDRM (Kawar et al., 2022), and DiffPIR (Zhu et al., 2023)) and in latent space (PSLD (Rout et al., 2023) and ReSample (Song et al., 2024)). This ensures our evaluation spans both discrete and continuous posterior sampling paradigms.

Benchmarks. We evaluate on the standard inverse problem benchmarks used in prior works. For high-resolution faces we use FFHQ at 256×256 (Karras et al., 2019), and for diverse natural images we use ImageNet at 256×256 (Deng et al., 2009). Performance is measured using three standard metrics: Learned Perceptual Image Patch Similarity (Zhang et al., 2018) (LPIPS), Peak Signal-to-Noise Ratio (PSNR), and Structural Similarity Index (SSIM) (Wang et al., 2004). All methods are evaluated at the same resolution and on the same images following G2D2 (Murata et al., 2024).

Tasks. We consider both linear and nonlinear inverse problems. Following prior works (e.g., G2D2 (Murata et al., 2024) and SGDD (Chu et al., 2025)), we evaluate on *Super Resolution (SR)*

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[2]In practice, the factor L can be absorbed into the likelihood weight or learning rate. If only positions with $\mathbf{z}_t^l = \mathbf{m}$ contribute, one may replace L by $|\{l : \mathbf{z}_t^l = \mathbf{m}\}|$. We keep the absorbed constant here for simplicity.

Table 2: **Quantitative results on general inverse problems.** We report results on two additional linear (random inpainting, motion deblurring) and nonlinear (HDR, nonlinear blur) tasks. Since G2D2 and SGDD do not evaluate on these tasks, we compare our *discrete* sampler against representative *continuous* baselines: DPS (pixel-space) and PSLD (latent-space)

Type	Method	Random Inpainting		Motion Deblur		HDR		Nonlinear Blur	
		LPIPS ↓	PSNR ↑	LPIPS ↓	PSNR ↑	LPIPS ↓	PSNR ↑	LPIPS ↓	PSNR ↑
					FFHQ				
Pixel	DPS	**0.203**	25.46	**0.246**	24.52	**0.264**	22.73	0.278	23.39
Latent	PSLD	0.221	**30.31**	0.336	22.31	–	–	–	–
Mask	**APS (ours)**	0.304	27.38	0.317	26.58	0.282	**23.89**	0.263	27.19
Discrete	**APS-L (ours)**	0.291	28.11	0.298	**27.98**	0.323	23.56	**0.262**	**28.46**
					ImageNet				
Pixel	DPS	**0.297**	23.52	0.423	18.96	0.503	19.23	**0.306**	22.49
Latent	PSLD	0.337	**31.30**	0.511	20.85	–	–	–	–
Mask	**APS (ours)**	0.378	24.59	0.410	23.37	0.345	21.92	0.330	24.18
Discrete	**APS-L (ours)**	0.338	25.39	**0.318**	25.19	0.346	**22.68**	0.309	**25.35**

| Reference Style | | Stylized Output | | Reference Style | | Stylized Output | |

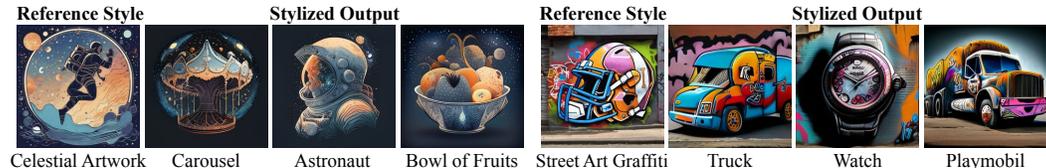Celestial Artwork Carousel Astronaut Bowl of Fruits Street Art Graffiti Truck Watch Playmobil

Figure 4: **Qualitative results on stylization.** We present four style–content combinations. For each case, our APS algorithm conditions on a single reference style image together with a text prompt.

(4×) and *Gaussian deblurring* on both FFHQ and ImageNet. Additionally, we evaluate on more linear (*random inpainting* and *motion blur*) and nonlinear (*high dynamic range (HDR) recovery* and *nonlinear deblurring*) tasks. Beyond inverse problems, we conduct experiments on *training-free stylization*, an emerging area largely dominated by continuous diffusion (Hertz et al., 2023; Wang et al., 2024; Rout et al., 2025c). To demonstrate scalability, we upsample the benchmark images to 1024×1024 and apply our posterior sampler to these scaled images (termed **APS-L** in discussion below). We defer implementation details to §B.1 and computational complexity to §B.5.

4.1 RESULTS ON LINEAR INVERSE PROBLEMS

Evaluation on FFHQ: Table 1 (a) shows that our approach outperforms prior discrete diffusion samplers across both super resolution and Gaussian deblurring tasks. For *super resolution*, APS reduces LPIPS by 13.65% and improves PSNR by 2.11%. On *Gaussian deblurring*, APS lowers LPIPS by 3.83% compared to G2D2 and raises PSNR by 5.88%. The large variant, APS-L, pushes performance even further, achieving up to a 31.36% gain over G2D2 in terms of LPIPS. These results show that our quantized expectation (§3.2.1) and anchored remasking (§3.2.2) strategies not only outperform discrete diffusion baselines but often surpass strong continuous-diffusion methods, such as DiffPIR (which uses pixel-space diffusion) and PSLD (which uses latent-space diffusion), resulting in higher perceptual and reconstruction quality.

Figure 2 (top two rows) presents qualitative comparisons for super resolution $(4\times)$ and Gaussian deblurring on FFHQ. DPS produces over-smoothed results with blurry facial details, while G2D2 often introduces artifacts and fails to restore a natural facial structure. Our APS sampler yields sharper textures and more faithful reconstructions, recovering details such as hair strands, facial contours, and eyeglass edges with higher perceptual quality. The large variant, APS-L, further enhances structure and realism, generating photo-realistic outputs with finer details and fewer artifacts. These results confirm that APS and APS-L deliver superior qualitative performance on face datasets, where perceptual fidelity is especially critical.

Evaluation on ImageNet: Table 1 (b) quantifies that our method achieves consistent improvements across both SR and Gaussian deblurring tasks on the ImageNet benchmark. For *super resolution*, APS reduces LPIPS by 7.16% compared to G2D2 and improves PSNR by 4.74%. On *Gaussian deblurring*, APS improves PSNR by 8.81% while maintaining comparable LPIPS. APS-L improves

LPIPS by up to 35.82% compared to G2D2. These results confirm that APS outperforms prior discrete posterior samplers and often surpasses continuous baselines such as DiffPIR and PSLD, delivering superior perceptual quality and reconstruction fidelity.

Figure 2 (bottom two rows) shows SR ($4\times$) and Gaussian deblurring results on ImageNet. Notably, DPS produces overly smooth outputs with a loss of fine details, while G2D2 introduces struggles to recover sharp edges. In contrast, APS reconstructs sharper textures (e.g., the fur of the dog and the feathers of the swan) and yields more natural color. The large variant, APS-L, further enhances structural fidelity, recovering finer details in challenging regions such as a frog's skin texture and a goat's fur. These examples highlight that our approach achieves superior perceptual quality and faithful structure reconstruction compared to both pixel-based and discrete diffusion baselines.

4.2 RESULTS ON GENERAL INVERSE PROBLEMS

Table 2 shows that APS generalizes effectively to more challenging linear (random inpainting, motion deblurring) and nonlinear (HDR, nonlinear blur) inverse problems on FFHQ and ImageNet. Unlike prior discrete samplers such as G2D2 and SGDD, which were demonstrated only on limited tasks, APS consistently achieves strong perceptual quality (lower LPIPS) and reconstruction fidelity (higher PSNR). For instance, on ImageNet motion deblurring APS-L attains 0.318 LPIPS and 25.19 PSNR, substantially outperforming continuous baselines DPS and PSLD. Similarly, in nonlinear tasks such as HDR and nonlinear blur, APS delivers sharper, more consistent reconstructions, closing the gap with continuous diffusion while operating within a purely discrete framework. These results highlight the broader applicability and robustness of our approach compared to existing discrete diffusion samplers. Figure 3 shows the qualitative results of general inverse problems.

4.3 RESULTS ON REFERENCE-BASED STYLIZATION

We compare APS with the discrete diffusion baseline MMaDA (Yang et al., 2025) and continuous methods (shaded gray). Table 3 reports quantitative results, while Figure 4 shows qualitative examples. APS improves over MMaDA on ImageReward (Xu et al., 2024), CLIP-T (Radford et al., 2021), and DINO (Caron et al., 2021) scores, demonstrating that our posterior sampler unlocks stylization capabilities absent in the base model. Notably, APS also

	ImageReward ↑	CLIP-T ↑	DINO ↑
IP-Adapter	-1.51	0.26	0.89
StyleAligned	0.01	0.31	0.85
InstantStyle	0.72	0.33	0.72
RB-Modulation	1.18	0.34	0.73
MMaDA	0.48	0.33	0.32
APS (ours)	0.63	0.34	0.41

Table 3: **Quantitative results on stylization.**

surpasses continuous baselines such as IP-Adapter (Ye et al., 2023) and StyleAligned (Hertz et al., 2023) on ImageReward and CLIP-T, despite relying on a weaker generative prior.

The appendix provides extensive details on our Anchored Posterior Sampling method. We present the theoretical derivation of variational bounds (§A), all implementation specifics and hyperparameters (Algorithm 1, Table 4), and a detailed ablation study (§B.3, Figure 5) demonstrating the impact of our innovations. We also showcase the superior computational efficiency (Table 7) and numerous additional qualitative results (§B) across complex inverse problems, high-resolution text-guided block inpainting (Figure 11), and extended stylization results (Figure 12 and 13).

5 CONCLUSION

We introduce **Anchored Posterior Sampling (APS)**, a training-free posterior sampler with discrete diffusion foundation models. Grounded in our theoretical bounds, APS enables the reuse of a pretrained denoiser *without* task-specific retraining. APS is built on two algorithmic innovations: *Quantized Expectation*, which provides fine-grained, gradient-like guidance in discrete spaces, and *Anchored Remasking*, which adaptively decodes important tokens early in the reverse process. Together, these yield an efficient, scalable, and training-free posterior sampler.

Extensive experiments on linear and nonlinear inverse problems demonstrate that APS achieves state-of-the-art results among discrete samplers and is competitive with continuous baselines—while operating at significantly lower inference cost. Beyond inverse problems, APS also unlocks new capabilities such as training-free stylization, underscoring the flexibility of discrete diffusion models when paired with effective posterior sampling. We believe this work establishes discrete diffusion as a practical and scalable alternative for posterior sampling, with promising extensions to video, multimodal generation, and other structured domains.

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A ADDITIONAL THEORETICAL RESULTS

This appendix develops complementary theory for masked discrete diffusion posterior sampling. We first derive a pathwise variational bound for *training* a likelihood-tilted reverse process (**Theorem A.1**), showing that $-\log p_\varphi(\mathbf{x}|\mathbf{y})$ is upper bounded by a reconstruction term, a sum of token-wise KL-divergence terms, and a sequence of measurement likelihood-based tilting terms. We then specialize this analysis to the *training-free* setting where the token-to-image decoder is shared between unconditional generation and posterior sampling, yielding a bound for test-time anchored posterior sampling (**Theorem A.2**). We provide theoretical insights drawn from each theorem in **Implication** subsections after the corresponding proofs.

Theorem A.1 (Discrete Diffusion Posterior Sampling(DDPS)). *Let $Z_{0:1} = \{Z_{t(i)}\}_{i=0}^{T}$ with $t(i) = i/T$ and $s(i) = (i-1)/T$ be the latent path of a masked discrete diffusion model, and let $q(Z_{0:1}|\mathbf{x})$ be the forward noising law from (3). Consider reverse kernels and a terminal decoder that factorize as*

$$p_\varphi(\mathbf{x}, Z_{0:1}|\mathbf{y}) = p_\varphi(Z_1|\mathbf{y})\, p_\varphi(\mathbf{x}|Z_0, \mathbf{y}) \prod_{i=1}^{T} p_\varphi\big(Z_{s(i)}|Z_{t(i)}, \mathbf{y}\big),$$

with token-wise reverse transitions given by the inference posterior in (4) tilted by the likelihood,

$$p_\varphi(Z_s^l|Z_t, \mathbf{y}) \propto q\big(Z_s^l|Z_t^l, \mathbf{x}_\varphi(Z_t)\big)\, q\big(\mathbf{y}|\mathbf{x}_\varphi(Z_t; Z_s)\big).$$

Then, for any $(\mathbf{x}, \mathbf{y})$,

$$-\log p_\varphi(\mathbf{x}|\mathbf{y}) \leq \mathcal{L}_{\mathrm{DDPS}}(\mathbf{x}, \mathbf{y}; \varphi) := \mathbb{E}_{q(Z_0|\mathbf{x})}[-\log p_\varphi(\mathbf{x}|Z_0)]$$

$$+ \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})} \Bigg[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \sum_{l=1}^{L} \mathrm{CE}\big(\mathbf{x}^l, \mathbf{x}_\varphi^l(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)}^l = \mathbf{m}\}}$$

$$- \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})} \Big[\sum_{l=1}^{L} \log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}^l)) \Big]$$

$$+ \log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y}) \Bigg]. \tag{8}$$

Proof. We present the derivation for sequence length $L = 1$; the extension to $L > 1$ follows by token-wise factorization as in (Sohl-Dickstein et al., 2015). Recall our parameterized reverse kernel (identical in form to (4) up to a likelihood tilt):

$$q(Z_s^l|Z_t^l, \mathbf{x}_\varphi(Z_t)) = \begin{cases} \mathrm{Cat}(Z_s^l; Z_t^l), & Z_t^l \neq \mathbf{m}, \\ \mathrm{Cat}\big(Z_s^l; \frac{\alpha_s - \alpha_t}{1 - \alpha_t}\mathbf{x}_\varphi(Z_t) + \frac{1 - \alpha_s}{1 - \alpha_t}\mathbf{m}\big), & Z_t^l = \mathbf{m}. \end{cases} \tag{9}$$

Starting from the conditional likelihood,

$$-\log p_\varphi(\mathbf{x}|\mathbf{y}) = -\log \int p_\varphi(\mathbf{x}, Z_{0:1}|\mathbf{y})\, \mathrm{d}Z_{0:1}$$

$$= -\log \int p_\varphi(\mathbf{x}, Z_{0:1}|\mathbf{y}) \frac{q(Z_{0:1}|\mathbf{x}, \mathbf{y})}{q(Z_{0:1}|\mathbf{x}, \mathbf{y})}\, \mathrm{d}Z_{0:1}.$$

By the conditional independence of the forward process (the noising path is conditionally independent of $\mathbf{y}$ given $\mathbf{x}$), we have $q(Z_{0:1}|\mathbf{x}, \mathbf{y}) = q(Z_{0:1}|\mathbf{x})$, hence

$$-\log p_\varphi(\mathbf{x}|\mathbf{y}) = -\log \mathbb{E}_{q(Z_{0:1}|\mathbf{x})} \left[\frac{p_\varphi(\mathbf{x}, Z_{0:1}|\mathbf{y})}{q(Z_{0:1}|\mathbf{x})} \right].$$

Applying Jensen's inequality,

$$-\log p_\varphi(\mathbf{x}|\mathbf{y}) \leq \mathbb{E}_{q(Z_{0:1}|\mathbf{x})} \left[-\log \frac{p_\varphi(\mathbf{x}, Z_{0:1}|\mathbf{y})}{q(Z_{0:1}|\mathbf{x})} \right].$$

Using the factorization $p_\varphi(\mathbf{x}, Z_{0:1}|\mathbf{y}) = p_\varphi(Z_1|\mathbf{y}) \, p_\varphi(\mathbf{x}|Z_0, \mathbf{y}) \prod_{i=1}^T p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})$ and the forward process factorization $q(Z_{0:1}|\mathbf{x}) = q(Z_1|\mathbf{x}) \prod_{i=1}^T q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})$, we obtain

$$- \log \frac{p_\varphi(\mathbf{x}, Z_{0:1}|\mathbf{y})}{q(Z_{0:1}|\mathbf{x})}$$

$$= - \log p_\varphi(\mathbf{x}|Z_0, \mathbf{y}) + \log \frac{q(Z_1|\mathbf{x})}{p_\varphi(Z_1|\mathbf{y})} + \sum_{i=1}^T \log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})}.$$

Taking expectation under $q(Z_{0:1}|\mathbf{x})$ yields

$$- \log p_\varphi(\mathbf{x}|\mathbf{y}) \le \mathbb{E}_{q(Z_{0:1}|\mathbf{x})}\big[- \log p_\varphi(\mathbf{x}|Z_0, \mathbf{y})\big] + \mathbb{E}_{q(Z_1|\mathbf{x})}\Big[\log \frac{q(Z_1|\mathbf{x})}{p_\varphi(Z_1|\mathbf{y})}\Big]$$

$$+ \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})} \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\Big[\log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})}\Big]. \tag{10}$$

Expanding the tilted reverse kernel using the parameterization:

$$p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y}) = \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)})) \, q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))}{\mathcal{Z}(Z_{t(i)}, \mathbf{y})}, \quad \text{where}$$

$$\mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y}) = \sum_{\mathbf{z}_s \in \mathcal{V}} q\big(\mathbf{z}_s|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)})\big) \, q\big(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; \mathbf{z}_s)\big),$$

we can rewrite the last term as

$$\sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})} \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\Big[\log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))} - \log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)})) + \log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y})\Big].$$

This yields the final decomposition:

$$- \log p_\varphi(\mathbf{x}|\mathbf{y}) \le \underbrace{\mathbb{E}_{q(Z_0|\mathbf{x})}[- \log p_\varphi(\mathbf{x}|Z_0, \mathbf{y})] + \mathrm{KL}\big(q(Z_1|\mathbf{x}) \,\|\, p_\varphi(Z_1|\mathbf{y})\big)}_{\text{reconstruction + boundary KL}}$$

$$+ \underbrace{\sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\Big[\mathrm{KL}\big(q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))\big)\Big]}_{\text{per-step KLs}}$$

$$- \underbrace{\sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}\big[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))\big]}_{\text{likelihood tilt terms}}$$

$$+ \underbrace{\sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\big[\log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y})\big]}_{\text{normalizing factor}} \coloneqq \mathcal{L}_{\mathrm{DDPS}}(\mathbf{x}, \mathbf{y}; \varphi).$$

The first two groups of terms correspond to the standard masked diffusion posterior NELBO objective (Sohl-Dickstein et al., 2015; Austin et al., 2021) denoted by:

$$\mathcal{L}_{\mathrm{PNELBO}}(\mathbf{x}, \mathbf{y}; \varphi) \coloneqq \mathbb{E}_{q(Z_0|\mathbf{x})}\big[- \log p_\varphi(\mathbf{x}|Z_0, \mathbf{y})\big]$$

$$+ \mathrm{KL}\big(q(Z_1|\mathbf{x}) \,\|\, p_\varphi(Z_1|\mathbf{y})\big)$$

$$+ \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\Big[\mathrm{KL}\big(q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))\big)\Big].$$

The likelihood term,

$$- \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}\big[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))\big],$$

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capture the effect of incorporating observations $\mathbf{y}$ into posterior sampling. Finally, we have an additive normalizing term that ensures the validity of the reverse transition (6).

We now bound the per-step KL terms. Having connected the decomposition to the NELBO, it suffices to compute, for each step i, the divergence

$$\mathrm{KL}\big(q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))\big).$$

Since masked diffusion induces a two–state posterior at each step (either remain masked or reveal the data token), we treat the two cases for $Z_{t(i)}$ separately.

Case I: $Z_{t(i)} = \mathbf{m}.$ From the masked diffusion forward process (3), when $Z_{t(i)} = \mathbf{m}$ the true posterior over $Z_{s(i)}$ has mass

$$q(Z_{s(i)} = \mathbf{m}|Z_{t(i)} = \mathbf{m}, \mathbf{x}) = \frac{1 - \alpha_{s(i)}}{1 - \alpha_{t(i)}}, \qquad q(Z_{s(i)} = \mathbf{x}|Z_{t(i)} = \mathbf{m}, \mathbf{x}) = \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}}.$$

Under the model with prediction $\mathbf{x}_\varphi(Z_{t(i)})$ (a categorical distribution) as in (4), the "unmask" branch is weighted by the model's probability of the true token, $\langle \mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x}\rangle$, while the mask probability is unchanged. Hence,

$$\mathrm{KL}\big(q(Z_{s(i)}|Z_{t(i)} = \mathbf{m}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)} = \mathbf{m}, \mathbf{x}_\varphi(Z_{t(i)}))\big)$$

$$= \frac{1 - \alpha_{s(i)}}{1 - \alpha_{t(i)}} \log \frac{\frac{1-\alpha_{s(i)}}{1-\alpha_{t(i)}}}{\frac{1-\alpha_{s(i)}}{1-\alpha_{t(i)}}} + \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \log \frac{\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}}{\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}} \langle \mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x}\rangle}$$

$$= \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \log \frac{1}{\langle \mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x}\rangle} = \frac{\alpha_{t(i)} - \alpha_{s(i)}}{1 - \alpha_{t(i)}} \log \langle \mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x}\rangle.$$

Equivalently, writing the cross-entropy with the one-hot target $\mathbf{x}$ as $\mathrm{CE}(\mathbf{x}, \mathbf{x}_\varphi(Z_{t(i)})) = -\log\langle \mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x}\rangle$,

$$\mathrm{KL}\big(q(Z_{s(i)}|Z_{t(i)} = \mathbf{m}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)} = \mathbf{m}, \mathbf{x}_\varphi(Z_{t(i)}))\big) = \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \mathrm{CE}(\mathbf{x}, \mathbf{x}_\varphi(Z_{t(i)})).$$

Case II: $Z_{t(i)} \neq \mathbf{m}.$ When the current token is already unmasked, the posterior is deterministic: $q(Z_{s(i)} = Z_{t(i)}|Z_{t(i)} \neq \mathbf{m}, \mathbf{x}) = 1$. Thus,

$$\mathrm{KL}\big(q(Z_{s(i)}|Z_{t(i)} \neq \mathbf{m}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)} \neq \mathbf{m}, \mathbf{x}_\varphi(Z_{t(i)}))\big) = 0.$$

Only masked coordinates contribute to the per-step KL, yielding for each step i,

$$\mathrm{KL}\big(q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))\big) = \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \mathrm{CE}\big(\mathbf{x}, \mathbf{x}_\varphi(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)} = \mathbf{m}\}}.$$

This recovers the standard masked-diffusion NELBO weighting (cf. (5)): per-step contributions are cross-entropies at masked positions, scaled by $(\alpha_{s(i)} - \alpha_{t(i)})/(1 - \alpha_{t(i)})$. Generalizing this to sequences with length $L > 1$ yields:

$$\mathrm{KL}\big(q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))\big) = \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \sum_{l=1}^{L} \mathrm{CE}\big(\mathbf{x}^l, \mathbf{x}_\varphi^l(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)}^l = \mathbf{m}\}}.$$

Combining this with the likelihood tilt terms, we get

$$\mathcal{L}_{\mathrm{DDPS}}(\mathbf{x}, \mathbf{y}; \varphi) = \mathbb{E}_{q(Z_0|\mathbf{x})}[-\log p_\varphi(\mathbf{x}|Z_0, \mathbf{y})] + \mathrm{KL}\big(q(Z_1|\mathbf{x}) \,\|\, p_\varphi(Z_1|\mathbf{y})\big)$$

$$+ \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \sum_{l=1}^{L} \mathrm{CE}\big(\mathbf{x}^l, \mathbf{x}_\varphi^l(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)}^l = \mathbf{m}\}}\right]$$

$$- \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}\big[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))\big]$$

$$+ \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\big[\log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y})\big].$$

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In masked diffusion Z_1 is typically the fully masked state or absorbing state, so $q(Z_1|\mathbf{x})$ is degenerate and, with $p_\varphi(Z_1|\mathbf{y}) = q(Z_1|\mathbf{x})$, the boundary KL vanishes. Thus,

$$
\begin{aligned}
\mathcal{L}_{\mathrm{DDPS}}(\mathbf{x}, \mathbf{y}; \varphi) = {}& \mathbb{E}_{q(Z_0|\mathbf{x})}[-\log p_\varphi(\mathbf{x}|Z_0, \mathbf{y})] \\
& + \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \sum_{l=1}^{L} \mathrm{CE}\big(\mathbf{x}^l, \mathbf{x}_\varphi^l(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)}^l = \mathbf{m}\}} \right] \\
& - \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}\big[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)})) \big] \\
& + \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\big[\log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y}) \big].
\end{aligned}
$$

Furthermore, $\mathbf{y}$ imposes a distribution over $\mathbf{x}$ from which we wish to sample. However, when Z_0 is given then $\mathbf{x}$ is uniquely determined by the decoder as $\mathbf{x} = \mathrm{Dec}(Z_0)$. Therefore,

$$
\begin{aligned}
\mathcal{L}_{\mathrm{DDPS}}(\mathbf{x}, \mathbf{y}; \varphi) = {}& \mathbb{E}_{q(Z_0|\mathbf{x})}[-\log p_\varphi(\mathbf{x}|Z_0)] \\
& + \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \sum_{l=1}^{L} \mathrm{CE}\big(\mathbf{x}^l, \mathbf{x}_\varphi^l(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)}^l = \mathbf{m}\}} \right] \\
& - \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}\big[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)})) \big] \\
& + \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\big[\log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y}) \big],
\end{aligned}
$$

which upon regrouping completes the proof of the statement. $\qquad\square$

Implications. Theorem A.1 provides a principled upper bound on the negative log-posterior likelihood in discrete masked diffusion models.

- *Trainable upper bound.* The pathwise upper bound $\mathcal{L}_{\mathrm{DDPS}}(\mathbf{x}, \mathbf{y}; \varphi)$ in (8) provides a principled training criterion for discrete diffusion posterior sampling.

- *Reduction to standard training when $\mathbf{y}$ is absent.* Setting the likelihood to a constant (no measurements) removes the tilt terms and recovers the masked-diffusion training objective: only the reconstruction term and per-step KL-divergence terms remain.

- *Masked-token supervision.* The per-step KL-divergence terms vanish on already-revealed tokens and reduce to weighted cross-entropies on masked tokens, focusing learning signal exactly where denoising must occur. The weights $(\alpha_{s(i)} - \alpha_{t(i)})/(1 - \alpha_{t(i)})$ expose how the noise schedule shapes gradient magnitude.

- *Data-consistency via tilt.* The additive terms $-\sum_i \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))]$ encourage reverse transitions that produce intermediate predictions consistent with the measurement model, integrating task specific information at every denoising step.

- *Boundary conditions.* With an absorbing mask state, the boundary KL-divergence is constant (often zero), so optimization concentrates on reconstruction, token-level KL-divergence terms, and measurement consistency.

- *Compatibility with efficient parameterizations.* Because (8) is written in terms of token-wise categoricals, it directly supports time-independent or lightweight parameterizations (e.g., shared denoisers), helping scalability to long sequences and high resolution.

Theorem A.2 (Test-time Anchored Posterior Sampling). *Let $Z_{0:1} = \{Z_{t(i)}\}_{i=0}^{T}$ with $t(i) = i/T$ and $s(i) = (i-1)/T$ denote the latent path of a masked discrete diffusion model, and let $q(Z_{0:1}|\mathbf{x})$ be the forward noising law from (3). Assume the decoder is shared between unconditional generation and posterior sampling ($p_\varphi(\mathbf{x}|Z_0) = p_\theta(\mathbf{x}|Z_0)$), and the unconditional reverse transitions are parameterized as in (4). Define $\mathcal{L}_{\mathrm{NELBO}}(\mathbf{x}; \theta)$ as in (5). Then, for any $(\mathbf{x}, \mathbf{y})$,*

$$
-\log p_\varphi(\mathbf{x}|\mathbf{y}) \ \le\ \mathcal{L}_{\mathrm{APS}}(\mathbf{x}, \mathbf{y}; \varphi),
$$

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where

$$\mathcal{L}_{\text{APS}}(\mathbf{x}, \mathbf{y}; \varphi) := \mathcal{L}_{\text{NELBO}}(\mathbf{x}; \theta) + \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \log \frac{\langle \mathbf{x}_\theta(Z_{t(i)}), \mathbf{x} \rangle}{\langle \mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x} \rangle} \mathbf{1}_{\{Z_{t(i)} = \mathbf{m}\}} \right.$$

$$- \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)})) \right]$$

$$\left. + \log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y}) \right].$$

Proof. From the proof of **Theorem A.1**, the conditional likelihood satisfies

$$- \log p_\varphi(\mathbf{x}|\mathbf{y}) \leq \mathbb{E}_{q(Z_0|\mathbf{x})}\left[-\log p_\varphi(\mathbf{x}|Z_0, \mathbf{y}) \right] + \mathbb{E}_{q(Z_1|\mathbf{x})}\left[\log \frac{q(Z_1|\mathbf{x})}{p_\varphi(Z_1|\mathbf{y})} \right]$$

$$+ \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})} \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})} \right]. \tag{11}$$

Compute each term inside summation. We now focus on the ith term inside the summation. Using the pretrained network $p_\theta(Z_{s(i)}|Z_{t(i)})$ as given in (4), we decompose the term as:

$$\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})} \right]$$

$$= \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}{p_\theta(Z_{s(i)}|Z_{t(i)})} + \log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})} \right].$$

Isolating the first-term as a KL-divergence term, this yields

$$\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})} \right]$$

$$= \underbrace{\text{KL}\Big(q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}) \,\|\, p_\theta(Z_{s(i)}|Z_{t(i)})\Big)}_{\text{divergence w.r.t. pretrained network}} + \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})} \right].$$

Since $p_\theta(Z_{s(i)}|Z_{t(i)})$ is parameterized via the pretrained network prediction $\mathbf{x}_\theta(Z_{t(i)})$, we get

$$\text{KL}\Big(q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}) \,\|\, p_\theta(Z_{s(i)}|Z_{t(i)})\Big) = \text{KL}\Big(q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\theta(Z_{t(i)}))\Big).$$

Following the argument of **Theorem A.1**, we distinguish two cases for $Z_{t(i)}$. For $Z_{t(i)} = \mathbf{m}$, the KL-divergence evaluates to

$$\text{KL}\Big(q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}) \,\|\, q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\theta(Z_{t(i)}))\Big) = \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \text{CE}\big(\mathbf{x}, \mathbf{x}_\theta(Z_{t(i)})\big),$$

while for $Z_{t(i)} \neq \mathbf{m}$ the KL-divergence vanishes. Thus we obtain

$$\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})} \right] \tag{12}$$

$$= \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \text{CE}\big(\mathbf{x}, \mathbf{x}_\theta(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)} = \mathbf{m}\}} + \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})} \right].$$

Substituting the above expression in the conditional likelihood (11), we get

$$- \log p_\varphi(\mathbf{x}|\mathbf{y}) \leq \mathbb{E}_{q(Z_0|\mathbf{x})}\left[-\log p_\varphi(\mathbf{x}|Z_0, \mathbf{y}) \right] + \mathbb{E}_{q(Z_1|\mathbf{x})}\left[\log \frac{q(Z_1|\mathbf{x})}{p_\varphi(Z_1|\mathbf{y})} \right]$$

$$+ \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \text{CE}\big(\mathbf{x}, \mathbf{x}_\theta(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)} = \mathbf{m}\}} \right]$$

$$+ \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})} \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})} \right]. \tag{13}$$

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Expected log-likelihood ratio under q**.** We now examine the expected difference in log-likelihoods between the prior and posterior transition distributions under the conditional law of the forward process. Recall that the reverse transition under our parameterization is

$$p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y}) = \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))\, q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))}{\mathcal{Z}(Z_{t(i)}, \mathbf{y})}, \quad \text{where}$$

$$\mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y}) = \sum_{\mathbf{z}_s \in \mathcal{V}} q(\mathbf{z}_s|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))\, q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; \mathbf{z}_s)).$$

Consider the expectation of the log-likelihood ratio under the forward posterior $q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})$:

$$\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})}\right].$$

Substituting the definition of p_φ, we obtain

$$\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})\mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y})}{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))\, q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))}\right]$$

$$= \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))} - \log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)})) + \log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y})\right]$$

$$= \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))}\right] - \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))\right]$$

$$+ \log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y}).$$

Since $p_\theta(Z_{s(i)}|Z_{t(i)})$ is represented by the network prediction $\mathbf{x}_\theta(Z_{t(i)})$, we can rewrite the first expectation as:

$$\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))}\right] = \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\theta(Z_{t(i)}))}{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))}\right].$$

Putting everything together, the expected log-likelihood ratio under q becomes

$$\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{p_\varphi(Z_{s(i)}|Z_{t(i)}, \mathbf{y})}\right]$$

$$= \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log \frac{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\theta(Z_{t(i)}))}{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x}_\varphi(Z_{t(i)}))}\right] - \mathbb{E}_{q(Z_{s(i)}|Z_{t(i)}, \mathbf{x})}\left[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))\right]$$

$$+ \log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y}). \tag{14}$$

Similarly to the proof of **Theorem A.1**, we consider two cases to compute the first expectation.

Case I: $Z_{t(i)} = \mathbf{m}.$ Using the masked diffusion inference posterior derived from (3), when $Z_{t(i)} = \mathbf{m}$ we have

$$q(Z_{s(i)} = \mathbf{m} \,|\, Z_{t(i)} = \mathbf{m}, \mathbf{x}) = \frac{1 - \alpha_{s(i)}}{1 - \alpha_{t(i)}}, \quad q(Z_{s(i)} = \mathbf{x} \,|\, Z_{t(i)} = \mathbf{m}, \mathbf{x}) = \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}}.$$

Under the pretrained model parameterization (4), the corresponding terms are

$$q(Z_{s(i)} = \mathbf{m} \,|\, Z_{t(i)} = \mathbf{m}, \mathbf{x}_\theta(Z_{t(i)})) = \frac{1 - \alpha_{s(i)}}{1 - \alpha_{t(i)}},$$

$$q(Z_{s(i)} = \mathbf{x} \,|\, Z_{t(i)} = \mathbf{m}, \mathbf{x}_\theta(Z_{t(i)})) = \frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \langle \mathbf{x}_\theta(Z_{t(i)}), \mathbf{x} \rangle,$$

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and likewise with $\mathbf{x}_\varphi(Z_{t(i)})$ replacing $\mathbf{x}_\theta(Z_{t(i)})$ for our parameterization. Hence,

$$\mathbb{E}_{q(Z_{s(i)}\,|\,Z_{t(i)}=\mathbf{m},\mathbf{x})}\left[\log\frac{q(Z_{s(i)}\,|\,Z_{t(i)}=\mathbf{m},\mathbf{x}_\theta(Z_{t(i)}))}{q(Z_{s(i)}\,|\,Z_{t(i)}=\mathbf{m},\mathbf{x}_\varphi(Z_{t(i)}))}\right]$$

$$=\frac{1-\alpha_{s(i)}}{1-\alpha_{t(i)}}\log\frac{\frac{1-\alpha_{s(i)}}{1-\alpha_{t(i)}}}{\frac{1-\alpha_{s(i)}}{1-\alpha_{t(i)}}}+\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}\log\frac{\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}\langle\mathbf{x}_\theta(Z_{t(i)}),\mathbf{x}\rangle}{\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}\langle\mathbf{x}_\varphi(Z_{t(i)}),\mathbf{x}\rangle}$$

$$=\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}\log\frac{\langle\mathbf{x}_\theta(Z_{t(i)}),\mathbf{x}\rangle}{\langle\mathbf{x}_\varphi(Z_{t(i)}),\mathbf{x}\rangle}.$$

Case II: $Z_{t(i)}\neq\mathbf{m}$. When the token is already revealed, the posterior is deterministic:

$$q(Z_{s(i)}=Z_{t(i)}\,|\,Z_{t(i)}\neq\mathbf{m},\mathbf{x})=1,$$

and this form is identical for the $\mathbf{x}_\theta$- and $\mathbf{x}_\varphi$-parameterized posteriors as well:

$$q(Z_{s(i)}=Z_{t(i)}\,|\,Z_{t(i)}\neq\mathbf{m},\mathbf{x}_\theta(Z_{t(i)}))=1,\quad q(Z_{s(i)}=Z_{t(i)}\,|\,Z_{t(i)}\neq\mathbf{m},\mathbf{x}_\varphi(Z_{t(i)}))=1.$$

Therefore,

$$\mathbb{E}_{q(Z_{s(i)}\,|\,Z_{t(i)}\neq\mathbf{m},\mathbf{x})}\left[\log\frac{q(Z_{s(i)}\,|\,Z_{t(i)},\mathbf{x}_\theta(Z_{t(i)}))}{q(Z_{s(i)}\,|\,Z_{t(i)},\mathbf{x}_\varphi(Z_{t(i)}))}\right]=0$$

Combining both cases,

$$\mathbb{E}_{q(Z_{s(i)}\,|\,Z_{t(i)},\mathbf{x})}\left[\log\frac{q(Z_{s(i)}\,|\,Z_{t(i)},\mathbf{x}_\theta(Z_{t(i)}))}{q(Z_{s(i)}\,|\,Z_{t(i)},\mathbf{x}_\varphi(Z_{t(i)}))}\right]=\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}\log\frac{\langle\mathbf{x}_\theta(Z_{t(i)}),\mathbf{x}\rangle}{\langle\mathbf{x}_\varphi(Z_{t(i)}),\mathbf{x}\rangle}\mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}},$$

which simplifies (14) as follows:

$$\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)},\mathbf{x})}\left[\log\frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{p_\varphi(Z_{s(i)}|Z_{t(i)},\mathbf{y})}\right]$$

$$=\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}\log\frac{\langle\mathbf{x}_\theta(Z_{t(i)}),\mathbf{x}\rangle}{\langle\mathbf{x}_\varphi(Z_{t(i)}),\mathbf{x}\rangle}\mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}-\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)},\mathbf{x})}\left[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)};Z_{s(i)}))\right]$$

$$+\log\mathcal{Z}_\varphi(Z_{t(i)},\mathbf{y}).\qquad(15)$$

Substituting (15) into (13), the conditional likelihood can be bounded as

$$-\log p_\varphi(\mathbf{x}|\mathbf{y})\leq\mathbb{E}_{q(Z_{0:1}|\mathbf{x})}[-\log p_\varphi(\mathbf{x}|Z_0,\mathbf{y})]+\mathbb{E}_{q(Z_1|\mathbf{x})}\left[\log\frac{q(Z_1|\mathbf{x})}{p_\varphi(Z_1|\mathbf{y})}\right]$$

$$+\sum_{i=1}^T\mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}\,\mathrm{CE}(\mathbf{x},\mathbf{x}_\theta(Z_{t(i)}))\,\mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}\right]$$

$$+\sum_{i=1}^T\mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\mathbb{E}_{q(Z_{s(i)}|Z_{t(i)},\mathbf{x})}\left[\log\frac{p_\theta(Z_{s(i)}|Z_{t(i)})}{p_\varphi(Z_{s(i)}|Z_{t(i)},\mathbf{y})}\right]$$

$$=\mathbb{E}_{q(Z_{0:1}|\mathbf{x})}[-\log p_\varphi(\mathbf{x}|Z_0,\mathbf{y})]+\mathbb{E}_{q(Z_1|\mathbf{x})}\left[\log\frac{q(Z_1|\mathbf{x})}{p_\varphi(Z_1|\mathbf{y})}\right]$$

$$+\sum_{i=1}^T\mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}\,\mathrm{CE}(\mathbf{x},\mathbf{x}_\theta(Z_{t(i)}))\,\mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}\right]$$

$$+\sum_{i=1}^T\mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)}-\alpha_{t(i)}}{1-\alpha_{t(i)}}\log\frac{\langle\mathbf{x}_\theta(Z_{t(i)}),\mathbf{x}\rangle}{\langle\mathbf{x}_\varphi(Z_{t(i)}),\mathbf{x}\rangle}\,\mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}\right]$$

$$-\sum_{i=1}^T\mathbb{E}_{q(Z_{t(i)},Z_{s(i)}|\mathbf{x})}\left[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)};Z_{s(i)}))\right]$$

$$+\sum_{i=1}^T\mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\log\mathcal{Z}_\varphi(Z_{t(i)},\mathbf{y})\right].\qquad(16)$$

Treatment of boundary conditions. In masked diffusion the terminal state Z_1 is absorbing (all-mask), hence $q(Z_1|\mathbf{x}) = p_\varphi(Z_1|\mathbf{y})$ and the boundary KL-divergence vanishes. Thus (16) becomes

$$
\begin{aligned}
-\log p_\varphi(\mathbf{x}|\mathbf{y}) \leq\; & \mathbb{E}_{q(Z_0|\mathbf{x})}\big[-\log p_\varphi(\mathbf{x}|Z_0, \mathbf{y})\big] \\
& + \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \operatorname{CE}\big(\mathbf{x}, \mathbf{x}_\theta(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}\right] \\
& + \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \log \frac{\langle\mathbf{x}_\theta(Z_{t(i)}), \mathbf{x}\rangle}{\langle\mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x}\rangle} \mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}\right] \\
& - \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}\Big[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))\Big] \\
& + \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\Big[\log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y})\Big].
\end{aligned}
\tag{17}
$$

Since $\mathbf{x}$ is a deterministic function of Z_0, the reconstruction term in (17) simplifies to:

$$
\begin{aligned}
-\log p_\varphi(\mathbf{x}|\mathbf{y}) \leq\; & \mathbb{E}_{q(Z_0|\mathbf{x})}\big[-\log p_\varphi(\mathbf{x}|Z_0)\big] \\
& + \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \operatorname{CE}\big(\mathbf{x}, \mathbf{x}_\theta(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}\right] \\
& + \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \log \frac{\langle\mathbf{x}_\theta(Z_{t(i)}), \mathbf{x}\rangle}{\langle\mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x}\rangle} \mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}\right] \\
& - \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}\Big[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))\Big] \\
& + \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\Big[\log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y})\Big].
\end{aligned}
\tag{18}
$$

Finally, the decoder is same for unconditional generation and posterior sampling, i.e., $p_\varphi(\mathbf{x}|Z_0) = p_\theta(\mathbf{x}|Z_0)$. Substituting this property in (18) yields

$$
\begin{aligned}
-\log p_\varphi(\mathbf{x}|\mathbf{y}) \leq\; & \mathbb{E}_{q(Z_0|\mathbf{x})}\big[-\log p_\theta(\mathbf{x}|Z_0)\big] \\
& + \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \operatorname{CE}\big(\mathbf{x}, \mathbf{x}_\theta(Z_{t(i)})\big) \mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}\right] \\
& + \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \log \frac{\langle\mathbf{x}_\theta(Z_{t(i)}), \mathbf{x}\rangle}{\langle\mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x}\rangle} \mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}}\right] \\
& - \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}\Big[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))\Big] \\
& + \sum_{i=1}^T \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\Big[\log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y})\Big] := \mathcal{L}_{\mathrm{APS}}(\mathbf{x}, \mathbf{y}; \varphi)
\end{aligned}
\tag{19}
$$

Since $\mathrm{CE}\big(\mathbf{x}, \mathbf{x}_\theta(Z_{t(i)})\big) = -\log\langle \mathbf{x}, \mathbf{x}_\theta(Z_{t(i)})\rangle$, the first two terms in (19) equals the standard NELBO (5) used to train the masked diffusion model. Therefore, we have

$$\mathcal{L}_{\mathrm{APS}}(\mathbf{x}, \mathbf{y}; \varphi) = \mathcal{L}_{\mathrm{NELBO}}(\mathbf{x}; \theta) + \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\frac{\alpha_{s(i)} - \alpha_{t(i)}}{1 - \alpha_{t(i)}} \log \frac{\langle \mathbf{x}_\theta(Z_{t(i)}), \mathbf{x}\rangle}{\langle \mathbf{x}_\varphi(Z_{t(i)}), \mathbf{x}\rangle} \mathbf{1}_{\{Z_{t(i)}=\mathbf{m}\}} \right]$$

$$- \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}, Z_{s(i)}|\mathbf{x})}\left[\log q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)})) \right]$$

$$+ \sum_{i=1}^{T} \mathbb{E}_{q(Z_{t(i)}|\mathbf{x})}\left[\log \mathcal{Z}_\varphi(Z_{t(i)}, \mathbf{y}) \right],$$

which upon regrouping completes the statement of the theorem. $\qquad\square$

Implications. **Theorem A.2** establishes a principled upper bound on the negative log-posterior likelihood when posterior sampling is performed without additional training.

- *Reuse of pretrained objective.* The bound $\mathcal{L}_{\mathrm{APS}}(\mathbf{x}, \mathbf{y}; \varphi)$ is expressed in terms of the standard masked diffusion $\mathcal{L}_{\mathrm{NELBO}}(\mathbf{x}; \theta)$, meaning that posterior sampling can be performed using pretrained masked diffusion models.

- *Adaptation gap.* The log-ratio correction term quantifies the mismatch between the pretrained transitions $\mathbf{x}_\theta$ and the proposed posterior transitions $\mathbf{x}_\varphi$, effective only at masked positions. This isolates the additional cost of posterior sampling.

- *Measurement consistency.* The final summation enforces alignment with the measurement likelihood $q(\mathbf{y}|\mathbf{x}_\varphi(Z_{t(i)}; Z_{s(i)}))$, ensuring that the sampler accounts for observations at each diffusion step.

- *Boundary conditions.* As in the training bound, the absorbing mask state renders the boundary KL-divergence constant (often zero), so the effective objective simplifies to the pretrained NELBO plus adaptation and measurement terms.

- *Test-time posterior sampling.* Together, the decomposition clarifies how posterior sampling can be performed without retraining: start from the pretrained NELBO and add corrections for model adaptation at masked positions while incorporating measurements via tilting.

B ADDITIONAL EXPERIMENTS

This section provides supplementary details and evaluations of our APS method. We first describe implementation details for both inverse problems and stylization (§B.1), followed by algorithmic analysis including pseudocode and hyperparameter studies (§B.2). We then examine the impact of design choices in our ablations (§B.3), evaluate using the same prior (§B.4), and discuss computational complexity (§B.5). Next, we outline the compared baselines (§B.7) and summarize benchmarks and metrics (§B.8). Finally, we present additional results (§B.9), limitations (§B.10), and conclude with reproducibility (§B.11) and ethics (§B.12).

B.1 IMPLEMENTATION DETAILS

B.1.1 INVERSE PROBLEMS

For inverse problems, we implement our test-time optimization using two main configurations corresponding to the APS (512×512) and APS-L (1024×1024) results reported in §4. The full reverse sampling process is discretized into 15 time steps following a cosine mask schedule, using a classifier-free guidance scale of 3.5.

At each of the 15 reverse steps, we perform an inner optimization loop to ensure measurement consistency. This loop consists of 100 optimization steps using the Adam optimizer with a learning rate of 1.0. The total loss function is a weighted sum of two components: (1) a reconstruction loss, which could be L1 or L2 norm, but we choose L1 norm: $\|\mathbf{y} - \mathcal{A}(\hat{\mathbf{x}})\|_1$ as it is known to generate sharper quality), and (2) a VGG perceptual loss with a coefficient of 10^{-3}.

The specific parameters for each degradation operator $\mathcal{A}(\cdot)$ vary by task, with all measurements simulated by adding Gaussian noise of $\sigma = 0.05$. For super resolution, we use a $4\times$ downsampling factor. For Gaussian Deblurring, the operator is a Gaussian kernel of size 61×61 with a standard deviation of 3.0. Motion Deblurring uses a kernel of the same size with an intensity parameter of 0.5 (on a scale of 0 for linear to 1 for highly nonlinear), corresponding to a moderately nonlinear motion path. For Inpainting, we randomly remove 70% of pixels. Nonlinear Deblurring kernels are generated using the `KernelWizard`[3] model from the `bkse`[4] library. Finally, High Dynamic Range (HDR) reconstruction is modeled by scaling the image data and clipping the result, following the operation $\text{clip}(\text{data} \times 2, -1, 1)$.

B.1.2 Reference-based Stylization

Our approach to training-free, reference-based stylization leverages the core APS framework by framing the task as a *highly nonlinear* inverse problem. Let $\hat{x}_{\text{ref}}$ denote the conditional reference image providing the style. The target measurement, a style vector y, is obtained by applying a pretrained Contrastive Style Descriptor (CSD) (Somepalli et al., 2024) model as our measurement operator $\mathcal{A}(\cdot)$ to this reference image, i.e., $y = \mathcal{A}(\hat{x}_{\text{ref}})$. At each reverse diffusion step t, APS aims to generate a sample whose style matches this target.

As discussed in §3.2, there are two main stages of APS. In the first stage (§3.2.1), we perform a differentiable forward pass by computing the expected codebook embedding $\bar{x}^l = \sum_{k=1}^{K} c_k x_\varphi^l(z_{t(i)})$ for $l = 1, \ldots, L$ using the model's output probabilities $x_\theta(z_t)^l$ as initial condition for $\varphi_{t(i)}^l$. The straight-through estimator is then used to obtain a differentiable image representation $\hat{x} = \mathcal{D}(\tilde{x})$. To guide the optimization, the measurement consistency loss is calculated as the cosine distance between the style vector of the generated image and the target style vector y as follows:

$$\mathcal{L}_{\text{style}}(\varphi_{t(i)}) = 1 - \frac{\langle \mathcal{A}(\hat{x}), y \rangle}{\|\mathcal{A}(\hat{x})\| \|y\|}.$$

For each reverse step, we perform 100 optimization steps using an Adam optimizer with a learning rate of 0.1. Gradients from this style loss are backpropagated through the frozen VQ-VAE decoder $\mathcal{D}(\cdot)$ and the straight-through estimator $\tilde{x}$ to update all the entries of the conditional probability table $\varphi_{t(i)} = \{\varphi_{t(i)}^l\}_{l=1}^{L}$.

In the second stage (§3.2.2), the posterior estimate $x_\varphi(z_{t(i)})$ obtained from the first stage is used adaptively unmask anchor tokens in the sequence. Both the processes continue over 15 total steps following a cosine mask schedule. The fully unmasked sequence satisfies the stylistic constraints without requiring any task-specific retraining of the foundation model.

B.1.3 High-Resolution Inference

To demonstrate the scalability of our method, we experiment with a higher-resolution setting. For a fair comparison on our 256×256 benchmark, we follow the upsampling protocol described in G2D2 (Murata et al., 2024). Specifically, for both our standard (512×512) and large-scale (APS-L, 1024×1024) models, we first upsample the benchmark images to the model's native resolution before applying the forward operator. For the APS-L configuration, this increases the number of visual tokens by a factor of four (from 1024 to 4096)[5]. The Transformer-based MMaDA (Yang et al., 2025) model accommodates this by processing a longer sequence without architectural changes. After the high-resolution reconstruction is complete, we downsample the output back to the benchmark's native 256×256 resolution for evaluation. As demonstrated in our experiments (§4.1, §4.2), this approach further improves performance, achieving substantial gains in both PSNR and LPIPS.

B.2 Algorithm Details

Algorithm 1 details our APS procedure. The process begins with a fully masked latent space, z_1, and iteratively refines the image over T reverse diffusion steps. Each step features an inner opti-

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[3] https://github.com/LeviBorodenko/motionblur
[4] https://github.com/VinAIResearch/blur-kernel-space-exploring
[5] The visual tokenizer (Yu et al., 2024) used in MMaDA (Yang et al., 2025) uses $16\times$ downscaling, generating $1024 = 32 \times 32$ tokens for an image of size 512×512.

Algorithm 1: Test-Time Anchored Posterior Sampling (APS)

1: **Input:** Diffusion steps T, measurement $\mathbf{y}$, denoiser $\mathbf{x}_\theta^{\text{logits}}(\cdot)$, operator $\mathcal{A}(\cdot)$, decoder $\mathcal{D}(\cdot)$.
2: **Tunable parameters:** Optimization steps M, learning rate η, loss coefficients $\lambda_{\text{p}}, \lambda_e$.
3: **Output:** Reconstructed image $\hat{\mathbf{x}}_{\text{final}}$.
4: Initialize latent state $\mathbf{z}_1 \leftarrow \{\mathbf{m}\}^L$ (all masked)
5: **for** $i = 1$ **to** T **do**
6:    $\varphi \leftarrow \mathbf{x}_\theta^{\text{logits}}(\mathbf{z}_{t(i)})$ ▷ Quantized Expectation §3.2.1
7:    **for** $m = 1$ **to** M **do**
8:       $\mathbf{x}_\varphi(\mathbf{z}_{t(i)}) = \text{Softmax}(\varphi)$
9:       $\bar{\mathbf{x}} \leftarrow \sum_{k=1}^K \mathbf{c}_k \mathbf{x}_\varphi(\mathbf{z}_{t(i)})$
10:      $\mathbf{x} = \mathcal{Q}_{\text{lfq}}(\bar{\mathbf{x}})$
11:      $\tilde{\mathbf{x}} \leftarrow \bar{\mathbf{x}} + \left[\mathbf{x} - \bar{\mathbf{x}}\right]_{\text{sg}}$
12:      $\hat{\mathbf{x}} \leftarrow \mathcal{D}(\tilde{\mathbf{x}})$
13:      $\mathcal{L} \leftarrow \mathcal{L}_{\text{reconstruction}}(\mathcal{A}(\hat{\mathbf{x}}), \mathbf{y}) + \lambda_{\text{p}}\mathcal{L}_{\text{perceptual}}(\mathcal{A}(\hat{\mathbf{x}}), \mathbf{y})$
14:      $\varphi \leftarrow \text{Adam}(\varphi, \nabla_\varphi \mathcal{L}, \eta)$
15:    **end for**
16:    $\mathbf{x}_\varphi^*(\mathbf{z}_{t(i)}) = \text{Softmax}(\varphi)$ ▷ Anchored Remasking §3.2.2
17:    $\bar{\mathbf{x}}^* \leftarrow \sum_{k=1}^K \mathbf{c}_k \mathbf{x}_\varphi^*(\mathbf{z}_{t(i)})$
18:    $\mathbf{x} = \mathcal{Q}_{\text{lfq}}(\bar{\mathbf{x}}^*)$
19:    $\kappa^l \leftarrow \langle (\mathbf{x}_\varphi^*(\mathbf{z}_{t(i)})^l, \mathbf{x}^l \rangle$ for $l = 1, \ldots, L$
20:    $\mathcal{P}_{t(i)} \leftarrow \{l : \kappa^l \geq \tau_{t(i)}\}$
21:    $\mathbf{z}_{s(i)} \leftarrow \text{UpdateState}(\mathbf{z}_{t(i)}, \mathbf{x}, \mathcal{P}_{t(i)})$
22: **end for**
23: $\hat{\mathbf{x}}_{\text{final}} \leftarrow \mathcal{D}(\mathbf{z}_0)$
24: **return** $\hat{\mathbf{x}}_{\text{final}}$

---

mization phase designed to align the model's predictions with the given measurement $\mathbf{y}$. To enable gradient-based optimization through the discrete quantization step which assigns each dimension of an embedding to its nearest value in $\{-1, 1\}$ we employ the straight-through estimator (STE) via the stop-gradient operator, $\text{sg}(\cdot)$.

This optimization is guided by a composite loss function. For reconstruction, we primarily use the Mean Absolute Error (MAE, or L1 loss), which we find produces perceptually superior results compared to the Mean Squared Error (MSE). While MSE corresponds to maximizing the Gaussian log-likelihood, MAE is typically more robust. This is supplemented with a VGG perceptual loss to enforce similarity in the feature domain, further improving visual quality. Finally, a small entropy term is included to regularize the posterior distribution. Once the optimization at a given step is complete, the Anchored Remasking strategy uses the model's confidence defined as the probability assigned to the chosen token for each position to selectively unmask tokens for the next iteration.

To determine the optimal weight for the perceptual loss, we conducted an ablation study on its coefficient, λ_{p}. As shown in Table 4, a value of 10^{-3} provides the optimal trade-off between reconstruction fidelity (PSNR) and perceptual quality (LPIPS) across our benchmarks.

Effect of perceptual loss Coefficient. Table 4 reports the effect of varying the perceptual loss coefficient on ImageNet and FFHQ. Small weights (10^{-5}, 10^{-4}) only marginally improve perceptual quality over the baseline, while larger weights (10^{-2}, 10^{-1}) overly emphasize perceptual similarity at the cost of distortion and structure. A coefficient of 10^{-3} provides the best balance: on ImageNet, it reduces LPIPS from 0.416 to 0.334 while maintaining PSNR 23.61 and SSIM 0.639, and on FFHQ, it achieves the lowest LPIPS (0.247) with a slight drop in PSNR (26.61) and SSIM (0.781). We therefore adopt 10^{-3} as the default across datasets. Notably, our approach introduces only this single hyperparameter, whereas G2D2 relies on multiple carefully tuned schedules (e.g., four different coefficients) that must be re-optimized for each task. In contrast, we use the same setting for all inverse problems, underscoring the robustness and simplicity of our posterior sampler relative to prior discrete diffusion methods.

Table 4: **Hyperparameter sweeps on perceptual loss coefficient (λ_p) on ImageNet and FFHQ.** Highlighted rows denote the chosen setting (1e-3), which offers a trade-off among perceptual quality (LPIPS), distortion (PSNR), and structure (SSIM).

(a) ImageNet					(b) FFHQ			
λ_p	LPIPS $\downarrow$	PSNR $\uparrow$	SSIM $\uparrow$		λ_p	LPIPS $\downarrow$	PSNR $\uparrow$	SSIM $\uparrow$
0.0	0.416	23.98	0.652		0.0	0.311	27.32	0.801
1e-5	0.402	24.05	**0.658**		1e-5	0.301	27.40	0.803
1e-4	0.380	**24.06**	0.655		1e-4	0.268	**27.67**	**0.812**
1e-3	0.334	23.61	0.639		1e-3	**0.247**	26.61	0.781
1e-2	**0.325**	22.06	0.566		5e-3	0.252	24.90	0.729
5e-2	0.328	21.44	0.543		1e-2	0.256	24.49	0.715
1e-1	0.327	21.29	0.539		1e-1	0.260	23.44	0.692

B.3 ABLATION STUDY

To better understand the impact of our algorithmic innovations, we conduct a systematic ablation study on ImageNet SR ($4\times$), as shown in Figure 5 (top row). Our analysis focuses on three core design choices: (i) *standard remasking*, which highlights the limitations of confidence-based token selection under the prior; (ii) *quantized expectation*, which addresses sampling bias and improves measurement consistency; and (iii) *anchored remasking*, which preserves informative tokens identified by optimization while suppressing spurious high-confidence background tokens. Together, these ablations disentangle the contributions of each component, providing both theoretical insight and qualitative evidence (Figure 5) into how APS achieves stable and measurement-consistent posterior sampling.

B.3.1 EFFECT OF STANDARD REMASKING

Standard remasking in MaskGIT (Chang et al., 2022) and MMaDA (Yang et al., 2025) proceeds as follows. At each iteration, the base denoiser produces a distribution over all tokens $\mathbf{x}_\theta(\mathbf{z}_{t(i)}) = \{\mathbf{x}_\theta^l(\mathbf{z}_{t(i)})\}_{l=1}^L$ given the current partially unmasked sequence $\mathbf{z}_t$. Sampled tokens $\mathbf{x} = \{\mathbf{x}^l\}_{l=1}^L$ are drawn independently at each position, and previously unmasked tokens are carried over to the next state $\mathbf{z}_s$. For masked positions, the confidence of each sampled token $\mathbf{x}^l$ is defined as

$$\kappa_t^l = \langle \mathbf{x}_\theta^l(\mathbf{z}_{t(i)}), \mathbf{x}^l \rangle.$$

Tokens to unmask are then selected based on an adaptive threshold schedule (e.g., cosine),

$$\mathcal{P}_t = \{l : \kappa_t^l \geq \tau_t\},$$

which favors unmasking the most confident tokens under the prior distribution $\mathbf{x}_\theta(\mathbf{z}_t)$.

In language generation, this corresponds to unmasking frequent, low-information tokens (e.g., articles or conjunctions) (Rout et al., 2025a). Analogously in images, the model tends to unmask background regions first, since they dominate training statistics and are easier to predict. As a result, informative foreground tokens remain masked until late in the process, limiting semantic guidance and increasing conditional entropy.

Qualitatively, Figure 5 (second row) illustrates this phenomenon. While the model confidently unmasks background tokens early, the salient object is revealed only much later, highlighting the limitation of standard remasking for posterior sampling.

B.3.2 EFFECT OF QUANTIZED EXPECTATION

Standard remasking suffers from two key issues: (i) sampling bias, and (ii) independent token-wise confidence. When sampling tokens directly from the unconditional distribution $\mathbf{x}_\theta(\mathbf{z}_t)$, the model may pick unrelated or spurious tokens, which—once unmasked—remain fixed in all future steps. This introduces inconsistency and often locks the model into poor generations.

To address this, we propose *quantized expectation* (§3.2.1). Instead of sampling, we tilt the unconditional distribution $\mathbf{x}_\theta(\mathbf{z}_t)$ towards the measurement likelihood, obtaining an approximate posterior

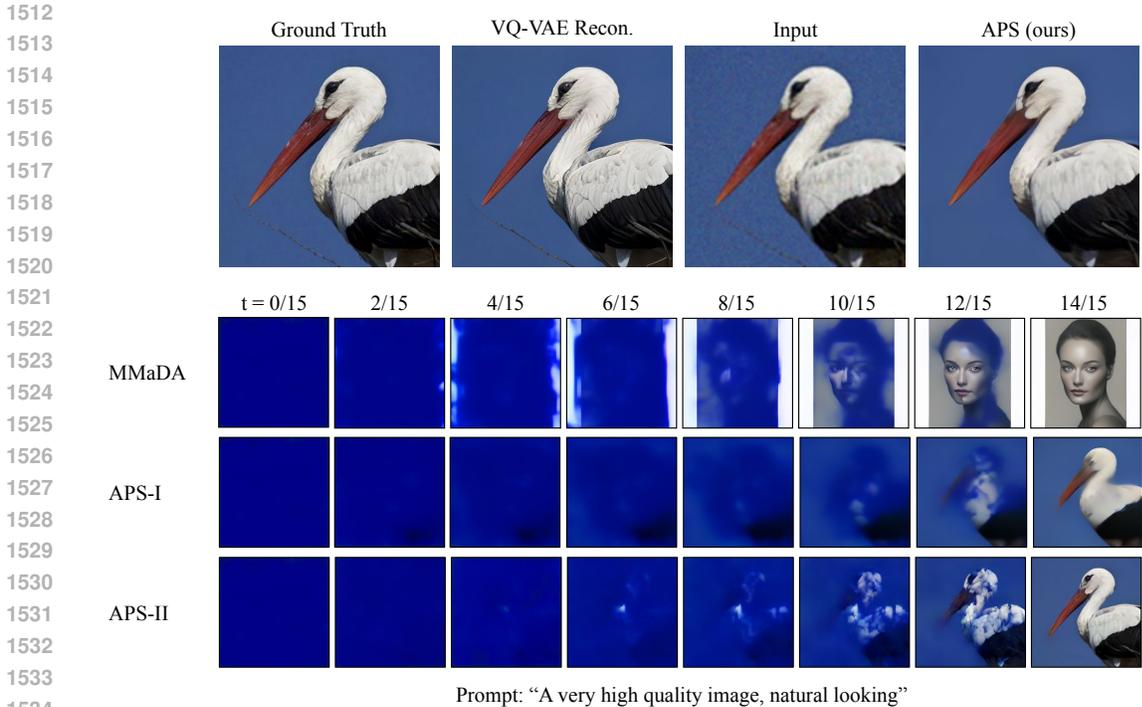Prompt: "A very high quality image, natural looking"

Figure 5: **Ablation study of design choices in APS.** We study the posterior sampling problem with the input of the noisy image of a bird (top row, third image from left). In the base case of using MMada (second row), since it does not take an image as an input (but only text), the final image is inconsistent with the noisy input image (image of person instead of bird). Thus, this shows the effect of *standard remasking*, where high-confidence background tokens are unmasked early. The third row (APS-I) adds *quantized expectation*, which mitigates sampling bias and improves consistency with the measurement, but still relies on prior-based confidence remasking. This results in blurry reconstructions. Finally, the fourth row (APS-II) combines *quantized expectation* with *anchored remasking*, preserving optimized anchor tokens while remasking uninformative background tokens. This combination yields the most stable and measurement-consistent generations, as discussed in §B.3. Indeed, we note that best case one could hope for is the direct VQ-VAE reconstruction (top row, second from left) of the Ground Truth image (top row, leftmost). We observe that APS (top row, rightmost) is comparable in quality to the VQ-VAE reconstruction.

$\mathbf{x}_\varphi(\mathbf{z}_t)$. We then optimize in the span of codebook embeddings by treating the tilted probabilities as coefficients in a linear combination and passing their expectation through the decoder using a straight-through estimator. The resulting embedding is then quantized back to the nearest valid token. This procedure implicitly maximizes the measurement likelihood, avoids sampling noise, and enables the discovery of tokens with zero prior probability mass but strong measurement consistency–leading to a better posterior sample.

We treat such tokens as "anchor tokens," since they minimize reconstruction error and provide critical guidance under the measurement operator. As shown in Figure 5 (third row, APS-I), quantized expectation corrects sampling bias and yields reconstructions that remain consistent with observations throughout the denoising trajectory.

B.3.3 EFFECT OF ANCHORED REMASKING

A limitation of confidence-based remasking under the prior is that anchor tokens, obtained through our optimization procedure, may receive near-zero probability mass under $\mathbf{x}_\theta(\mathbf{z}_t)$. As a result, the model would discard these informative tokens in favor of background tokens, which the prior predicts with high confidence. This reintroduces the very bias we aim to avoid.

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Table 5: **Effect of Optimization Steps (Super-Resolution (4×) on FFHQ).** Performance saturates beyond 100 optimization steps, confirming robustness and efficiency.

# Optimization Steps (M)	PSNR ↑	SSIM ↑	LPIPS ↓
0	9.49	0.1855	0.6747
10	21.47	0.5541	0.4298
50	24.80	0.7130	0.2730
100	25.71	0.7422	0.2468
200	25.72	0.7389	0.2474

Table 6: **Quantitative results for super resolution (4×) on FFHQ.** APS consistently outperforms G2D2 across standard evaluation metrics: PSNR, SSIM and LPIPS. All the methods have a reconstruction loss by default. Both G2D2 and APS use the same base discrete diffusion model VQ-Diffusion (Gu et al., 2022), which has a VQ-VAE (Van Den Oord et al., 2017) tokenizer as opposed to the LFQ (Yu et al., 2024) tokenizer used in MMaDA (Yang et al., 2025). The runtime of G2D2 and APS is nearly identical (see Table 7).

Method	PSNR ↑	SSIM ↑	LPIPS ↓
G2D2 (original)	25.33	0.706	0.353
G2D2 (w/ perceptual 1e-3)	25.46	0.720	0.349
G2D2 (w/ perceptual 1e-2)	25.34	0.707	0.357
APS (Ours)	**26.80**	**0.759**	**0.310**

To address this, we compute token confidence using the posterior estimate $\mathbf{x}_\varphi(\mathbf{z}_t)$ rather than the unconditional prior. In this way, anchor tokens identified via quantized expectation are preserved, while low-likelihood background tokens are remasked. Qualitatively, this effect is evident at $t = 6/15$ in Figure 5 (fourth row, APS-II): unlike the standard prior-based strategy (second row), which prematurely unmasks background pixels, our approach commits to anchor tokens aligned with the bird's body. This ensures that the background (blue sky in the measurement) is correctly down-weighted, as it is inconsistent with the prior white background generated by $\mathbf{x}_\theta$. Subsequent steps therefore refine the image conditioned on these anchor tokens, reducing conditional entropy and producing reconstructions that remain faithful to the measurements.

B.3.4 EFFECT OF OPTIMIZATION STEPS.

Table 5 presents an ablation study on the number of inner optimization steps used during test-time anchoring in **Algorithm 1**. We observe that performance improves rapidly up to 50 steps and saturates beyond 100 steps, confirming that our method converges efficiently without requiring additional iterations. Importantly, all optimization steps backpropagate only through the lightweight VQ-VAE decoder rather than the large 8B-parameter MMaDA backbone, keeping computational overhead minimal.

B.4 EVALUATION UNDER IDENTICAL DISCRETE DIFFUSION PRIOR

We compare APS to G2D2 using the *official* G2D2 codebase (§B.7) on 100 images from the FFHQ validation set. Both methods use an identical compute budget: 100 reverse diffusion steps, each with 30 inner optimization steps. As shown in Table 6, APS consistently improves over G2D2 across all metrics: PSNR improves from 25.46 to 26.80, SSIM from 0.717 to 0.759, and LPIPS decreases from 0.350 to 0.310. Since the generative prior and compute budget are identical, these performance gains arise purely from our algorithmic innovations: quantized expectation (§3.2.1), anchored remasking (§3.2.2), and the use of a perceptual loss. This experiment demonstrates that APS is not only effective but also compatible with both *mask-based* and *uniform* discrete diffusion frameworks.

Table 7: **Quantitative results on sampling efficiency of continuous and discrete samplers.** PDM/LDM denote pixel-/latent-space continuous diffusion models, respectively; VQ-Diffusion and MMaDA are discrete diffusion models. Rows shaded gray report runtimes on a single NVIDIA A6000 GPU copied from G2D2 (Murata et al., 2024); rows shaded orange are measured by us on a single NVIDIA A100 GPU.

Method	Model	Resolution	GPU (GiB)	Time (s)	#Steps
DPS (Chung et al., 2023)	PDM	256	10.7	277	1000
DDRM (Kawar et al., 2022)	PDM	256	5.8	4	20
PSLD (Rout et al., 2023)	LDM	512	20.9	738	1000
ReSample (Song et al., 2024)	LDM	256	7.1	555	500
G2D2 (Murata et al., 2024)	VQ-Diffusion	256	4.7	194	100
DPS (Chung et al., 2023)	PDM	256	10.7	180	1000
LDPS (Rout et al., 2023)	LDM	256	15.4	190	1000
PSLD (Rout et al., 2023)	LDM	256	15.5	194	1000
G2D2 (Murata et al., 2024)	VQ-Diffusion	256	4.7	107	100
APS (ours)	VQ-Diffusion	256	4.6	**106**	100
APS (ours)	MMaDA	256	19.2	**55**	15
PSLD (Rout et al., 2023)	LDM	512	20.9	720	1000
APS (ours)	MMaDA	512	26.2	**121**	15
APS-L (ours)	MMaDA	1024	51.8	**484**	15

B.5 COMPUTATIONAL COMPLEXITY

Analysis. Table 7 highlights the *sampling efficiency* of our discrete posterior sampler (APS) relative to both continuous-diffusion baselines and the prior discrete sampler G2D2.

Against continuous (pixel/latent) diffusion. Pixel-space samplers (DPS, DDRM) either require very long Markov chains (e.g., DPS: 1000 steps, 277 s) or sacrifice quality when shortened; latent-space samplers (PSLD, ReSample) still need 500–1000 steps and hundreds of seconds per image at 256×256–512×512 resolutions (e.g., PSLD: 738 s at 512×512). In contrast, APS runs with only 15 reverse steps on the MMaDA backbone: 55 s at 256×256 and 121 s at 512×512 on a single A100—$66\times$ *shorter chain* for comparable or better perceptual quality. Crucially, at 512×512 APS matches the quality of PSLD as shown in Table 1 while being $\sim 6\times$ faster (PSLD: $\sim$720–740 s vs. APS: 121 s). When scaling the sequence length to $L{=}4096$ tokens (corresponding resolution 1024×1024), APS-L completes in 484 s and becomes significantly more accurate: ImageNet Gaussian deblur improves by 22.7% LPIPS and 9.6% PSNR over PSLD—with $\sim 1.5\times$ less time.

Against prior discrete diffusion (G2D2). Under the same budget, APS matches or improves G2D2's runtime while yielding better reconstructions: at 256×256 with VQ-Diffusion both methods use 100 steps, but APS yields higher quality as given in Table 6. Moreover, our 1024×1024 configuration (APS-L) demonstrates better test-time scaling behavior compared continuous diffusion: we keep 15 steps and still obtain substantial quality gains at reasonable cost (484 s).

Importantly, continuous methods struggle to match this performance without prohibitive runtimes. PSLD (Rout et al., 2023) already takes nearly 12 minutes to process a single 512×512 image and more complex methods such as P2L (Chung et al., 2024) take around 30 minutes for the same resolution. Therefore, training-free posterior sampling using continuous diffusion at very high resolutions such as 1024×1024 becomes computationally prohibitive.

B.6 ANALYSIS OF ROBUSTNESS TO MEASUREMENT NOISE

Table 8 evaluates the noise robustness of APS against G2D2 under Gaussian noise with varying standard deviations. APS consistently improves PSNR and SSIM while reducing LPIPS across all noise levels, indicating superior visual fidelity and structural consistency. The performance gap re-

Table 8: **Noise robustness analysis for 256×256 Super-Resolution (4x) on FFHQ.** APS consistently outperforms G2D2 across varying noise levels ($\sigma \in \{0.01, 0.02, 0.03, 0.04, 0.05\}$) in PSNR, SSIM, and LPIPS. The performance gain of APS remains consistent across different noise levels.

Measurement Noise (σ)	Method	PSNR ↑	SSIM ↑	LPIPS ↓
0.01	G2D2	25.98	0.739	0.328
	APS	**27.52**	**0.784**	**0.268**
0.02	G2D2	25.90	0.738	0.333
	APS	**27.42**	**0.781**	**0.273**
0.03	G2D2	25.88	0.733	0.338
	APS	**27.31**	**0.774**	**0.281**
0.04	G2D2	25.64	0.726	0.341
	APS	**27.04**	**0.764**	**0.290**
0.05	G2D2	25.46	0.717	0.350
	APS	**26.80**	**0.759**	**0.310**

mains stable as σ increases, showing that APS maintains high reconstruction quality even under strong measurement corruption. These results, along with experiments spanning different tokenizers (LFQ, VQ-VAE), datasets (FFHQ, ImageNet, StyleAligned), and tasks (inpainting, deblurring, super-resolution), demonstrate the robustness and versatility of our approach.

B.7 COMPARED BASELINES

We compare our method against state-of-the-art posterior samplers using pixel-/latent-space continuous and discrete diffusion models. Each baseline is evaluated under the same data as ours. We follow the experimental setup from G2D2 and reuse the baseline implementations to ensure a fair comparison. To address the resolution mismatch between our model and the benchmark datasets, we adopt the protocol from G2D2 (Murata et al., 2024). The benchmark images are first upsampled to match the input resolution of our base model MMaDA (Yang et al., 2025). The forward corruption operator and our posterior sampling method are then applied in this high-resolution space. Finally, the resulting output is downsampled to the original 256×256 resolution for a fair evaluation. A brief description of each baseline and links to available source code are provided below:

- **DPS** (Chung et al., 2023): A *continuous* diffusion-based method operating in pixel space that solves noisy inverse problems by employing a one-step gradient update in the pixel domain. Source: https://github.com/DPS2022/diffusion-posterior-sampling

- **DDRM** (Kawar et al., 2022): A *continuous* diffusion-based method in pixel space, evaluated using the same base models as DPS. Source: https://github.com/bahjat-kawar/ddrm

- **DiffPIR** (Zhu et al., 2023): A pixel-space *continuous* diffusion-based method for plug-and-play image restoration. Source: https://github.com/yuanzhi-zhu/DiffPIR

- **DAPS** (Zhang et al., 2025): A *continuous* diffusion-based method that employs a decoupled noise annealing strategy to solve inverse problems. Source: https://github.com/zhangbingliang2019/DAPS

- **PSLD** (Rout et al., 2023): A latent-space *continuous* diffusion method that solves inverse problems by performing a one-step gradient update in the latent space and optimizing towards the fixed point of a VAE. Source: https://github.com/LituRout/PSLD

- **ReSample** (Song et al., 2024): A *continuous* latent-space diffusion method that enforces a hard data-consistency constraint during sampling. Source: https://github.com/soominkwon/resample

- **G2D2** (Murata et al., 2024): A *discrete* diffusion posterior sampler that uses a star-shaped noising process and Gumbel-Softmax continuous relaxation to enable gradient guidance

in discrete space. We follow the exact implementation and hyperparameters provided in the original paper. Specifically, we use 100 reverse diffusion steps and 30 optimization steps per reverse step. The learning rate and the coefficient for the KL-divergence loss are scheduled logarithmically, as proposed. Source: https://github.com/sony/g2d2

- **SGDD** (Chu et al., 2025): A *discrete* diffusion posterior sampler that uses a split Gibbs sampler, reweights probabilities by Hamming distance, and employs rejection sampling via Metropolis–Hastings. Source: https://github.com/chuwd19/Split-Gibbs-Discrete-Diffusion-Posterior-Sampling

B.8 BENCHMARKS & METRICS

The APS method is evaluated on standard inverse problem benchmarks and is also shown to generalize to more complex tasks, including non-linear inverse problems and training-free stylization. The evaluation uses two main datasets to cover diverse image types and resolutions, with performance measured using Learned Perceptual Image Patch Similarity (LPIPS) ($\downarrow$: lower the better), Peak Signal-to-Noise Ratio (PSNR) ($\uparrow$: higher the better), and Structural Similarity Index (SSIM) ($\uparrow$: higher the better).

FFHQ (Flickr-Faces-HQ) (Karras et al., 2019):

- **Dataset Focus**: High-resolution face images.
- **Evaluation Set**: To maintain a fair comparison with prior work, specifically SGDD (Chu et al., 2025), G2D2 (Murata et al., 2024) and DAPS (Zhang et al., 2025), our APS algorithm is evaluated on **100 images** (indices $0, 1, \ldots, 99$) from the FFHQ validation set.
- **Tasks**: The evaluation includes (1) linear inverse problems: SR ($4\times$), Gaussian Deblurring, random inpainting, and motion deblur and (2) nonlinear inverse problems: high dynamic range (HDR) and nonlinear blur.

ImageNet (Deng et al., 2009):

- **Dataset Focus**: Diverse natural images.
- **Evaluation Set**: Following the experimental setup by G2D2, a subset of **100 images** is selected from the validation set, ensuring diverse class representation by sampling from classes with indices $0, 10, \ldots, 990$. The specific image list is publicly available in the following text file: imagenet_val_1k.txt $\rightarrow$ https://github.com/XingangPan/deep-generative-prior/.
- **Tasks**: The evaluation includes the same linear and nonlinear inverse problems as in FFHQ.

Licenses and Usage. Both FFHQ and ImageNet datasets used in this work are publicly available and licensed for research use. FFHQ dataset, including its documentation and metadata, is distributed by NVIDIA Corporation under the Creative Commons Attribution-NonCommercial-ShareAlike 4.0 International (CC BY-NC-SA 4.0) license. ImageNet data is provided free of charge to researchers, for non-commercial research and educational purposes.

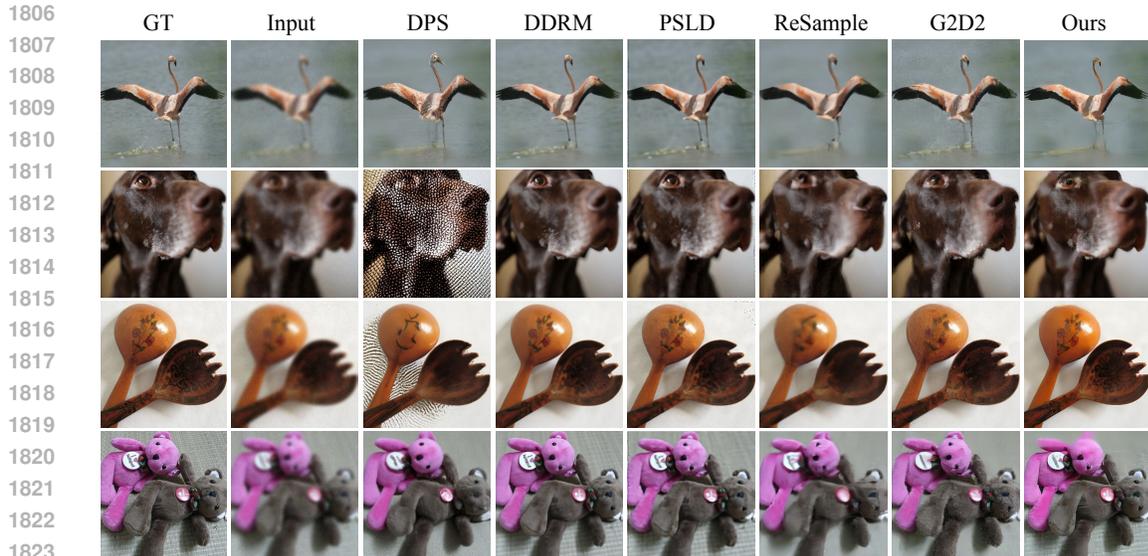## B.9 ADDITIONAL RESULTS

B.9.1 GENERAL INVERSE PROBLEMS

Figure 6 illustrates super resolution ($4\times$) results on ImageNet (Deng et al., 2009). Competing methods—DPS (Chung et al., 2023), DDRM (Kawar et al., 2022), PSLD (Rout et al., 2023), and Re-Sample (Song et al., 2024)—recover coarse structures but often yield blurry textures or color shifts, while G2D2 (Murata et al., 2024) sharpens details at the cost of noticeable artifacts. In contrast, APS produces sharper and more natural reconstructions across both object and animal categories, closely adhering to the ground truth.

Similarly, Figure 7 shows results for Gaussian deblurring. Continuous methods again capture overall structure but leave residual blur or noise, and G2D2 (Murata et al., 2024) partially enhances details yet struggles with fine textures. APS delivers cleaner and more faithful reconstructions, effectively balancing sharpness and natural appearance across diverse scenes.
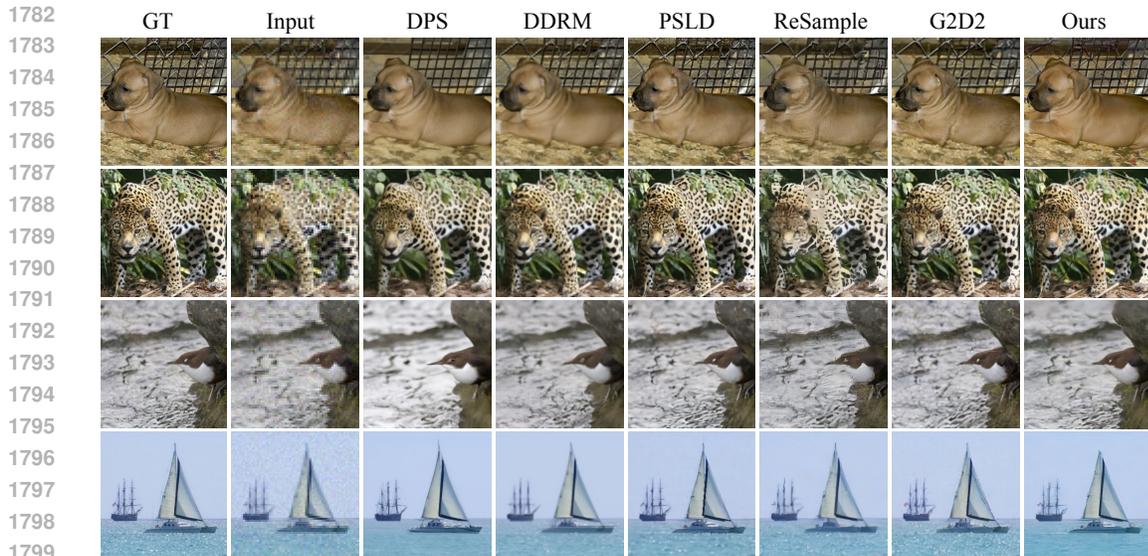
Figure 6: **Additional qualitative results for super resolution** ($4\times$) **on ImageNet.** Compared to continuous baselines (DPS, DDRM, PSLD, ReSample) and the discrete baseline G2D2, APS produces sharper details and more faithful reconstructions across diverse examples. For instance, in the second row, the leopard's eyes are reconstructed with fine detail, noticeably better than the baselines.

Figure 7: **Additional qualitative results for Gaussian deblurring on ImageNet.** Compared to continuous baselines (DPS, DDRM, PSLD, ReSample) and the discrete baseline G2D2, our APS sampler achieves sharper textures, less artifacts, and more faithful reconstructions across diverse examples. For instance, in the third row, both the floral pattern on the outside of the wooden spoon and the artistic pattern inside it are accurately preserved by our method, whereas most baselines either miss or misrepresent these details.

Figure 8 compares APS against continuous (DPS (Chung et al., 2023), DDRM (Kawar et al., 2022), PSLD (Rout et al., 2023), ReSample (Song et al., 2024)) and discrete (G2D2 (Murata et al., 2024)) approaches on FFHQ super resolution. Continuous methods capture overall facial structure but tend to oversmooth, leaving blurred or distorted skin textures, while ReSample introduces strong artifacts. G2D2 sharpens details but produces unnatural appearances. APS, by contrast, reconstructs sharper
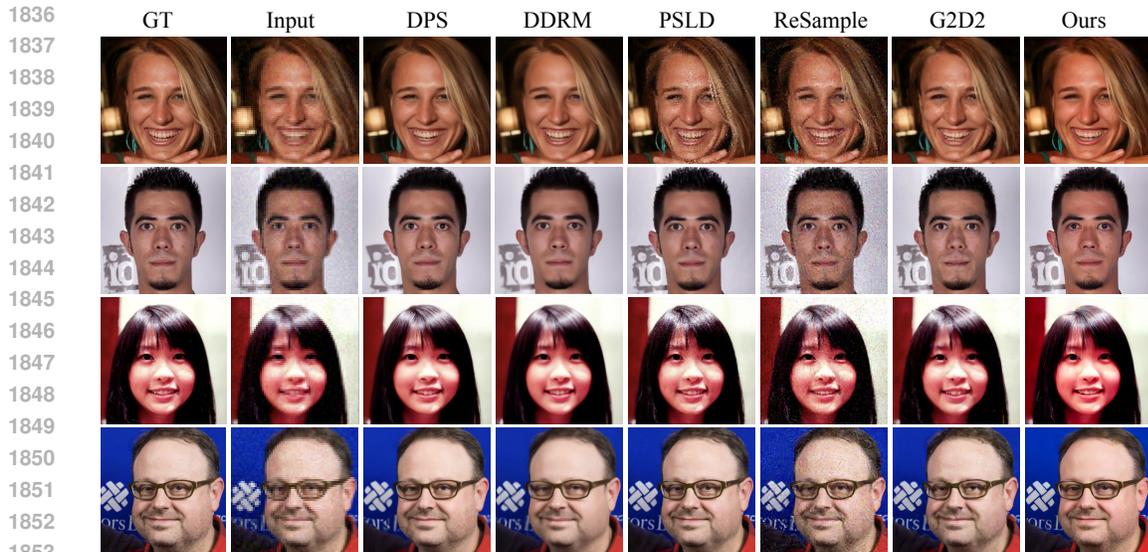
34

GT	Input	DPS	DDRM	PSLD	ReSample	G2D2	Ours

Figure 8: **Additional qualitative results for super resolution** ($4\times$) **on FFHQ.** Continuous baselines (DPS, DDRM, PSLD, ReSample) generate plausible but oversmoothed faces, while the discrete baseline G2D2 often introduces artifacts. In contrast, our APS algorithm reconstructs sharper, more natural faces that closely align with the ground truth. For example, in the second row, our method successfully recovers the small mole on the person's left cheek, a detail overlooked by the baselines.

features with natural skin tones and clean edges, yielding perceptually faithful faces across diverse examples and demonstrating clear advantages for high-resolution face restoration.

Figure 9 presents Gaussian deblurring on FFHQ (Karras et al., 2019). DPS (Chung et al., 2023) and DDRM (Kawar et al., 2022) again oversmooth, suppressing fine facial detail; PSLD (Rout et al., 2023) and ReSample (Song et al., 2024) introduce ringing and plastic-like skin; and G2D2 (Murata et al., 2024) struggles to remove noise from the noisy measurements (Input), creating halos along edges. APS recovers crisp structures such as hair, eyeglass frames, and lip contours while preserving natural highlights and avoiding artifacts, producing reconstructions that are perceptually closer to the ground truth and aligned with our quantitative improvements.

Figure 10 shows additional qualitative results on complex linear and nonlinear inverse problems on FHHQ dataset (Karras et al., 2019), showcasing the performance of both APS and APS-L.

B.9.2 TEXT-GUIDED LARGE BLOCK INPAINTING

Large block inpainting is a particularly challenging setting for generative models, as it requires filling in large missing regions with semantically coherent and high-fidelity content guided by text descriptions. One interesting application of large block inpainting is virtual try-on (Han et al., 2018; Zhu et al., 2024), where models must realistically generate clothing or accessories consistent with both a reference garment and the overall body pose.

In a typical real-world fashion catalog, the full body images naturally have rectangular aspect ratios. Since most existing multimodal foundation models are trained on square images (e.g., 512×512 for MMaDA), we fine-tuned MMaDA on a collection of 1024×512 full-body images curated from a fashion dataset (Zhu et al., 2024), following preprocessing with segmentation-based cropping and padding to standardize framing. This adaptation enables our base model to better handle rectangular image structures. For training, we have randomly selected 100K images from this dataset, providing a diverse and challenging testbed for inpainting at scale.

Comparison. We compare APS against large-scale continuous diffusion baselines: Imagen3 (Baldridge et al., 2024), Flux (Black Forest Labs, 2024), and HDPainter (Manukyan et al., 2025). As illustrated in Figure 11, our method generates realistic clothing textures, with stronger alignment to the reference prompts and fewer artifacts (red boxes highlight failure regions of com-
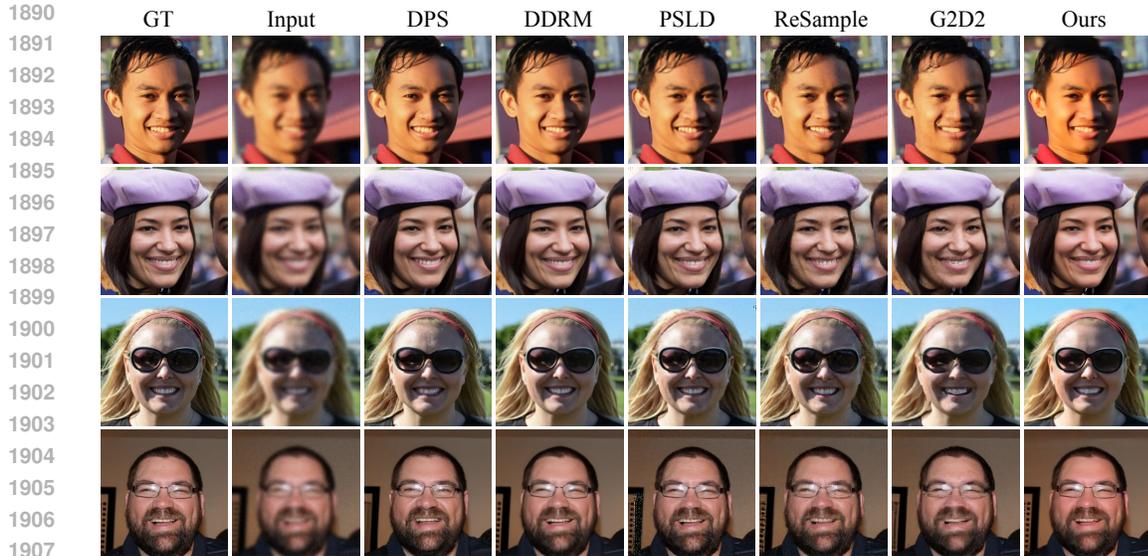
Figure 9: **Additional qualitative results for Gaussian deblurring on FFHQ.** Our proposed approach recovers sharper facial details and edges with fewer artifacts (e.g., reduced ringing and texture distortions), leading to more natural reconstructions. For instance, in the first row, our method accurately reconstructs the hair strands and their shadow on the face, whereas the prior baselines fail to capture these details as precisely.

peting methods). In contrast, Imagen3 and Flux often introduce distorted or inconsistent garment regions, while HD-painter produces less faithful completions with mismatched styles. APS leverages discrete diffusion's ability to directly reweight categorical distributions under posterior guidance, yielding visually compelling and semantically accurate completions.

B.9.3 STYLIZATION

Figure 12 presents additional qualitative results on reference-based stylization. In each case, our APS optimizer conditions on a single reference style image and a text prompt describing the desired content. The outputs demonstrate that APS effectively transfers diverse artistic styles—including tattoo art, steampunk mechanical, psychedelic art, and tribal tattoo—while preserving semantic fidelity to the target content. These results highlight the robustness and versatility of APS in handling a wide range of style-content combinations.

Figure 13 provides a qualitative comparison of our full method (MMaDA + APS) against the base model (MMaDA) and state-of-the-art continuous diffusion approaches. This experiment is designed to demonstrate the novel capability of discrete diffusion models for challenging nonlinear style transfer, not solely to outperform continuous alternatives. We observe that competing methods struggle to balance style fidelity with content alignment. Training-free methods like StyleAligned (Hertz et al., 2023) and InstantStyle (Wang et al., 2024) often drift towards generic textures. Conversely, the training-based StyleDrop (Sohn et al., 2023) tends to overfit to superficial color patterns, which compromises semantic coherence with the text prompt. Our base model, MMaDA (Yang et al., 2025), maintains reasonable content fidelity but fails to transfer fine-grained style attributes, such as material textures or stroke-level details. In contrast, our full method (MMaDA + APS) consistently produces outputs that preserve the reference style while maintaining strong semantic alignment. For instance, our result for the "letter" prompt retains the intricate, flowing smoke design, while the "milkshake" example accurately captures the specified retro diner aesthetic. These results highlight the effectiveness of anchored posterior sampling in discrete diffusion models for complex style transfer tasks.
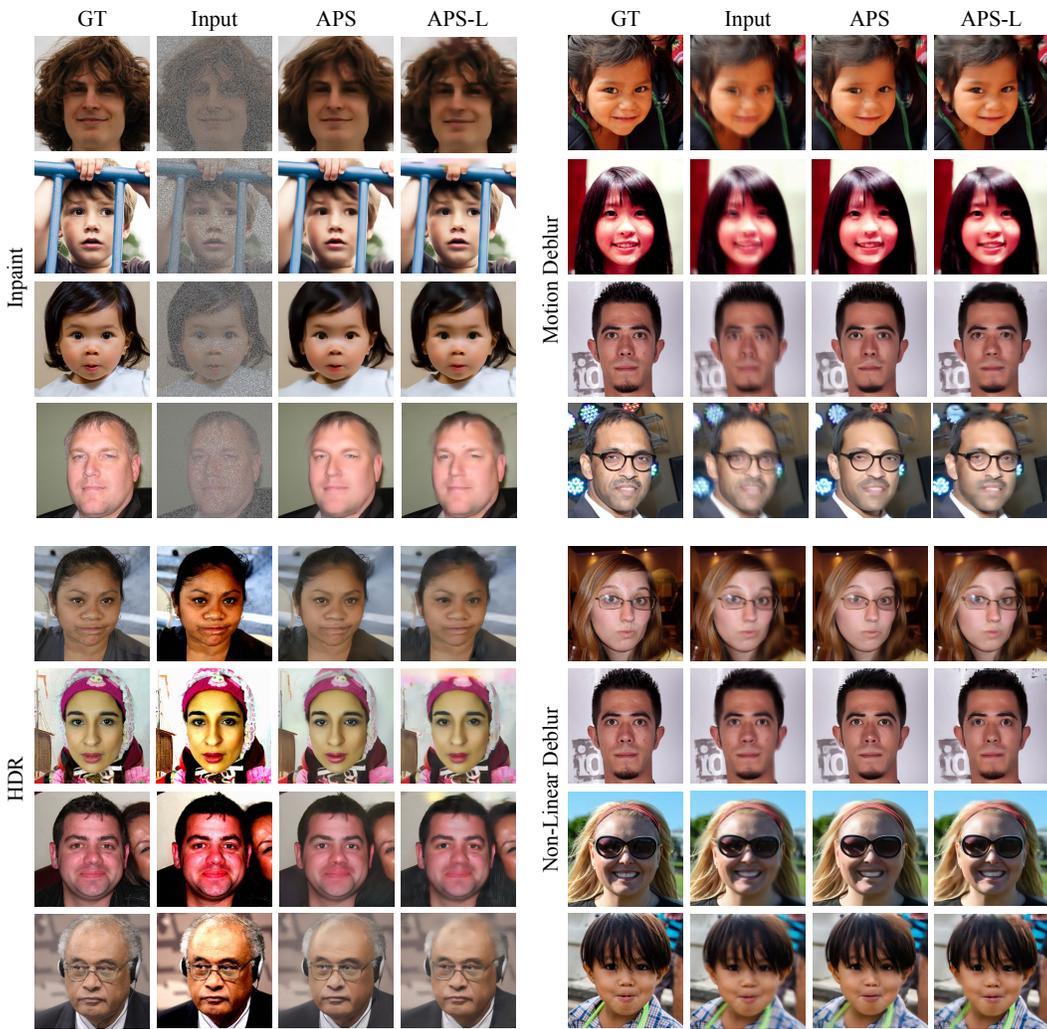
Figure 10: **Qualitative results on FFHQ** for linear (top 4 rows: inpaint and motion deblur) and nonlinear (bottom 4 rows: HDR and non-linear deblur) inverse problems .

Figure 11: **Text-guided large-block inpainting on high-resolution (1024×512) fashion images.** APS generates coherent garment completions with stronger prompt alignment compared to prior methods (Imagen 3 (Baldridge et al., 2024), Flux (Black Forest Labs, 2024), and HD-Painter (Manukyan et al., 2025)). Red boxes highlight incorrect or prompt-inconsistent synthesis in competing approaches. In the first row, the prompt specifies an untucked, oversized T-shirt with vibrant colors—details missed by Imagen 3, Flux, and HD-Painter, respectively. In the second row, our approach correctly buttons the shirt all the way to the top.
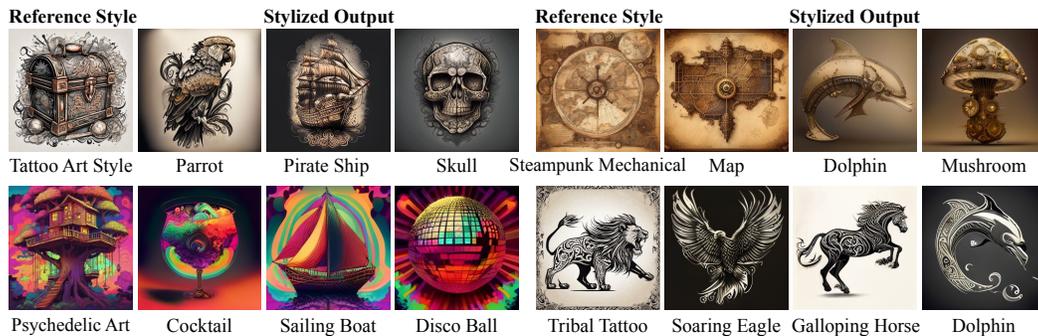

Figure 12: **Additional qualitative results on reference-based stylization.** We show four style-content combinations. For each, our APS optimizer conditions on a single Reference Style image and a text prompt describing the content to generate the Stylized Output images.

B.9.4 Additional Quantitative Results

We perform a larger-scale evaluation for $4\times$ super resolution on FFHQ, extending our analysis to 1000 samples. Our results are compared against the numbers reported for the same task in G2D2 (Murata et al., 2024). Importantly, our observation in the main draft extends to the larger-scale setting and our APS algorithm consistently outperforms G2D2 in all metrics.

B.10 Limitations

Despite achieving state-of-the-art performance among discrete diffusion samplers on (1) complex (linear and nonlinear) inverse problems and (2) reference-based stylization tasks, APS exhibits the following limitations:
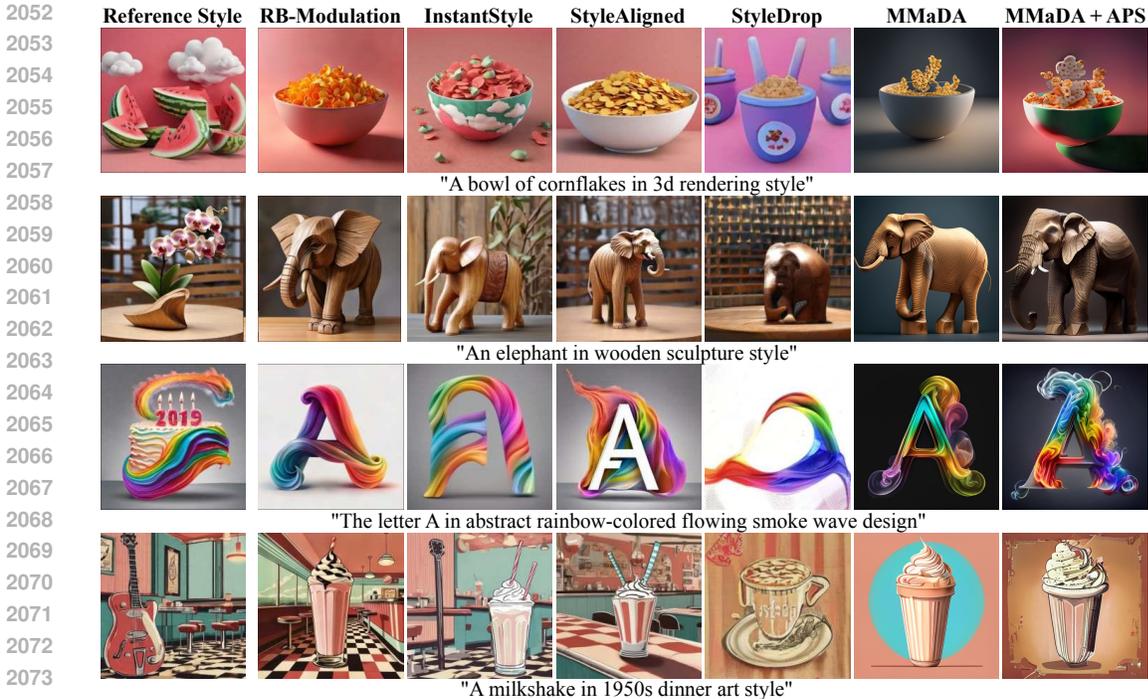
Figure 13: **Additional qualitative comparison on reference-based stylization.** We compare our full method (MMaDA + APS) with the base model (MMaDA) and several state-of-the-art continuous diffusion methods across four style-prompt pairs. For each row, all methods use the same Style Reference image (left) and text prompt (shown below the images).

Table 9: **Super Resolution ($4\times$) on FFHQ.** Performance comparison of APS against prior works. Continuous methods are shaded gray.

Type	Method	LPIPS ↓	PSNR ↑	SSIM ↑
Pixel-domain	DPS	0.238	26.07	0.756
	DDRM	0.252	28.09	0.804
LDM	PSLD	0.282	27.12	0.757
	ReSample	0.508	23.07	0.445
Discrete	G2D2	0.265	27.29	0.763
	G2D2 w/ Markov noise process	0.369	25.15	0.699
Mask	**APS**	**0.232**	**27.81**	**0.808**

1. **Tokenizer quality.** MMaDA uses MagVIT-v2 (Yu et al., 2024) tokenizer which has limited reconstruction quality compared to modern visual tokenizers (Black Forest Labs, 2024). Therefore, future improvements in discrete visual tokenizers could directly benefit APS.

2. **Base model performance.** Discrete diffusion backbones are still in an early stage of development and, at present, underperform large-scale continuous diffusion foundation models such as Flux (Black Forest Labs, 2024), SD3.5 (Esser et al., 2024), and Imagen (Baldridge et al., 2024) in unconditional generative quality. Nevertheless, our theoretical and empirical results indicate that discrete diffusion shows promising potential for posterior sampling and could, with further advances, become a viable alternative to the continuous models that dominate current practice.

3. **Stylization dependence.** The performance of APS in stylization tasks depends both on the pretrained discrete diffusion backbone and the quality of the style feature extractor (e.g., CSD). If the style extractor has not been trained on a particular style, our sampler struggles to transfer it faithfully, limiting its applicability to out-of-distribution styles.

Reference Style	Stylized Output	Reference Style	Stylized Output	Reference Style	Stylized Output	Reference Style	Stylized Output
Geometric Shapes	Rainbow	Infographic	Sofa	Retro Hipster	Coffee Cup	Neon Graffiti	Pair of Rings

Figure 14: **Failure cases of APS.** Under extreme circumstances such as out-of-distribution styles or highly nonlinear measurement operators, our method can sometimes fail, producing over-smoothed reconstructions or noticeable artifacts.

Failure Cases. Figure 14 illustrates failure cases of our approach in stylization. We observe that APS sometimes produces over-smoothed outputs when the reference style is out-of-distribution, or introduces artifacts when the measurement operator is poorly aligned with the pretrained backbone. These examples highlight opportunities for improving robustness and generalization.

B.11 REPRODUCIBILITY STATEMENT

Our experiments are built upon the publicly available MMaDA codebase (Yang et al., 2025). All modifications and implementation details are described in Appendix B.1, which includes the pseudocode in Algorithm 1 and the specific parameters used for every experiment. Furthermore, Appendix B.2 provides comprehensive ablation studies and hyperparameter sweeps (Table 4). The experiments utilize the widely-used public datasets FFHQ (Karras et al., 2019) and ImageNet (Deng et al., 2009). The combination of a public codebase and datasets, along with our detailed Algorithm 1 and parameters (§B.2), should ensure that our results are readily reproducible.

B.12 ETHICS STATEMENT

The proposed method for controlled image editing contributes to the democratization of advanced image editing. While this has positive societal benefits, we acknowledge the dual-use nature of generative models and the potential for misuse or misinterpretation of their outputs.

A primary concern arises in the context of inverse problems such as super resolution, inpainting, and deblurring. Our method generates a plausible, high-quality image, but this output is a sample from a posterior distribution and does not represent a unique or guaranteed reconstruction of the original ground truth. A misunderstanding of this core limitation could lead to dangerous assumptions, particularly in sensitive applications like forensic analysis, where a generated image might be misinterpreted as factual evidence. We therefore emphasize that the method's intended purpose is to enhance perceptual realism for creative or aesthetic applications, and it is not suitable for tasks requiring high-fidelity reconstruction or person identification.

In stylization tasks, there is a risk of unintentional information leakage, where details from the source content image could persist in the stylized output. This could lead to the inadvertent disclosure of sensitive information that a user did not intend to share.