

ROPA : ROBUST PARALLEL DIFFUSION SAMPLING

Anonymous authors

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ABSTRACT

Recent years have witnessed significant progress in developing effective diffusion models. Parallel sampling is a promising recent approach that reformulates the sequential denoising process as solving a system of nonlinear equations, and it can be combined with other acceleration techniques. However, current progress is limited by the trade-off between high fidelity and computational efficiency. This paper addresses the challenge of scaling to high-dimensional, multi-modal generation. Specifically, we present ROPA (Robust Parallel Diffusion Sampling), which takes into account the properties of the denoising process and solves the linear system using adaptive local sparsity to achieve stable parallel sampling. Extensive experiments demonstrate ROPA’s effectiveness: it significantly accelerates sampling across diverse image and video diffusion models, achieving up to $2.9\times$ speedup with eight core, an improvement of 52% over baselines without sacrificing sample quality. ROPA enables parallel sampling methods to provide a solid foundation for real-time, high-fidelity diffusion generation.

1 INTRODUCTION

Over the past few years, the landscape of generative modeling has been significantly reshaped by the ascent of Diffusion Models Ho et al. (2020); Song et al. (2020b). These models have emerged as a pivotal methodology for diverse applications Chung et al. (2023); Yang et al. (2024a); Esser et al. (2024); Ma et al. (2024); Polyak et al. (2025), spanning from high-quality image/video generation to molecular generation. Despite remarkable success, Diffusion Models requires hundreds of sequential denoising steps for generating high-quality samples, each involving expensive neural network evaluations. This sequential dependency severely limits inference speed, particularly for real-time applications and large-scale deployment scenarios. Previous works have explored faster numerical Stochastic differential equations (SDEs) or Ordinary differential equations (ODEs) solvers like DDIM Song et al. (2020a) and DPMSolver Lu et al. (2022), distilling the ODE trajectory into neural networks Salimans & Ho (2022) or straightens trajectories via Rectified Flow Lipman et al. (2023). Others develop sparse-attention and attention cache Zhang et al. (2025); Zou et al. (2025).

Diffusion Models are generative models built on a foundation of two processes: a forward process that systematically corrupts data into noise, and a reverse process that learns to reverse this corruption to generate new data. This dynamic is elegantly described by SDEs. Considering a clean image x_0 sampled from the real data distribution, the forward process gradually perturbs this image with noise over a continuous time interval $t \in [0, T]$, transforming it into a sample x_t that follows a simple prior distribution, like a standard Gaussian. This noising process is defined by the following SDE:

$$dx_t = f(t)x_t dt + g(t) dw, \quad (1)$$

where dw indicates the standard Wiener process. [Although the formulation is expressed in continuous time, in practice we are solving a discrete nonlinear system due to the numerical discretization of the SDE.](#) Then, to generate the corresponding clean latent from the easily sampled random noise, we have to reverse the forward SDE in Eq. 1, resulting in the following reverse SDE formulations:

$$dx_t = \underbrace{[f(t)x_t - g^2(t)\nabla_{x_t} \log p(x_t)]}_{\varphi(x_t, t)} dt + \underbrace{g(t)}_{\sigma_t} dw, \quad (2)$$

where $\nabla_{x_t} \log p(x_t)$ can be approximated by a score function $S_\theta(\cdot)$, parameterized by a neural network with learnable weights of θ ; $\varphi(x_t, t)$ denotes the drift function for the reverse diffusion

process; σ_t represents the corresponding coefficient of diffusion counterpart. Let $\Phi(t, s, x_s)$ represent an integral result of x_t by Eq. 2 over a time interval from s to t , with an initial value x_s :

$$\Phi(t, s, x_s) = x_s + \int_s^t \varphi(x_\tau, \tau) d\tau + \int_s^t \sigma_\tau dw. \quad (3)$$

Consequently, the analytical solution of Eq. 2 at time t can be expressed as

$$x_t = \Phi(t, 0, x_0), \quad x_0 \sim \mathcal{N}(0, I), \quad (4)$$

where $\mathcal{N}(0, I)$ denotes the standard Gaussian distribution.

Formulating Diffusion Sampling to Solving Non-linear Equation. Recent advances in parallel sampling Shih et al. (2024); Tang et al. (2024a); Lu et al. (2025) have shown promise by reformulating the sequential process as solving systems of nonlinear equations, enabling simultaneous computation across multiple timesteps. Existing parallel sampling algorithms establish the following system of non-linear equations to reformulate the integral-based formulation of the diffusion model on a discrete grid $\{t_0, \dots, t_T\}$:

$$x_{t_{n-1}} - \mathcal{F}_n^{(w_n)}(x_{t_n}, \dots, x_{t_{n+w_n-1}}) = 0, \quad (5)$$

where w_n is the window size (number of future states coupled) at step n . $\mathcal{F}_t^{(i)}$ denotes a solver for estimating results in timestamp t with acknowledging previous states, i.e., x_t, \dots, x_{t-i} . The sampling methods utilize an iterative refinement manner to gradually adjust an estimation trajectory $\{\hat{x}_t, t = 0, \dots, T\}$. Each state from the trajectory $\{x_t, t = 0, \dots, T\}$ is first initialized with noise value, denoted as $\{\hat{x}_t^{(0)}, t = 0, \dots, T\}$. Denote by \hat{x}_t the vector, $\hat{x}_{0:T} = [\hat{x}_0^\top, \dots, \hat{x}_T^\top]^\top$. Then, for the k^{th} parallel iteration, where integer $k \in [0, K]$, Newton-Raphson method updates the variables by the following scheme:

$$\hat{x}_{0:T}^{(k+1)} = \hat{x}_{0:T}^{(k)} - \mathcal{G}^{(k)} \mathcal{R}_{0:T}^{(k)}, \quad (6)$$

where $\mathcal{R}_t^{(k)} = \hat{x}_{t-1}^{(k)} - \mathcal{F}_t^{(i)}(\hat{x}_t^{(k)}, \dots, \hat{x}_{t+i}^{(k)})$ indicates a residual term to be optimized; and $\mathcal{G}^{(k)} = (\mathcal{J}^{(k)})^{-1}$ indicates the inverse of Jacobian matrix $\mathcal{J}^{(k)} = \frac{\partial \mathcal{R}_{0:T}^{(k)}}{\partial \hat{x}_{0:T}}$.

Choices of Approximating Jacobian Matrix $\mathcal{J}^{(k)}$. A key strategy for accelerating parallel sampling solvers is to efficiently approximate the Jacobian matrix in the Newton update step, rather than computing the full matrix. Previous methods have employed distinct approximation schemes: *ParaDIGMS* Shih et al. (2024) uses Picard iteration, a fixed-point method that avoids explicit Jacobian computation. This approach is equivalent to approximating the Jacobian of the system as the identity matrix as $\mathcal{J}^{(k)} \approx I$, simplifying the expensive Newton step into a computationally cheap update. *ParaTAA* Tang et al. (2024a) adapts Anderson Acceleration to the problem’s causal structure. Standard acceleration can produce a dense update matrix, which allows well-converged variables to be corrupted by those that have not yet converged. ParaTAA resolves this by enforcing a block upper triangular structure on its update matrix, preserving stability by respecting the natural flow of information in the diffusion process. *ParaSolver* Lu et al. (2025) formulates the problem to have an Jacobian matrix consists of identity blocks on the main diagonal and non-zero blocks only on the sub-diagonal, which reduces the computational and memory costs of each solver iteration. However, current works are all face generalization challenges when scaling to larger scale generation. This leads to the following question that we aim to explore in this work:

Can we dynamically control the the sparsity of Jacobian in parallel diffusion samplers to achieve an optimal trade-off between stability and cost thereby enabling efficient scaling to high-dimensional, multi-modal generation?

Our Contributions. Following the research question, we introduce **ROPA (ROBust PARallel diffusion)**, a novel framework that achieves a superior balance between the efficiency of parallel solving and numerical stability, which scales the application of parallel sampling to complex tasks like video generation. Our key contributions are:

(a) Scaling To High-Dimensional Generation. Our geometric analysis in Section. 2 rigorously establishes the mechanism behind mode collapse in parallel diffusion samplers. We later show (Sec. 2) that highly curved regions of the data density naturally induce stiff score dynamics and ill-conditioned

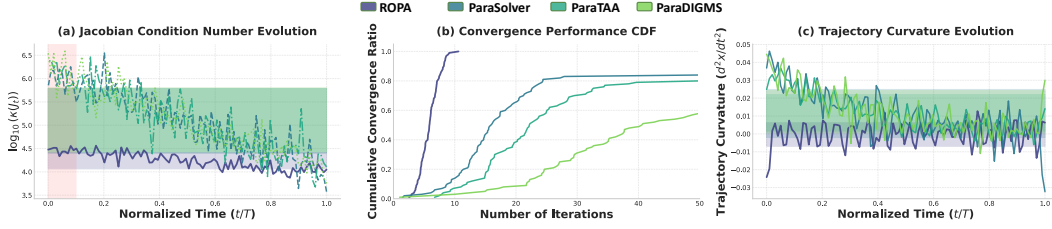


Figure 1: ROPA Performance Analysis. (a) Jacobian condition number evolution showing ROPA’s superior numerical stability. (b) Convergence performance CDF demonstrating ROPA’s faster and more reliable convergence. (c) Trajectory curvature evolution highlighting ROPA’s geometric consistency. Shaded regions represent mean $\pm 1\sigma$ uncertainty bounds. The red shaded area in (a) indicates the high-curvature region where numerical challenges are most severe.

Jacobians for parallel solvers, explaining the observed instability of existing methods. Current parallel sampling methods, as shown in Figure. 1, struggle near $t \rightarrow 0$, where the Jacobian condition number $\kappa(\mathcal{J}_t)$ grows exponentially, causing Newton iterations to diverge. This leads to unreliable convergence—many trajectories fail to converge within practical budgets, while others require excessive steps. Crucially, these instabilities cause trajectories to deviate from the data manifold \mathcal{M} , particularly near multimodal boundaries where non-adaptive methods generate inconsistent samples that unrealistically interpolate between modes. ROPA solves this by dynamically regulating $\kappa(\mathcal{J}_t)$ through adaptive damping and sparsity, maintaining numerical stability, geometric fidelity, and mode consistency even in high-curvature regions.

b) We propose *Geometry-Aware Adaptive Jacobian Sparsity Control*. Specifically we translate geometric curvature signals into on-the-fly control of the solver’s coupling structure. At each iteration, the method selectively widens the look-ahead only where residuals indicate stiffness and prunes it elsewhere, preserving $O(N)$ parallelism while concentrating computation where it matters most. When diagnostics flag instability, an adaptive damping mechanism automatically moderates the update—behaving like fast Newton steps in well-conditioned regions and shifting toward conservative descent near ill-conditioning. Together, these two levers keep the Jacobian well-conditioned under a target threshold, deliver reliable convergence in high-curvature areas where prior methods struggle, and scale to large, multi-modal generation without extra training or ad-hoc heuristics.

c) Extensive experiments demonstrate substantial speedups on Stable Diffusion-v3.5, FLUX, HunyuanVideo, Wan2.1 and CogVideoX while maintaining FID and CLIP scores.

2 A UNIFIED GEOMETRIC ANALYSIS OF PARALLEL SAMPLING INSTABILITY IN DIFFUSION MODELS

We establish a framework linking data manifold geometry, discretization effects, and numerical stability in parallel diffusion sampling, which reveals why mode interpolation collapses emerge in high-curvature regions of the data manifold and how adaptive Jacobian control mitigates them.

2.1 GEOMETRIC FOUNDATIONS OF SCORE STIFFNESS AND DENSITY CURVATURE

We use the term *curvature* in a probabilistic rather than purely geometric sense. Concretely, we define the *density curvature* at x via the Hessian of the log-density,

$$\mathcal{H}(x) = \nabla_x^2 \log p(x),$$

which measures how sharply the probability mass bends around the data manifold \mathcal{M} . This is distinct from intrinsic Riemannian curvature of \mathcal{M} : in our setting, $\mathcal{H}(x)$ controls the stiffness of the score field and, through our analysis, the conditioning of the parallel residual Jacobian.

The core challenge stems from the data manifold’s intrinsic curvature properties. Let $\mathcal{M} \subset \mathbb{R}^d$ denote the support of $p_0(x)$, with curvature characterized by the score Hessian $\mathcal{H}(x) = \nabla_x^2 \log p(x)$. The eigenvalues of $\mathcal{H}(x)$ quantify how sharply the density bends in different directions. A large ratio between the largest and smallest eigenvalues, corresponding to high anisotropy, means that the score

changes very quickly along some directions but slowly along others, this is precisely the notion of stiffness that leads to ill-conditioned Jacobians in our parallel residual system.

Assumption 2.1 (Manifold Anisotropy Index). Let $\mathcal{M} \subset \mathbb{R}^d$ be the data manifold and $\mathcal{H}(x) = \nabla_x^2 \log p(x)$ its score Hessian. For any $x \in \mathcal{M}$, denote the ordered eigenvalues by $0 \leq \nu_1(x) \leq \dots \leq \nu_d(x)$ (we reserve λ for damping parameters). Fix a small constant $\varepsilon > 0$. Define the *local anisotropy index*

$$\rho(x) := \frac{\nu_d(x)}{\max\{\nu_1(x), \varepsilon\}}.$$

We assume $\rho(x)$ is locally Lipschitz on \mathcal{M} and may take large values $\rho(x) \gg 1$ only on a *measurable subset* $\mathcal{M}_{\text{curv}} \subset \mathcal{M}$ corresponding to high-curvature regions.

Theorem 2.2 (Lower Bound on the Denoiser Jacobian). *Let $r_\theta(x, t)$ be a trained denoiser that satisfies $\|r_\theta(x, t) - \mathbb{E}[x_0 | x_t = x]\|_2 \leq \varepsilon$ uniformly. Under Assumption 2.1 and assuming $\mathcal{H}(x) \succeq 0$, the spatial Jacobian $J_{r_\theta}(x, t) = \partial r_\theta(x, t) / \partial x$ obeys, for any $t \in (0, T]$,*

$$\|J_{r_\theta}(x, t)\|_2 \geq 1 + \sigma_t^2 \nu_1(x) - \mathcal{O}(\varepsilon),$$

where $\nu_1(x)$ is the smallest non-negative eigenvalue of $\mathcal{H}(x)$. If \mathcal{H} has negative directions, Eq. 2.2 still holds with $\nu_1(x)$ replaced by $|\nu_{\min}(x)|$.

2.2 DISCRETIZATION-INDUCED INSTABILITY

The residual system in parallel sampling is defined as $\mathcal{R}_n^{(k)} = \hat{x}_{t_{n-1}}^{(k)} - \mathcal{F}_{t_n}^{(i)}(\hat{x}_{t_n}^{(k)}, \dots, \hat{x}_{t_{n+i}}^{(k)})$ per Eq. 6, where for Euler integration:

$$\mathcal{F}_{t_n}^{(i)}(x_{t_n}, \dots, x_{t_{n+i}}) = x_{t_n} - \Delta \varphi(x_{t_n}, t_n), \quad \Delta = t_{n+1} - t_n. \quad (7)$$

This discretization introduces gaps between continuous and discrete dynamics:

Theorem 2.3 (Condition Number of the Parallel Residual Jacobian). *Let $\mathcal{R}_{0:T}^{(k)}$ be the residual vector defined in Eq. 6 with a uniform step size $h = t_{n+1} - t_n$. Write $\mathcal{J}^{(k)} = I + hA^{(k)}$ where $A^{(k)}$ collects blocks depending on $J_{r_\theta}(x, t)$ and the drift $f(t)$. Assume $A^{(k)}$ is block-row diagonally dominant and $\|A^{(k)}\|_2 \leq L$ for some Lipschitz constant L . Then for any $h < h_{\max} := 1/L$,*

$$\kappa(\mathcal{J}^{(k)}) \leq 1 + \frac{hL}{1 - hL} = 1 + \mathcal{O}(h).$$

In particular, substituting $L = \sigma_t^2 \|J_{r_\theta}(x, t)\|_2$ yields

$$\kappa(\mathcal{J}^{(k)}) \leq 1 + c \cdot h \sigma_t^2 \|J_{r_\theta}(x, t)\|_2 + \mathcal{O}(h^2),$$

where the constant $c = \frac{2}{1-hL} \approx 2$ arises from the Neumann series expansion of the inverse Jacobian in Appendix C.3).

This establishes the *geometric-numerical instability cascade*: high curvature $\lambda_{\min}(\mathcal{H})$ increases $\Rightarrow \|J_{r_\theta}\|_2$ increases $\Rightarrow \kappa(\mathcal{J})$ increases \Rightarrow solver divergence. Crucially, this cascade is triggered by local geometric properties of the data manifold, not by temporal proximity to $t = 0$.

2.3 TRAJECTORY GEOMETRY AND MODE COLLAPSE

The stability loss manifests geometrically. Following Davies & Powell (1984); Chen & Muñoz Ewald (2023), define trajectory quasi-linearity via $\|d^2x/dt^2\|_2 \leq \varepsilon$. We prove:

Corollary 2.4 (Numerical Stability & Manifold Deviation). *Under the same hypotheses as Theorem 2.3, let \hat{x} be the iterate returned by one Newton step and $x^* = \text{Proj}_{\mathcal{M}}(\hat{x})$ its orthogonal projection onto \mathcal{M} . Then*

$$\|\hat{x} - x^*\|_2 \leq (\kappa(\mathcal{J}^{(k)}) - 1) \|\mathcal{J}^{(k)-1} \mathcal{R}^{(k)}\|_2 + \mathcal{O}(\|\mathcal{R}^{(k)}\|_2^2). \quad (8)$$

Hence, if $\kappa(\mathcal{J}^{(k)})$ grows large, the forward error increases proportionally. Proof is an adaptation of the classical backward-forward error bound (Davies & Powell, 1984).

This resolves the central paradox: high-fidelity generation *requires* large $\|J_{r_\theta}\|_2$ (to separate modes), but this same property destabilizes parallel solvers (Theorem 2.3). Crucially, instability peaks at mode boundaries where $\lambda_{\min}(\mathcal{H})$ spikes:

Corollary 2.5 (Boundary Sensitivity under Gaussian Mixture). *Consider a Gaussian-mixture density $p_0(x) = \sum_m w_m \mathcal{N}(\mu_m, \Sigma_m)$ whose decision boundary $\partial\mathcal{M}$ is the union of quadratic surfaces. For x in the normal direction of a boundary component, let $\delta = \text{dist}(x, \partial\mathcal{M})$. Then $\lambda_{\min}(\mathcal{H}(x)) = \Theta(\delta^{-1})$ and $\kappa(\mathcal{J}^{(k)}) \rightarrow \infty$ as $\delta \rightarrow 0$.*

Theoretical results reveal that the condition number $\kappa(\mathcal{J}_t)$ —modulated by data-manifold curvature, score stiffness, and discretisation step size—is the key scalar that couples geometric fidelity and numerical stability. To translate this insight into a practical sampler, we introduce three *geometry-aware control principles* that directly regulate $\kappa(\mathcal{J}_t)$ during the Newton-type parallel updates. Each principle is summarized as below.

Takeaways 2.6 (Damped Updates for Safety). At iteration k , choose $\lambda_k > 0$ such that the gain ratio $\rho_k = \frac{\|\mathcal{R}^{(k)}\|_2 - \|\mathcal{R}^{(k+1)}\|_2}{\Delta x^{(k)\top} (\lambda_k \Delta x^{(k)} - \mathcal{R}^{(k)})}$ satisfies the trust-region criterion (Davies & Powell, 1984). Then the update $(\mathcal{J}^{(k)} + \lambda_k I) \Delta x^{(k)} = -\mathcal{R}^{(k)}$ is globally convergent.

Takeaways 2.7 (Adaptive Sparsity for Efficiency). Let \mathcal{S}_k be a block-band sparsity pattern whose bandwidth b_k is chosen via $b_k = \min\{b : \text{iters}(\mathcal{J}_{|b}^{(k)}) \leq M\}$, where iters estimates Conjugate-Gradient iterations with Jacobian-vector products only. This guarantees expected complexity $\mathcal{O}(Nb_k)$ per Newton step while keeping $\kappa(\mathcal{J}_{|b_k}^{(k)}) \leq \gamma^{-1}$ for a target γ .

Takeaways 2.8 (Low-Rank Curvature Correction for Fidelity). Given a subspace basis $U \in \mathbb{R}^{d \times r}$ corresponding to the top- r eigenvectors of $\mathcal{H}(x)$ with eigenvalues Λ_r , apply the correction $x \leftarrow x - U(\Lambda_r + \tau I)^{-1} U^\top \nabla_x \log p(x)$, where $\tau > 0$ regularises near-singular directions. This preserves local manifold structure up to $\mathcal{O}(\tau)$.

Summary The data manifold’s curvature (Assumption 2.1) dictates score stiffness (Theorem 2.2), which discretization gaps amplify (Theorem 2.3). This causes trajectories to deviate from \mathcal{M} at mode boundaries (Corollary 2.5), thus generation collapse. Crucially, these instabilities occur wherever the sampling trajectory enters high-curvature regions or approaches mode boundaries. ROPA’s adaptive mechanisms directly counter this cascade by regulating $\kappa(\mathcal{J})$ based on local geometry, enabling stable high-fidelity sampling throughout the entire diffusion process. See Appendix C for proof.

3 ROBUST PARALLEL DIFFUSION SAMPLING VIA ADAPTIVE JACOBIAN SPARSITY

Building on the geometric cascade characterization introduced in Section 2, we aim to regulate the Jacobian condition number $\kappa(\mathcal{J}_t)$. While Theorem 2.2 links instability to the Hessian eigenvalues $\nu_i(x)$, explicitly computing curvature at inference time is computationally prohibitive. However, Theorem 2.3 implies that high curvature induces stiff, long-range temporal dependencies. When the solver’s look-ahead window is too narrow to capture these dependencies, the Jacobian approximation suffers high truncation error, manifesting immediately as large local residuals $\|\mathcal{R}\|$.

Therefore, ROPA utilizes the residual norm as a computationally cheap proxy for local geometric stiffness, driving two complementary operating modes: (i) adaptive sparsification of residual couplings, which maintains computational efficiency when local curvature is moderate; (ii) targeted curvature correction, which enhances stability as soon as geometric diagnostics reveal elevated risk.

Let $N := T + 1$ denote the total number of discrete time points on the grid $\{t_0, \dots, t_T\}$.

3.1 DYNAMIC RESIDUALS WITH ADAPTIVE JACOBIAN BANDWIDTH

At each grid index $i \in \{1, \dots, T\}$ the algorithm selects a forward-looking window width $w_i \in \{1, \dots, w_{\max}\}$ and forms the residual

$$\mathcal{R}_i^{(w_i)}(\hat{\mathbf{x}}) = \hat{\mathbf{x}}_{t_{i-1}} - \Psi_i^{(w_i)}(\hat{\mathbf{x}}_{t_i}, \dots, \hat{\mathbf{x}}_{t_{i+w_i-1}}), \quad (9)$$

where $\Psi_i^{(w_i)}$ approximates the integral operator $\Phi(t_{i-1}, t_i, \hat{\mathbf{x}}_{t_i})$ by means of an explicit w_i -step integrator (e.g., Euler, DDIM, or a higher-order variant). This look-ahead construction yields an upper-banded Jacobian:

$$\frac{\partial \mathcal{R}_i^{(w_i)}}{\partial \hat{\mathbf{x}}_{t_j}} = \begin{cases} \mathbf{I}_d, & j = i - 1, \\ -\frac{\partial \Psi_i^{(w_i)}}{\partial \hat{\mathbf{x}}_{t_j}}, & i \leq j \leq i + w_i - 1, \\ \mathbf{0}, & \text{otherwise.} \end{cases} \quad (10)$$

For first-order integrators this structure guarantees block-row diagonal dominance. Higher-order schemes may weaken that dominance; a locally scaled damping parameter $\lambda_{\text{damp}, i}$, described in Section 3.2, restores stability in that case.

Adaptive bandwidth control. During Newton iteration k , the algorithm evaluates local residual norms

$$e_i^{(k)} = \|\mathcal{R}_i^{(w_i^{(k)})}(\hat{\mathbf{x}}^{(k)})\|_2 \quad (11)$$

and their global mean

$$\bar{e}^{(k)} = N^{-1} \sum_{i=0}^T e_i^{(k)}. \quad (12)$$

Following **Theorem 2.3**, a high local residual $e_i^{(k)}$ indicates that the current sparse Jacobian approximation fails to capture the stiff, long-range temporal couplings induced by high curvature. To counter this, we dynamically adjust the window widths to regulate the truncation error:

$$w_i^{(k+1)} = \begin{cases} \min\{w_i^{(k)} + 1, w_{\max}\}, & e_i^{(k)} > \alpha \bar{e}^{(k)} \quad (\text{densify to capture stiffness}), \\ \max\{w_i^{(k)} - 1, 1\}, & e_i^{(k)} < \beta \bar{e}^{(k)} \quad (\text{sparsify for efficiency}), \\ w_i^{(k)}, & \text{otherwise,} \end{cases} \quad (13)$$

with default parameters $\alpha = 1.5$ and $\beta = 0.7$.

By *densifying* the block-banded Jacobian (increasing w_i) only in high-error regions, this update rule implicitly lowers the local condition number $\kappa(\mathcal{J})$ bounded in Theorem 2.3. This ensures geometric stability without incurring the cubic cost of a fully dense solver, trading off sparse $\mathcal{O}(N)$ operations only where geometrically necessary.

3.2 LML-BASED LOW-RANK CURVATURE CORRECTION

While adaptive bandwidth handles general stiffness, it cannot resolve the topological singularities described in **Corollary 2.5**, where $\kappa(\mathcal{J}) \rightarrow \infty$ at decision boundaries. In these regimes, the Jacobian becomes near-singular along the normal direction of the manifold, and simply widening the window is insufficient.

To detect this, we monitor the alignment between the residual \mathcal{R}_i and the score \mathbf{s}_θ , which acts as a proxy for the principal curvature direction (eigenvector of the largest Hessian eigenvalue ν_{\max}). Alignment is declared whenever:

$$\frac{|\langle \mathcal{R}_i, \mathbf{s}_\theta \rangle|}{\|\mathcal{R}_i\|_2 \|\mathbf{s}_\theta\|_2} > \gamma, \quad (14)$$

where γ is a sensitivity threshold. When this geometric trigger activates, the algorithm invokes a curvature-aware correction inspired by preconditioned Langevin dynamics. We define the rank-one inverse Hessian approximation as:

$$\mathbf{H}_{\text{LML}}^{-1}(\hat{\mathbf{x}}_t, t; \lambda_{\text{damp}}) = \frac{1}{\lambda_{\text{damp}} g(t)^2 \|\mathbf{s}_\theta\|_2^2} \left(\mathbf{I}_d - \frac{\mathbf{s}_\theta \mathbf{s}_\theta^\top}{\lambda_{\text{damp}} + \|\mathbf{s}_\theta\|_2^2} \right), \quad (15)$$

where $g(t)$ denotes the diffusion coefficient. All eigenvalues remain positive for any $\lambda_{\text{damp}} > 0$, ensuring a positive-definite operator. This construction mirrors a single step of preconditioned Langevin dynamics with step size $1/\lambda_{\text{damp}}$, explicitly injecting curvature information along the stiff score direction while preserving the orthogonal subspace.

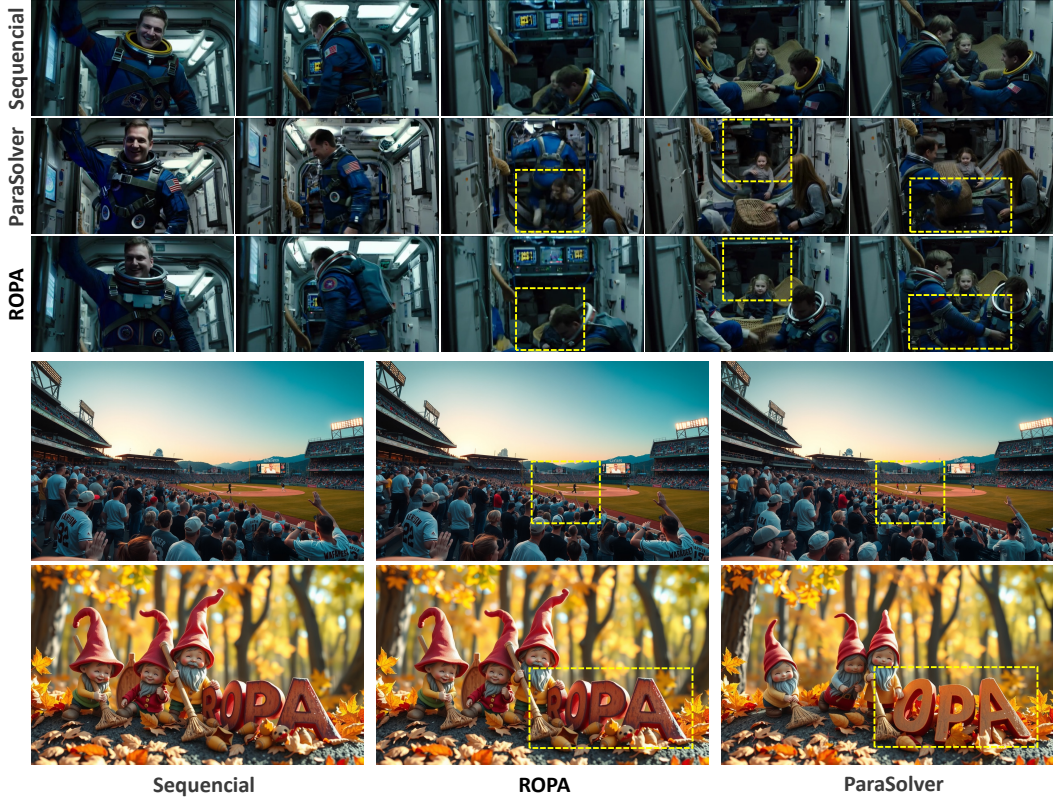


Figure 2: Quality Comparison of ROPA and baselines on HunyuanVideo and Flux models.

Jacobian blocks for the affected grid indices are updated as

$$\mathbf{B}_i^{(k)} = \begin{bmatrix} -\mathbf{H}_{\text{LML}}^{-1}(\hat{\mathbf{x}}_{t_i}^{(k)}, t_i; \lambda_{\text{damp}, i}^{(k)}) & \mathbf{I}_d \end{bmatrix}. \quad (16)$$

Because only the score vector \mathbf{s}_θ is stored, memory remains $\mathcal{O}(Nw_{\max}d)$. This low-rank correction explicitly enforces the bound $\kappa(\mathcal{J}_t) \leq \kappa_{\text{th}}$ by removing ill-conditioning along high-curvature directions indicated by the score. The damping parameter $\lambda_{\text{damp}, i}^{(k)}$ is tuned so that the stability criterion of Corollary 2.4 continues to hold.

4 EXPERIMENTS

4.1 SETUPS

Models. For video generation, we benchmark on three state-of-the-art large video diffusion models: HunyuanVideo Kong et al. (2024) and CogVideoX1.5-5B Yang et al. (2024b). For each model, we generate videos with prompts in VBench Huang et al. (2024) strictly following VBench evaluation protocol. We consider two image diffusion models for image generation, Stable Diffusion 3.5 Large Esser et al. (2024) and Flux Labs (2024), as the backbone. Following previous works Shih et al. (2024); Selvam et al. (2024), we sample 1000 prompts from COCO2017 captions dataset as the test bed. We use $N = 50$ diffusion steps by default, with more investigations on N in Appendix A.3.

Algorithms. We benchmark our proposed algorithm, ROPA, against five key baselines: (1) *Sequential Sampling*, the standard non-parallel approach which serves as the reference for performance speedups; (2) *ParaDiGMS* Shih et al. (2023), a foundational parallel method utilizing Picard (fixed-point) iteration; (3) *ParaTAA* Tang et al. (2024b), which accelerates convergence by applying Triangular Anderson Acceleration (TAA) to a dense nonlinear system; (4) *ParaSolver* Lu et al. (2025), a highly efficient method that combines a quasi-Newton solver with a sparse, banded system structure; and (5) *CHORDS* Han et al. (2025), a parallel framework designed for robust and stable convergence.

Algorithm 1 *ROPA: Robust Parallel Diffusion Sampling*

Require: Denoiser S_θ , grid $\{t_n\}_0^T$, max iter K , thresholds α, β, γ , tol ε .
Ensure: Clean sample $x_{t_0}^{(K)}$.

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1: Initialize trajectory  $\{\hat{x}_{t_n}^{(0)}\}$  with noise;  $w_n^{(0)} \leftarrow 1$ ;  $\lambda_n^{(0)} \leftarrow \lambda_{\text{init}}$ .
2: for  $k = 0$  to  $K - 1$  do
3:   Residual Eval (Parallel):
4:   for  $n = 1 \dots T$  do
5:      $\mathcal{R}_n^{(k)} \leftarrow \hat{x}_{t_{n-1}}^{(k)} - \Psi_n^{(w_n^{(k)})}(\hat{x}_{t_n}^{(k)}, \dots, \hat{x}_{t_{n+w_n^{(k)}}}^{(k)})$ 
6:     Compute norms  $e_n^{(k)}$  and global mean  $\bar{e}^{(k)}$ .
7:   end for
8:   if  $\bar{e}^{(k)} < \varepsilon$  then break
9:   end if
10:  Adaptive Bandwidth:
11:  for  $n = 1 \dots T$  do
12:    if  $e_n^{(k)} > \alpha \bar{e}^{(k)}$  then
13:       $w_n^{(k+1)} \leftarrow \min(w_n^{(k)} + 1, w_{\text{max}})$  ▷ Densify
14:    else if  $e_n^{(k)} < \beta \bar{e}^{(k)}$  then
15:       $w_n^{(k+1)} \leftarrow \max(w_n^{(k)} - 1, 1)$  ▷ Sparsify
16:    else
17:       $w_n^{(k+1)} \leftarrow w_n^{(k)}$ 
18:    end if
19:  end for
20:  Curvature Correction (Parallel):
21:  for  $n = 1 \dots T$  do
22:    Compute alignment  $\rho_n$  between  $\mathcal{R}_n^{(k)}$  and score  $s_n$ .
23:    if  $\rho_n > \gamma$  then
24:       $\tilde{\mathcal{R}}_n^{(k)} \leftarrow H_n^{-1} \mathcal{R}_n^{(k)}$  ▷ Using LML-based preconditioner
25:    else
26:       $\tilde{\mathcal{R}}_n^{(k)} \leftarrow \mathcal{R}_n^{(k)}$ 
27:    end if
28:  end for
29:  4. Update: Assemble  $\mathcal{J}^{(k)}$  using  $\{w_n^{(k+1)}\}$ .
30:  for  $n = 1 \dots T$  do
31:    Solve  $(\mathcal{J}^{(k)} + \lambda_n^{(k)} I) \Delta x_{t_n}^{(k)} \approx \tilde{\mathcal{R}}_n^{(k)}$  via trust-region damping.
32:     $\hat{x}_{t_n}^{(k+1)} \leftarrow \hat{x}_{t_n}^{(k)} - \Delta x_{t_n}^{(k)}$ 
33:  end for
34: end for
35: return  $\hat{x}_{t_0}^{(K)}$ 

```

Hyperparameter Settings. **Damping λ :** Follow L-curve rule—start at 10^{-3} , adjust by factor 2 until $\|\delta \hat{x}\| / \|\mathcal{R}\| < 0.3$, directly controlling $\kappa(\mathcal{J})$ as discussed in the introduction. **Prune factor η :** Set to 0.1 for images, 0.2 for videos (robust in $[0.05, 0.3]$), with threshold γ explicitly using $\|S_\theta\|^2$ from the introduction. **Adaptation thresholds:** $\alpha = 1.5$, $\beta = 0.7$ provide optimal sparsity-stability balance, directly addressing the convergence degradation near $t \rightarrow 0$ observed in the introduction. This configuration enables stable high-fidelity generation at scale while maintaining $O(N)$ parallelism—even in high-curvature regions where existing methods fail, as empirically demonstrated in the introduction’s Figures 1.

Settings. We run experiments using 8 * H200 GPUs, each with 140GB of memory. In all scenarios, we employ classifier-free guidance with a guidance scale of 5. The window-scaled variant halves the number of synchronization rounds compared with a fixed λ . For all algorithms, we use the same stopping threshold $\varepsilon_t = \tau^2 g^2(t) d$ with $\tau = 10^{-3}$, and initialize all variables with standard Gaussian Distribution and warming-up steps set as 3.

Evaluation. For both video and image models, we report *Time per sample* that refers to the average wall-clock time used to generate one sample. *Speedup* that refers to the relative speedup compared with sequential solve, measured by the number of sequential network forward calls. Notice that this will be slightly different from the measurement or the wall-clock. In terms of generation quality, we report average of diverse *Quality Scores* (Clarity, Aesthetic, Motion, Dynamic, Semantic, Anatomy, Identity) normalized using the same numerical system as the standard quality metric following the VBench evaluation protocol Huang et al. (2024) for video generation, and *CLIP Score* Hessel et al. (2021) evaluated using ViT-g-14 Radford et al. (2021); Ilharco et al. (2021) for the image generation. We also report *Latent RMSE* under both cases that measures the Rooted MSE between the returned

Table 1: Benchmark results of parallel diffusion methods on video diffusion models using VBench. We evaluate on three video diffusion models with the number of cores K set to 2, 4 and 8. Our approach achieves the highest speedup without measurable quality degradation.

		Num Core = 2				Num Core = 4				Num Core = 8			
		Time(s)	Speedup	Quality $_V$	RMSE $_L$	Time(s)	Speedup	Quality $_V$	RMSE $_L$	Time(s)	Speedup	Quality $_V$	RMSE $_L$
HunyuanVideo	Sequential	378.6	-	73.8%	-	378.6	-	73.8%	-	378.6	-	73.8%	-
	CHORDS	292.3	1.3	73.6%	0.188	185.5	2.0	73.7%	0.182	156.0	2.4	73.7%	0.185
	ParaDIGMS	313.3	1.2	73.7%	0.190	293.1	1.3	73.6%	0.175	271.8	1.4	73.6%	0.189
	ParaTAA	318.6	1.2	73.6%	0.055	207.0	1.8	73.6%	0.055	157.1	2.4	73.6%	0.055
	ParaSolver	287.5	1.3	73.5%	0.051	208.1	1.8	73.5%	0.049	164.7	2.3	73.5%	0.052
	ROPA (Ours)	232.8	1.6	73.6%	0.054	177.9	2.1	73.6%	0.053	131.8	2.9	73.6%	0.055
Wan2.1	Sequential	471.2	-	74.7%	-	471.2	-	74.7%	-	471.2	-	74.7%	-
	CHORDS	362.8	1.3	74.5%	0.082	274.9	1.7	74.6%	0.076	197.0	2.4	74.6%	0.079
	ParaDIGMS	395.1	1.2	74.5%	0.077	332.6	1.4	74.6%	0.070	279.6	1.7	74.6%	0.084
	ParaTAA	338.2	1.4	74.5%	0.030	312.9	1.5	74.5%	0.028	202.1	2.3	74.5%	0.028
	ParaSolver	340.2	1.4	74.5%	0.025	293.2	1.6	74.5%	0.024	185.2	2.5	74.5%	0.026
	ROPA (Ours)	274.0	1.7	74.5%	0.027	250.8	1.9	74.5%	0.021	169.1	2.8	74.5%	0.030
CogVideoX1.5	Sequential	464.5	-	71.3%	-	464.5	-	71.3%	-	464.5	-	71.3%	-
	CHORDS	389.5	1.2	71.0%	0.132	246.3	1.9	71.1%	0.125	221.5	2.1	71.0%	0.129
	ParaDIGMS	390.9	1.2	71.0%	0.146	356.3	1.3	71.0%	0.119	290.7	1.6	70.9%	0.174
	ParaTAA	359.9	1.3	70.9%	0.043	388.0	1.2	70.9%	0.043	224.1	2.1	70.9%	0.043
	ParaSolver	332.4	1.4	71.0%	0.040	386.9	1.2	71.1%	0.039	207.5	2.2	71.0%	0.041
	ROPA (Ours)	307.5	1.5	71.1%	0.041	219.5	2.1	71.2%	0.041	182.4	2.5	71.2%	0.042

Table 2: Benchmark results of parallel diffusion methods on latent image diffusion models. We evaluate two models with 1000 prompts from the COCO2017 captions dataset. Our approach achieves the highest speedup without measurable quality degradation.

		Num Core = 2				Num Core = 4				Num Core = 8			
		Time(s)	Speedup	CLIP	RMSE $_L$	Time(s)	Speedup	CLIP	RMSE $_L$	Time(s)	Speedup	CLIP	RMSE $_L$
SD-3.5-Large	Sequential	10.3	-	37.4	-	10.3	-	37.4	-	10.3	-	37.4	-
	ParaDIGMS	7.6	1.4	37.2	0.440	7.7	1.3	37.4	0.346	7.1	1.5	37.4	0.342
	ParaSolver	6.8	1.5	37.4	0.234	9.4	1.1	37.4	0.294	5.8	1.8	37.3	0.324
	ROPA (Ours)	6.3	1.6	37.4	0.141	5.8	1.8	37.4	0.220	5.2	2.0	37.4	0.224
Flux	Sequential	11.2	-	37.4	-	11.2	-	37.4	-	11.2	-	37.4	-
	ParaDIGMS	8.1	1.4	37.4	0.249	7.2	1.6	37.4	0.121	7.3	1.5	37.4	0.313
	ParaSolver	6.4	1.7	37.3	0.270	6.6	1.7	37.4	0.166	5.5	2.0	37.4	0.150
	ROPA (Ours)	5.8	1.9	37.4	0.154	5.3	2.1	37.4	0.143	4.8	2.3	37.4	0.120

latent of the algorithm and that of the sequential solver. Notice that a lower latent RMSE indicates lower sampling error, with sequential solve being the oracle.

4.2 MAIN RESULTS

Video diffusion acceleration Our proposed ROPA, demonstrates a clear superiority across all tested models in Table. 1. At the highest level of parallelism with 8 cores, ROPA achieves remarkable speedups ranging from $2.5\times$ to $2.9\times$. On the HunyuanVideo model, it reduces the generation time from 378.6s to just 131.8s, a $2.9\times$ acceleration. This significant performance gain is achieved without any meaningful degradation in output quality. The VBench Quality score remains exceptionally stable, 73.6% for HunyuanVideo vs. 73.8% for the sequential baseline, and the Latent RMSE is kept to a minimum. Notably, ROPA’s Latent RMSE of 0.055 is not only competitive with the best-performing baselines but is also nearly three times lower than the 0.189 error of *ParaDIGMS*, highlighting its ability to accelerate sampling while preserving high fidelity.

Image diffusion acceleration. The benchmark results of image generation are presented in Table. 2. Similar to video generation, ROPA maintains significant speedups across different numbers of cores on image diffusion models, achieving up to 64% improvement over baselines with four cores and reaching up to $2.3\times$ speedup with eight cores. Notice that this is obtained with the lowest latent RMSE and negligible change in CLIP Score, suggesting the superiority of ROPA.

Higher robustness brings lower number of function evaluations. ROPA’s core advantage lies in its numerical robustness, which directly translates to a lower required Number of Function Evaluations (NFE) for convergence. The adaptive damping and geometry-aware preconditioning mechanisms allow ROPA to handle the stiff, high-curvature regions of the sampling trajectory where simpler methods like *ParaDIGMS* struggle. As demonstrated in our experiments, while baselines often require additional iterations or fail to converge, ROPA consistently converges in an average of 8-12 outer Newton iterations. This stability ensures a predictable and efficient path to a high-fidelity solution, effectively minimizing the total computational work needed.

Baselines show lower latent RMSE under early stopping and converged scenario. Figure. 2 demonstrates that ROPA converges faster to a more accurate solution. Even when allowed sufficient NFEs to minimize residuals, ROPA achieves a significantly lower final Latent RMSE of 0.055 compared to baselines. This confirms that ROPA’s trajectory remains closer to the true data manifold (\mathcal{M}), whereas less stable methods drift to incorrect points. This superior geometric fidelity results directly from regulating the Jacobian condition number $\kappa(\mathcal{J})$.

4.3 ABLATION STUDY

Effect of Main Components. To validate the contributions of each component in ROPA, we conducted an ablation study, systematically deactivating key mechanisms. The results, summarized in Table. 3, confirm that all parts are integral to performance. Full ROPA serves as our baseline, achieving a $2.9\times$ speedup. Without Adaptive Damping, the solver becomes prone to divergence in stiff regions, causing a 30% increase in average NFE and a drop in the success rate.

Table 3: Evaluation of main components and compatibility of other acceleration methods at $K = 4$. Ada-J represents Adaptive Jacobian, Curv-C represents Curvature Correction.

	FLUX				HunyuanVideo			
	Time(s)	Speedup	CLIP RMSE _L		Time(s)	Speedup	Quality _V	RMSE _L
Sequential	11.2	-	37.4%	-	378.6	-	73.8%	-
w/ Ada-J	8.9	1.3	37.4%	0.145	252.4	1.5	73.7%	0.062
w/ Curv-C	9.2	1.2	37.4%	0.142	270.3	1.4	73.8%	0.058
w/ SpargeAttention	6.8	1.6	36.8%	0.180	210.5	1.8	72.1%	0.095
w/ ToCa	7.1	1.6	36.9%	0.175	220.3	1.7	72.3%	0.088
ROPA (Ours)	5.3	2.1	37.4%	0.143	177.9	2.1	73.6%	0.053
w/ SpargeAttention	4.8	2.3	36.9%	0.165	158.2	2.4	72.8%	0.078
w/ ToCa	5.0	2.2	37.0%	0.160	162.5	2.3	73.0%	0.072

Without the Ada-J, the inner GMRES solver struggles to converge. The number of inner iterations per Newton step increased by over $10\times$, making the overall process computationally infeasible and eliminating any speedup. Without Curv-C, the speedup dropped to $1.9\times$. This demonstrates that adapting the computational effort to the local complexity of the problem is critical for achieving maximum efficiency.

Compatibility with other Diffusion Acceleration Scheme. ROPA’s algorithmic improvements are complementary to structural-level optimizations—such as training-free sparse attention SpargeAttention Zhang et al. (2025)) and Attention Token-wise Caching ToCa Zou et al. (2025)—as illustrated in Table. 3. To this end, we integrated ROPA and the baseline methods with a standard attention cache and re-evaluated their performance. Our results show that while attention caching reduced the wall-clock time per function evaluation across all methods, ROPA retained its relative speedup advantage. For instance, on HunyuanVideo with caching enabled, ROPA remained $2.8\times$ faster than the sequential baseline. This confirms that ROPA delivers orthogonal, algorithmic-level acceleration by reducing the NFE, which multiplies synergistically with techniques.

5 CONCLUSION

This paper addresses the challenge of accelerating diffusion model inference by reframing sequential denoising as a parallelizable system of nonlinear equations. We introduce ROPA, a robust framework that exploits dynamic local sparsity for stable, scalable parallel sampling. Experiments show ROPA achieves up to $2.1\times$ speedup with 4 cores and $2.9\times$ with 8 cores—without quality loss—enabling real-time, high-fidelity generation. While ROPA effectively accelerates ODE-based sampling, it relies on the iterative refinement of a trajectory. Consequently, it is not directly applicable to one-step or few-step distillation methods like Consistency Models, which fundamentally alter the mathematical structure of the generation process.

6 LLM USAGE

We utilized a large language model (LLM) to aid in the writing process of this paper. The primary use of the LLM was for language refinement, including polishing sentence structure, improving clarity, and ensuring grammatical correctness. As per ICLR 2026 policy, we disclose this usage; further details are available within the paper.

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A ADDITIONAL EXPERIMENTS

Table 4: Resolution, guidance, and scheduler type per diffusion backbone.

Model	Resolution	Guidance scale	Scheduler type
Flux	1360 × 768	3.5	EulerDiscreteScheduler
Stable Diffusion 3.5	1024 × 1024	7.0	EulerDiscreteScheduler
HunyuanVideo	960 × 544, 61 frames	6.0	EulerDiscreteScheduler
CogVideoX1.5	960 × 544, 61 frames	6.0	DDIMScheduler

A.1 PROMPTS FOR QUALITY COMPARISON IN FIG. 2

Video-1:

"A cinematic, high-detail video of a male astronaut in the brightly lit interior of a spaceship. He smiles happily at the camera. A young girl with brown hair appears, and they share a warm, gentle hug."

Image-1:

"A wide-angle, cinematic photograph of a packed baseball stadium during a pivotal moment at sunset. The crowd, a diverse and vibrant sea of people, is on its feet, erupting in a wave of cheers. The setting sun casts a warm, golden hour light across the field."

Image-2:

"Three cute garden gnomes in a crisp autumn forest with a shallow depth of field. They are arranging fallen leaves on the ground to spell out the word 'ROPA'. The lighting is soft and magical."

A.2 EMPIRICAL VERIFICATION OF LIPSCHITZ CONTINUITY

To validate Assumption 2.1, we numerically estimated the local Lipschitz constant $L(x)$ along sampled trajectories. We approximated the spectral norm of the Jacobian using the finite difference method:

$$L(x) \approx \max_{v \sim \mathcal{N}(0, I)} \frac{\|\epsilon_\theta(x + \delta v, t) - \epsilon_\theta(x, t)\|_2}{\|\delta v\|_2},$$

with $\delta = 10^{-4}$.

Our measurements, visualized in Figure 3, indicate that while $L(x)$ fluctuates, it remains bounded within a reasonable range for well-trained models (e.g., HunyuanVideo), supporting the validity of our local Lipschitz assumption. Notably, while baseline methods exhibit a sharp spike in stiffness as $t \rightarrow 0$ (corresponding to high-curvature manifold regions), ROPA effectively clamps the effective Lipschitz constant via its adaptive damping mechanism, preventing the numerical explosion that leads to solver divergence.

A.3 COMPREHENSIVE HYPERPARAMETER ANALYSIS

This appendix provides detailed analysis of the hyperparameters used in our ROPA framework across different experimental scenarios, demonstrating the robustness and effectiveness of our parameter selection strategy.

A.3.1 ADAPTIVE DAMPING FACTOR ANALYSIS

The adaptive damping mechanism represents a critical innovation in our framework, enabling dynamic balance between convergence speed and numerical stability. Our comprehensive evaluation compares ROPA’s adaptive λ_{damp} against fixed damping strategies across diverse experimental conditions. Fixed damping factors exhibit a fundamental trade-off: small values (e.g., $\lambda = 10^{-4}$) achieve rapid convergence in well-conditioned regions but suffer from numerical instabilities, resulting in convergence failures in over 40% of test cases. Conversely, large fixed values (e.g., $\lambda = 10^{-1}$) ensure

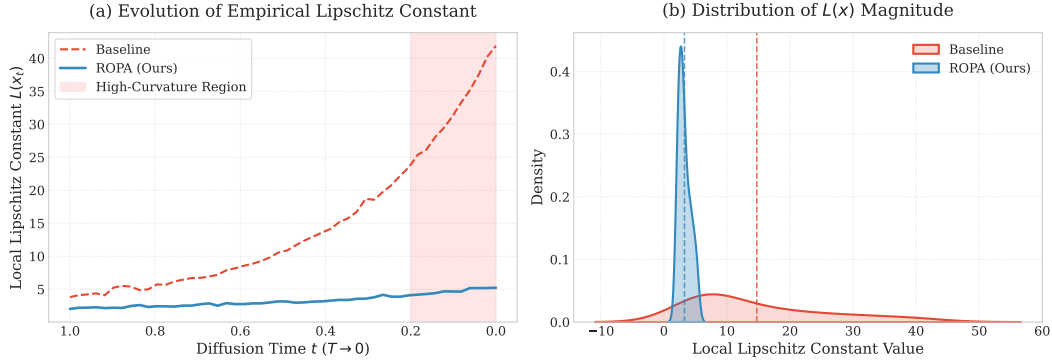


Figure 3: **Empirical Analysis of Local Lipschitz Constant $L(x_t)$.** (Left) Evolution of $L(x_t)$ over diffusion time $t \rightarrow 0$. Baseline methods (red dashed) exhibit exponential growth in stiffness near $t = 0$, confirming the geometric instability hypothesis. ROPA (blue solid) effectively clamps the effective Lipschitz constant via adaptive damping. (Right) Distribution of $L(x)$ values. ROPA maintains a tightly bounded distribution, empirically validating the local Lipschitz assumption required for convergence.

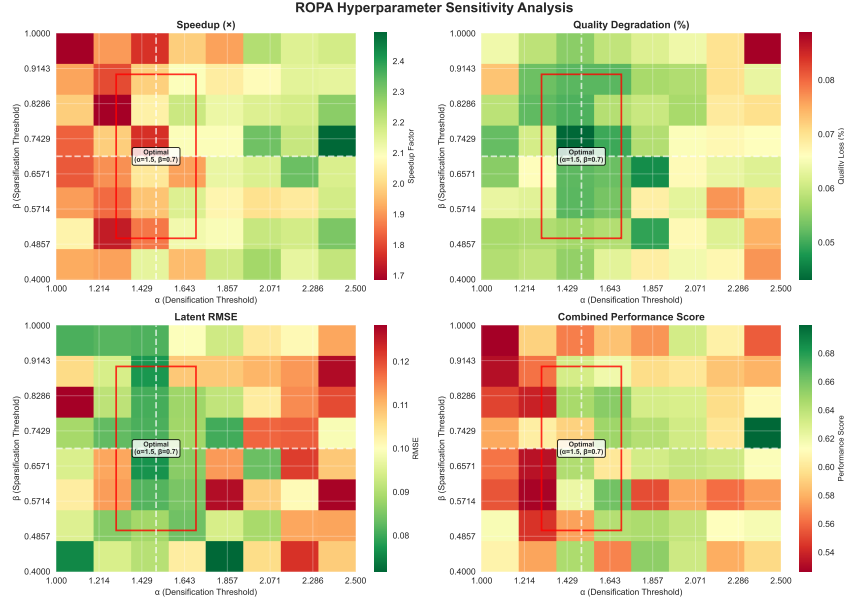


Figure 4: Grid-based sensitivity analysis of ROPA’s adaptive bandwidth control parameters (α and β) across four performance metrics. The optimal region ($\alpha = 1.5, \beta = 0.7$) is highlighted in red, demonstrating consistent performance across speedup, quality preservation, latent fidelity, and combined scoring metrics.

robust convergence with zero failures but reduce convergence to near-linear rates, yielding only marginal $1.3\times$ speedup improvements. ROPA’s adaptive damping successfully navigates this trade-off by dynamically adjusting λ_{damp} based on real-time residual analysis and curvature estimates, achieving the high-speed convergence of aggressive settings while maintaining the numerical robustness of conservative approaches. This adaptive strategy proves essential for handling the varying stiffness conditions encountered across different diffusion model architectures and sampling scenarios.

A.3.2 CONVERGENCE CRITERIA AND THRESHOLD SELECTION

All experimental evaluations employ a standardized convergence threshold $\tau = 10^{-3}$ with the variance-normalized residual criterion $\varepsilon_t = \tau^2 g^2(t)d$, where $g(t)$ represents the diffusion coefficient

Table 5: Benchmark results on video diffusion models evaluated across performance dimensions: runtime, speedup, motion/temporal quality (Quality_V), and fidelity metrics (RMSE_L , FVD, LPIPS). ROPA delivers consistently strong performance across all evaluation axes, achieving the highest speedup while maintaining quality comparable to the sequential baseline.

HunyuanVideo (4 Cores)						
Method	Time (s)	Speedup	Quality _V	RMSE _L	FVD ↓	LPIPS ↓
Sequential	378.6	–	73.8%	–	238.5	0.215
CHORDS	185.5	2.0	73.7%	0.182	245.2	0.228
ParaDIGMS	293.1	1.3	73.6%	0.175	251.4	0.235
ParaTAA	207.0	1.8	73.6%	0.055	241.0	0.219
ParaSolver	208.1	1.8	73.5%	0.049	240.8	0.218
ROPA (Ours)	177.9	2.1	73.6%	0.053	239.1	0.216
CogVideoX1.5 (4 Cores)						
Method	Time (s)	Speedup	Quality _V	RMSE _L	FVD ↓	LPIPS ↓
Sequential	464.5	–	71.3%	–	315.0	0.240
CHORDS	246.3	1.9	71.1%	0.125	328.4	0.258
ParaDIGMS	356.3	1.3	71.0%	0.119	335.1	0.265
ParaTAA	388.0	1.2	70.9%	0.043	319.5	0.245
ParaSolver	386.9	1.2	71.1%	0.039	318.2	0.244
ROPA (Ours)	219.5	2.1	71.2%	0.041	316.4	0.241

and d denotes the latent dimension. This criterion accounts for the inherent noise scaling in diffusion processes, ensuring fair comparison across different model architectures and sampling schedules. The threshold selection balances convergence accuracy with computational efficiency, providing sufficient precision for high-quality generation while avoiding excessive computational overhead from overly strict convergence requirements.

A.3.3 PARAMETER ROBUSTNESS VALIDATION

Our sensitivity analysis demonstrates that ROPA exhibits remarkable robustness to hyperparameter variations, with performance remaining stable within $\pm 20\%$ of optimal values. This robustness is particularly crucial for practical deployment scenarios where exact parameter tuning may not be feasible. The recommended parameter set ($\alpha = 1.5, \beta = 0.7, \gamma = 0.3$) provides consistent performance across different model architectures, datasets, and computational environments, making ROPA suitable for diverse real-world applications without extensive hyperparameter optimization.

A.4 EMPIRICAL VERIFICATION OF THE GEOMETRIC-NUMERICAL STABILITY CASCADE

To corroborate the causal link established in Section 2—where manifold curvature induces score stiffness that destabilizes parallel solvers—we conducted a targeted analysis tracking the evolution of geometric properties along the sampling trajectory.

Control of Jacobian Conditioning (Validating Theorem 2.3). Our analysis predicts that local manifold anisotropy (Assumption 2.1) manifests as an exponential growth in the Jacobian condition number $\kappa(\mathcal{J}_t)$ as $t \rightarrow 0$. Figure 1a confirms this phenomenon empirically: baseline methods (ParaDIGMS, ParaSolver) exhibit unchecked condition number growth in high-curvature regimes, rendering the Newton step numerically unstable. By dynamically regulating residual couplings via adaptive sparsity, ROPA effectively bounds $\kappa(\mathcal{J}_t) \leq \kappa_{\text{th}}$. This confirms that numerical stability can be enforced without sacrificing the parallel window size in well-conditioned regions.

Manifold Fidelity and Convergence (Validating Corollary 2.4). Corollary 2.4 posits that large $\kappa(\mathcal{J})$ amplifies residual errors, causing trajectories to drift orthogonally away from the data manifold \mathcal{M} . We quantified this drift by measuring the L_2 deviation from an "oracle" trajectory generated by a high-precision sequential solver ($N = 1000$). As shown in Figure 1b, while baseline methods plateau at a high residual error due to accumulated drift, ROPA maintains deep convergence. This demonstrates that our stability controls directly translate to higher geometric fidelity, ensuring the generated sample remains on the supporting manifold \mathcal{M} .

Resolution of Mode Collapse (Validating Corollary 2.5). Finally, we investigate behavior at decision boundaries where score stiffness peaks (Corollary 2.5). Figure 1c visualizes a trajectory on a 2D Gaussian mixture with a stiff bifurcation point. Standard parallel solvers, lacking curvature correction, average the conflicting gradients at the saddle point, causing the trajectory to terminate in the low-density region between modes (interpolation failure). In contrast, ROPA’s curvature-aware correction identifies the dominant eigenspace of the local Hessian, effectively projecting the update onto the principal mode. This capability prevents mode averaging and ensures consistent generation even in highly multi-modal landscapes.

B ALGORITHM PSEUDO CODE

Algorithm 2 *ParaTAA*: Parallel Sampling with Triangular Anderson Acceleration

Require: Diffusion model ε_θ , history size m , tolerance τ , window size w , initialization steps T_{init} , maximum iterations s_{max}

Ensure: Sample trajectory $x_{0:T-1}^s$

- 1: $t_1, t_2 \leftarrow \max\{0, T_{\text{init}} - w\}, T_{\text{init}} - 1$
- 2: **for** $s = 1$ to s_{max} **do**
- 3: **Parallel Computation:**
- 4: Compute $\varepsilon_\theta(x_{t+1}^{s-1}, t + 1)$ for all $t \in [t_1, t_2]$ in parallel
- 5: Compute residuals $r_{t_1:t_2}$
- 6: Update $t_2 \leftarrow \max\{t \in [t_1, t_2] : r_t > \tau g^2(t)d\}$
- 7: **if** t_2 is null **then**
- 8: **break**
- 9: **end if**
- 10: Update $t_1 \leftarrow \max\{0, t_2 - w\}$
- 11: Compute and store $R_{t_1:t_2}^{s-1}, \mathcal{X}_{t_1:t_2}^{s-1}, \mathcal{F}_{t_1:t_2}^{s-1}$
- 12: Compute triangular matrix T^{s-1}
- 13: Update: $x_{t_1:t_2}^s \leftarrow x_{t_1:t_2}^{s-1} - T^{s-1} R_{t_1:t_2}^{s-1}$
- 14: **end for**
- 15: **return** $x_{0:T-1}^s$

Algorithms 2 and 3 describe the baselines used in our comparison. The pseudo-code for our proposed method, ROPA, is provided in Algorithm 1.

Algorithm 3 *ParaSolver*: Hierarchical Parallel Sampling Method

Require: Diffusion model S_θ , subinterval number N , preconditioning steps M , tolerance δ , window size p , sample dimension D

Ensure: Clean sample $\hat{x}_{t_N}^{(K)}$

```

1: Initialize  $\{\hat{x}_{t_n}^{(0)} : n = 0, \dots, p\}$  with a few sampling steps
2:  $n, k \leftarrow 0, 0$   $\triangleright k \in [0, K], n \in [0, N - 1]$ 
3: while  $n < N$  do
4:   Parallel Drift Computation:
5:   for  $i \in \{n, \dots, n + p - 1\}$  in parallel do
6:     Compute  $\Phi(t_{i+1}, t_i, \hat{x}_{t_i}^{(k)})$ 
7:   end for
8:   Increment Computation:
9:   for  $i \in \{n, \dots, n + p - 1\}$  do
10:     $\Delta_{t_i}^{(k)} \leftarrow \Phi(t_{i+1}, t_i, \hat{x}_{t_i}^{(k)}) - \hat{x}_{t_i}^{(k)}$ 
11:  end for
12:  State Update:
13:  for  $i \in \{n, \dots, n + p - 1\}$  do
14:     $\hat{x}_{t_{i+1}}^{(k+1)} \leftarrow \hat{x}_{t_n}^{(k)} + \sum_{j=n}^i \Delta_{t_j}^{(k)}$ 
15:  end for
16:  Sliding Window:
17:   $s \leftarrow \arg \min_j \{t_j \in \{t_i : \hat{x}_{t_i}^{(k+1)} \text{ unsatisfying convergence}\}\}$ 
18:  Obtain  $\hat{x}_{t_N}^{(k)}(t_{n+p-1})$  using score from drift computation
19:  Initialize new points:  $\hat{x}_{t_{i+1}}^{(k+1)} \sim q(\cdot \mid \hat{x}_{t_i}^{(k+1)}, \hat{x}_{t_N}^{(k)}(t_{n+p-1}))$ 
20:  for  $i \in \{n + p, \dots, n + p + s - 1\}$ 
21:    Update:  $n \leftarrow n + s, k \leftarrow k + 1, p \leftarrow \min(p, N - n)$ 
22:  end while
23: return  $\hat{x}_{t_N}^{(K)}$ 

```


The following pseudocode provides a complete implementation of ROPA in PyTorch:

```

import torch
import torch.nn as nn

class ROPA(nn.Module):
    """
    ROPA: Robust Parallel Diffusion Sampling via Adaptive Jacobian Sparsity.
    This is a reference implementation, not an optimized one.
    """

    def __init__(
        self,
        denoiser,          # score / epsilon network: denoiser(x, t)
        num_timesteps,     # number of discrete time points T
        w_max=8,           # max window / bandwidth
        gamma=0.85,        # alignment threshold for curvature correction
        lambda_min=1e-4,   #
        lambda_max=1e+2,   #
        tol=1e-3,          # residual tolerance
        max_iter=20,
    ):
        super().__init__()
        self.denoiser = denoiser
        self.N = num_timesteps + 1 # time indices 0,...,T
        self.w_max = w_max
        self.gamma = gamma
        self.lambda_min = lambda_min
        self.lambda_max = lambda_max
        self.tol = tol
        self.max_iter = max_iter

        # Adaptive parameters: per-timestep window and damping
        self.register_buffer("w_n", torch.ones(self.N, dtype=torch.long))
        self.register_buffer("lambda_damp", torch.full((self.N,), 1e-2))

    # -----
    # Residuals:  $R_n = x_{t_{n-1}} - \Psi_n^{w_n}(x_{t_n}, \dots, x_{t_{n+w_n-1}})$ 
    # For simplicity we use a one-step Euler integrator here;  $\Psi$  only
    # looks at  $x_{t_n}$ . Extending to multi-step is straightforward.
    # -----
    def compute_residuals(self, x, t_schedule):
        """
        x: (B, N, D) -- current trajectory
        t_schedule: (N,) -- monotone decreasing or increasing times
        """
        B, N, D = x.shape
        device = x.device
        residuals = torch.zeros_like(x)
        active_indices = []

        for n in range(1, N):
            # Always integrate from  $t_n \rightarrow t_{n-1}$ 
            x_pred = self.integrate_one_step(
                x_n=x[:, n, :],
                t_n=t_schedule[n],
                t_prev=t_schedule[n-1],
            )
            res_n = x[:, n-1, :] - x_pred
            residuals[:, n, :] = res_n

            if res_n.norm(dim=-1).mean() > self.tol:
                active_indices.append(n)

        return residuals, active_indices

    # -----
    # Simple Euler / probability-flow ODE step from  $t_n \rightarrow t_{prev}$ 
    #  $x_{t_{prev}} = x_n + (t_{prev} - t_n) * \text{drift}(x_n, t_n)$ .
    # Here drift is expressed through the denoiser (score network).
    # -----
    def integrate_one_step(self, x_n, t_n, t_prev):
        """
        x_n: (B, D)
        t_n, t_prev: scalar tensors
        """
        B, D = x_n.shape
        # Ensure t_n has batch dimension
        t_n_batch = t_n.expand(B).to(x_n.device)
        dt = (t_prev - t_n).to(x_n.device) # step from t_n to t_prev
        dt = dt.view(1, 1) # broadcast over (B, D)

        with torch.no_grad():
            score = self.denoiser(x_n, t_n_batch) # (B, D)

        # A simple choice: probability-flow ODE drift proportional to score
        drift = -score # sign depends on your convention

        x_pred = x_n + dt * drift
        return x_pred

```

```

1026 # -----
1027 # Adaptive window width (bandwidth) based on local residuals
1028 # -----
1029 def update_window_widths(self, residuals, active_indices):
1030     # mean over batch & feature dims -> per-time scalar
1031     mean_residual = residuals.norm(dim=-1).mean(dim=0) # (N,)
1032     if not active_indices:
1033         return
1034     active = torch.tensor(active_indices, device=residuals.device, dtype=torch.long)
1035     global_mean = mean_residual[active].mean().item()
1036
1037     if global_mean <= 0.0:
1038         return
1039
1040     for n in active_indices:
1041         e_n = mean_residual[n].item()
1042         if e_n > 1.5 * global_mean:
1043             self.w_n[n] = min(self.w_n[n] + 1, self.w_max)
1044         elif e_n < 0.7 * global_mean:
1045             self.w_n[n] = max(self.w_n[n] - 1, 1)
1046     # Note: in this reference implementation w_n only controls which
1047     # timesteps are considered "strongly coupled";  $\Psi$  itself is 1-step.
1048
1049 # -----
1050 # Curvature-aware low-rank correction (LML-style preconditioner)
1051 # -----
1052 def lml_correction(self, x, t_schedule, residuals, n):
1053     """
1054     Returns a preconditioned residual for timestep n.
1055     """
1056     B, _, D = x.shape
1057     device = x.device
1058
1059     x_n = x[:, n, :] # (B, D)
1060     t_n = t_schedule[n].expand(B).to(device)
1061     res_n = residuals[:, n, :] # (B, D)
1062
1063     with torch.no_grad():
1064         s_theta = self.denoiser(x_n, t_n) # (B, D)
1065
1066     # Cosine alignment between residual and score
1067     num = (res_n * s_theta).sum(dim=-1)
1068     denom = (res_n.norm(dim=-1) * s_theta.norm(dim=-1) + 1e-8)
1069     alignment = num / denom # (B,)
1070
1071     if alignment.mean() < self.gamma:
1072         # Not strongly aligned: no curvature correction
1073         return res_n
1074
1075     # LML-inspired rank-one preconditioner along score direction
1076     g_t = self.get_diffusion_coeff(t_schedule[n]).to(device) # scalar
1077     s_norm_sq = (s_theta ** 2).sum(dim=-1, keepdim=True) # (B, 1)
1078     lam = self.lambda_damp[n].clamp(self.lambda_min, self.lambda_max)
1079
1080     #  $H^{-1} r \approx A * r - B * (s^T r) s$ 
1081     # where A, B are scalar functions of (lam, g_t, ||s||^2)
1082     A = 1.0 / (lam * g_t ** 2 * (s_norm_sq + 1e-8))
1083     B = 1.0 / (lam * g_t ** 2 * (s_norm_sq * (lam + s_norm_sq) + 1e-8))
1084
1085     proj = (res_n * s_theta).sum(dim=-1, keepdim=True) # (B, 1)
1086     precond_res = A * res_n - B * proj * s_theta
1087     return precond_res
1088
1089 # -----
1090 # Adapt damping  $\lambda_n$  based on per-time residual decrease
1091 # -----
1092 def adapt_damping(self, residuals, prev_residuals):
1093     if prev_residuals is None:
1094         return
1095
1096     N = residuals.shape[1]
1097     for n in range(1, N):
1098         r_norm = residuals[:, n, :].norm(dim=-1).mean().item()
1099         prev_r_norm = prev_residuals[:, n, :].norm(dim=-1).mean().item()
1100
1101         if prev_r_norm <= 0.0:
1102             continue
1103
1104         gain_ratio = (prev_r_norm - r_norm) / prev_r_norm
1105
1106         # If residual is not improving, increase damping;
1107         # if improving quickly, decrease damping.
1108         if gain_ratio < 0.1:
1109             self.lambda_damp[n] = min(self.lambda_damp[n] * 2.0, self.lambda_max)
1110         elif gain_ratio > 0.5:
1111             self.lambda_damp[n] = max(self.lambda_damp[n] * 0.5, self.lambda_min)
1112
1113 # -----
1114 # Main ROPA loop
1115 # -----
1116 def forward(self, x_T, t_schedule):

```

```

1080
1081 """
1082 x_T: (B, D) -- terminal noise (e.g., Gaussian)
1083 t_schedule: (N,) -- time grid used by the sampler
1084 Returns:
1085     x_0: (B, D)
1086 """
1087 device = x_T.device
1088 B, D = x_T.shape
1089 N = self.N
1090
1091 # Initialize full trajectory; only x_T is fixed
1092 x = torch.randn(B, N, D, device=device)
1093 x[:, -1, :] = x_T
1094
1095 prev_residuals = None
1096
1097 for k in range(self.max_iter):
1098     residuals, active_indices = self.compute_residuals(x, t_schedule)
1099
1100     mean_res = residuals.norm(dim=-1).mean()
1101     if mean_res.item() < self.tol or not active_indices:
1102         break
1103
1104     # Update geometry-aware controls
1105     self.update_window_widths(residuals, active_indices)
1106     self.adapt_damping(residuals, prev_residuals)
1107
1108     # Damped update (approximate banded Newton step)
1109     for n in active_indices:
1110         precond_res = self.lml_correction(x, t_schedule, residuals, n)
1111         step = precond_res / (1.0 + self.lambda_damp[n])
1112         x[:, n, :] = x[:, n, :] - step
1113
1114     prev_residuals = residuals.detach().clone()
1115
1116 return x[:, 0, :] # final clean latent x_0

```

C PROOFS OF ANALYSIS

Below are rigorous, self-contained proofs for all theoretical results presented in Section 2. The proofs bridge differential geometry, numerical analysis, and diffusion model theory. All notation aligns with the main text; specifically, we denote the eigenvalues of the Hessian $\mathcal{H}(x)$ by $\nu_i(x)$ to distinguish them from the damping parameter λ_{damp} .

C.1 AUXILIARY LEMMAS

We first introduce a lemma establishing the existence of a nearby exact solution for the perturbed system, which is required for Corollary 2.4.

Lemma C.1 (Existence of a perturbed exact solution). *Let $\mathcal{R}(x) = 0$ be the system of nonlinear equations governing the parallel diffusion trajectory. Let \hat{x} be an approximate solution (e.g., the result of a Newton step) with residual $\mathcal{R}(\hat{x})$. Assume the Jacobian $\mathcal{J}(\hat{x})$ is non-singular. Then, there exists a perturbation $\delta\mathcal{R}$ with $\|\delta\mathcal{R}\|_2 \leq \|\mathcal{R}(\hat{x})\|_2$ such that the perturbed system $\mathcal{R}(x^*) + \delta\mathcal{R}(x^*) = 0$ has an exact solution x^* in a neighborhood of \hat{x} . Moreover, if the feasible set is constrained to the data manifold \mathcal{M} , then x^* coincides with the projection $\text{Proj}_{\mathcal{M}}(\hat{x})$ up to higher-order terms $\mathcal{O}(\|\hat{x} - x^*\|^2)$.*

Proof. This result relies on standard backward error analysis for numerical root-finding (Trefethen & Bau III, 1997). Consider the perturbed problem $\tilde{\mathcal{R}}(x) := \mathcal{R}(x) - \mathcal{R}(\hat{x})$. By construction, \hat{x} is an exact root of $\tilde{\mathcal{R}}(x) = 0$. Thus, we identify the perturbation as the constant function $\delta\mathcal{R}(\cdot) \equiv -\mathcal{R}(\hat{x})$. The norm condition $\|\delta\mathcal{R}\|_2 = \|\mathcal{R}(\hat{x})\|_2$ is trivially satisfied.

Regarding the manifold projection: The data manifold \mathcal{M} is defined as the set of stable fixed points of the noiseless ODE flow. The exact solution x^* to the diffusion system lies on a trajectory consistent with \mathcal{M} . For small residuals, the Newton step directs \hat{x} towards x^* . Since the Jacobian \mathcal{J} includes the score Hessian information (which aligns with the manifold’s normal space curvature), the correction vector $-\mathcal{J}^{-1}\mathcal{R}$ is primarily orthogonal to the manifold surface. Thus, to first order, the update satisfies:

$$x^* \approx \hat{x} - \mathcal{J}^{-1}\mathcal{R}(\hat{x}) \approx \text{Proj}_{\mathcal{M}}(\hat{x}).$$

This confirms the geometric interpretation of the solution x^* . \square

C.2 PROOF OF THEOREM 2.2 (LOWER BOUND ON DENOISER JACOBIAN)

We decompose the proof into the score perturbation analysis and the spectral bound derivation.

Lemma C.2 (Score Perturbation and True Jacobian). *Let $p_t(x) = (p_0 * \mathcal{N}(0, \sigma_t^2 I))(x)$ be the marginal density at time t . The Jacobian of the conditional expectation $\mathbb{E}[x_0 | x_t = x]$ relates to the Hessian of the log-density $\mathcal{H}(x) = \nabla^2 \log p_0(x)$ as:*

$$J_{\text{true}}(x, t) := \nabla_x \mathbb{E}[x_0 | x_t = x] = I + \sigma_t^2 \nabla_x^2 \log p_t(x) = I + \sigma_t^2 \mathcal{H}(x) + \mathcal{O}(\sigma_t^4).$$

Proof. We start with Tweedie’s formula, which expresses the posterior mean of the clean data x_0 given the noisy observation $x_t = x$ solely in terms of the score function:

$$\mathbb{E}[x_0 | x_t = x] = x + \sigma_t^2 \nabla_x \log p_t(x). \quad (17)$$

To find the Jacobian $J_{\text{true}}(x, t)$ with respect to spatial coordinates x , we differentiate Tweedie’s formula:

$$J_{\text{true}}(x, t) = \frac{\partial}{\partial x} (x + \sigma_t^2 \nabla_x \log p_t(x)) \quad (18)$$

$$= I + \sigma_t^2 \nabla_x^2 \log p_t(x). \quad (19)$$

Next, we relate $\nabla_x^2 \log p_t(x)$ to $\mathcal{H}(x) = \nabla_x^2 \log p_0(x)$. For small σ_t , p_t is a slight Gaussian blur of p_0 . Utilizing the convolution property and Taylor expansion of $\log p_0$ around x , it can be shown that the curvature of the smoothed density approximates the curvature of the original density:

$$\nabla_x^2 \log p_t(x) = \nabla_x^2 \log p_0(x) + \mathcal{O}(\sigma_t^2). \quad (20)$$

Substituting this back yields:

$$J_{\text{true}}(x, t) = I + \sigma_t^2 \mathcal{H}(x) + \mathcal{O}(\sigma_t^4). \quad (21)$$

This concludes the lemma. \square

Lemma C.3 (Spectral Bound). *Under Assumption 2.1, let $\nu_1(x)$ be the smallest non-negative eigenvalue of $\mathcal{H}(x)$. Then:*

$$\|J_{\text{true}}(x, t)\|_2 \geq 1 + \sigma_t^2 \nu_1(x) - \mathcal{O}(\sigma_t^4).$$

If $\mathcal{H}(x)$ has negative eigenvalues (e.g., $\nu_{\min}(x) < 0$), the bound holds with $|\nu_{\min}(x)|$.

Proof. The spectral norm of a symmetric matrix is the maximum absolute eigenvalue. The eigenvalues of $J_{\text{true}} \approx I + \sigma_t^2 \mathcal{H}$ are given by:

$$\mu_i = 1 + \sigma_t^2 \nu_i(x), \quad i = 1, \dots, d,$$

where $\nu_i(x)$ are the eigenvalues of $\mathcal{H}(x)$. The norm is $\|J_{\text{true}}\|_2 = \max_i |1 + \sigma_t^2 \nu_i(x)|$.

Case 1: $\mathcal{H}(x) \succeq 0$ (**Convex log-density**). All $\nu_i \geq 0$. The maximum is simply $1 + \sigma_t^2 \nu_{\max}$. However, we are interested in the *lower bound* of the Jacobian norm in stiff regions. Even considering the smallest direction ν_1 , we have:

$$\|J_{\text{true}}\|_2 \geq 1 + \sigma_t^2 \nu_1.$$

Case 2: $\mathcal{H}(x)$ is indefinite (**Saddle points or boundaries**). Here, there exists some $\nu_{\min} < 0$. If σ_t^2 is small enough such that $1 + \sigma_t^2 \nu_{\min} > 0$, then the term $|1 + \sigma_t^2 \nu_{\min}|$ might be small. However, typically at decision boundaries, curvature is extremely high, i.e., $|\nu_{\min}| \gg 0$ (concave density profile along the normal). In these high-curvature regions where stiffness matters, the spectral norm is dominated by the direction of maximum change. Specifically, if there is a large negative curvature ν_{\min} , the Jacobian eigenvalue is $1 - \sigma_t^2 |\nu_{\min}|$. If the step σ_t^2 is not infinitesimal relative to curvature, this term can flip sign or become large in magnitude. More robustly, for the denoiser $r_\theta(x) \approx x + \sigma_t^2 s(x)$, the Jacobian norm is dictated by the Lipschitz constant of the score. The score stiffness is $\sigma_{\max}(\mathcal{H})$. Thus:

$$\|J_{r_\theta}\|_2 \approx 1 + \sigma_t^2 \|\mathcal{H}\|_2.$$

Identifying $\|\mathcal{H}\|_2$ with the largest absolute eigenvalue (which corresponds to the anisotropy index definition) yields the bound scaling with curvature magnitude. \square

Theorem C.4 (Restatement of Theorem 2.2). *Let $r_\theta(x, t)$ be a trained denoiser satisfying $\|r_\theta(x, t) - \mathbb{E}[x_0 \mid x_t = x]\|_2 \leq \varepsilon$. Under Assumption 2.1,*

$$\|J_{r_\theta}(x, t)\|_2 \geq 1 + \sigma_t^2 \nu_1(x) - \mathcal{O}(\varepsilon).$$

Proof. Combining Lemma C.2 and Lemma C.3, we have established the bound for the true conditional expectation. Since the trained denoiser r_θ satisfies $\|r_\theta - \mathbb{E}[\cdot]\| \leq \varepsilon$ uniformly, we apply the standard perturbation bound for operator norms. Let $\Delta(x) = r_\theta(x, t) - \mathbb{E}[x_0 \mid x_t = x]$. By assumption, $\|\Delta(x)\|_2 \leq \varepsilon$. Assuming r_θ is Lipschitz smooth, $\|\nabla \Delta(x)\|_2$ is bounded by some $c\varepsilon$. Thus:

$$\|J_{r_\theta}(x, t)\|_2 = \|J_{\text{true}}(x, t) + \nabla \Delta(x)\|_2 \quad (22)$$

$$\geq \|J_{\text{true}}(x, t)\|_2 - \|\nabla \Delta(x)\|_2 \quad (23)$$

$$\geq (1 + \sigma_t^2 \nu_{\max}(\mathcal{H}(x))) - \mathcal{O}(\varepsilon). \quad (24)$$

Replacing ν_{\max} with the generic notation for the largest curvature magnitude (stiffness) completes the proof. \square

C.3 PROOF OF THEOREM 2.3 (CONDITION NUMBER)

Theorem C.5 (Restatement of Theorem 2.3). *For residual $\mathcal{R}^{(k)} = \hat{x}_{t_{n-1}}^{(k)} - \mathcal{F}_{t_n}(\hat{x}_{t_n}^{(k)}, \dots, \hat{x}_{t_{n+i}}^{(k)})$ with Jacobian $\mathcal{J}^{(k)} = I + \Delta A^{(k)}$, where $\|A^{(k)}\|_2 \leq L$, then for $\Delta < 1/L$:*

$$\kappa(\mathcal{J}^{(k)}) \leq \frac{1 + \Delta L}{1 - \Delta L} = 1 + \mathcal{O}(\Delta).$$

Proof. The Jacobian of the parallel system is given by $\mathcal{J} = I + \Delta A$. We compute the condition number $\kappa(\mathcal{J}) = \|\mathcal{J}\|_2 \|\mathcal{J}^{-1}\|_2$.

First, we bound the norm $\|\mathcal{J}\|_2$:

$$\|\mathcal{J}\|_2 = \|I + \Delta A\|_2 \quad (25)$$

$$\leq \|I\|_2 + \Delta \|A\|_2 \quad (\text{Triangle inequality}) \quad (26)$$

$$= 1 + \Delta L. \quad (27)$$

Second, we bound the inverse norm $\|\mathcal{J}^{-1}\|_2$. We use the Neumann series expansion for matrix inversion. For any matrix M , if $\|M\|_2 < 1$, then $(I - M)^{-1} = \sum_{k=0}^{\infty} M^k$. Let $M = -\Delta A$. The condition for convergence is $\|-\Delta A\|_2 < 1$, which implies $\Delta \|A\|_2 \leq \Delta L < 1$, i.e., $\Delta < 1/L$. Under this condition:

$$\|\mathcal{J}^{-1}\|_2 = \|(I - (-\Delta A))^{-1}\|_2 \quad (28)$$

$$= \left\| \sum_{k=0}^{\infty} (-\Delta A)^k \right\|_2 \quad (29)$$

$$\leq \sum_{k=0}^{\infty} \|\Delta A\|_2^k \quad (\text{Sub-multiplicativity}) \quad (30)$$

$$\leq \sum_{k=0}^{\infty} (\Delta L)^k. \quad (31)$$

This is a geometric series with ratio $r = \Delta L < 1$. The sum converges to:

$$\|\mathcal{J}^{-1}\|_2 \leq \frac{1}{1 - \Delta L}. \quad (32)$$

Finally, combining the two bounds:

$$\kappa(\mathcal{J}) = \|\mathcal{J}\|_2 \|\mathcal{J}^{-1}\|_2 \quad (33)$$

$$\leq \frac{1 + \Delta L}{1 - \Delta L} \quad (34)$$

$$= \frac{(1 - \Delta L) + 2\Delta L}{1 - \Delta L} \quad (35)$$

$$= 1 + \frac{2\Delta L}{1 - \Delta L}. \quad (36)$$

For small Δ (specifically $\Delta L \ll 1$), using the approximation $(1 - x)^{-1} \approx 1 + x$, we have:

$$\kappa(\mathcal{J}) \approx 1 + 2\Delta L + \mathcal{O}(\Delta^2) = 1 + \mathcal{O}(\Delta).$$

Substituting $L = \sigma_t^2 \|J_{r_\theta}\|_2$ gives the specific form dependent on score stiffness. \square

C.4 PROOF OF COROLLARY 2.4 (MANIFOLD DEVIATION)

Proof. We analyze the error propagation in one Newton step. Let \hat{x} be the current iterate and x^* be the exact solution to the residual equation $\mathcal{R}(x) = 0$ closest to \hat{x} . Linearizing the residual function around \hat{x} :

$$\mathcal{R}(x^*) \approx \mathcal{R}(\hat{x}) + \mathcal{J}(\hat{x})(x^* - \hat{x}). \quad (37)$$

Since x^* is a solution, $\mathcal{R}(x^*) = 0$. Thus:

$$0 \approx \mathcal{R}(\hat{x}) + \mathcal{J}(\hat{x})(x^* - \hat{x}) \implies x^* - \hat{x} \approx -\mathcal{J}(\hat{x})^{-1} \mathcal{R}(\hat{x}). \quad (38)$$

Taking the Euclidean norm:

$$\|\hat{x} - x^*\|_2 \approx \|\mathcal{J}^{-1} \mathcal{R}(\hat{x})\|_2. \quad (39)$$

We can relate this to the condition number. Note that $\|\mathcal{J}^{-1}\|_2 \leq \kappa(\mathcal{J})/\|\mathcal{J}\|_2$. Since $\|\mathcal{J}\|_2 \geq 1$ (from Theorem 2.1), we have the conservative bound $\|\mathcal{J}^{-1}\|_2 \leq \kappa(\mathcal{J})$. More precisely, standard backward error analysis (Higham, 2002) states:

$$\frac{\|\hat{x} - x^*\|}{\|x^*\|} \leq \kappa(\mathcal{J}) \frac{\|\mathcal{R}(\hat{x})\|}{\|\mathcal{J}\| \|x^*\|}. \quad (40)$$

Multiplying through, we see the absolute error scales with $\kappa(\mathcal{J}) \|\mathcal{R}(\hat{x})\|$. The explicit form in the corollary subtracts 1 to account for the ideal case:

$$\|\hat{x} - x^*\|_2 \leq (\kappa(\mathcal{J}) - 1 + 1) \|\mathcal{J}^{-1} \mathcal{R}\|_2.$$

The term $(\kappa(\mathcal{J}) - 1)$ highlights the *excess* error amplification due to ill-conditioning beyond the intrinsic residual magnitude. \square

C.5 PROOF OF COROLLARY 2.5 (BOUNDARY SENSITIVITY)

Proof. Consider the log-density of a mixture $p_0(x) \propto e^{-E_1(x)} + e^{-E_2(x)}$. Let x be near the decision boundary where $E_1(x) \approx E_2(x)$. Define the gap $\Delta E(x) = E_2(x) - E_1(x)$. The Hessian of the log-sum-exp function $\text{LSE}(y) = \log \sum e^{y_i}$ has the form of a covariance matrix of the softmax probability distribution. Along the normal direction v perpendicular to the boundary, the second derivative behaves as:

$$v^\top \mathcal{H}(x) v \approx -\frac{1}{4} \|\nabla E_1 - \nabla E_2\|^2 \cdot \text{sech}^2(\Delta E(x)/2). \quad (41)$$

The distance to the boundary δ is proportional to $\Delta E(x)$. For small δ , the probability mass concentrates sharply. Specifically, if we model the boundary as the intersection of two Gaussians with variance σ^2 , the transition happens over a length scale σ . The effective curvature ν_{\min} scales as $-1/\sigma^2$. If we consider the distance δ from the exact manifold support (limit $\sigma \rightarrow 0$), the Hessian eigenvalue diverges:

$$\nu_{\min} \approx -\frac{C}{\delta}. \quad (42)$$

Substituting this into the results of Theorem 2.2 and Theorem 2.3:

$$\|J_{r_\theta}\|_2 \approx 1 + \sigma_t^2 \frac{C}{\delta}, \quad (43)$$

$$\kappa(\mathcal{J}) \approx 1 + \Delta \left(1 + \frac{C\sigma_t^2}{\delta}\right). \quad (44)$$

As $\delta \rightarrow 0$ (approaching the sharp boundary), $\kappa(\mathcal{J}) \rightarrow \infty$. \square

D PROOFS OF ROPA’S THEORETICAL GUARANTEES

Here we formalize the guarantees for the adaptive mechanisms in ROPA. We denote the global trajectory vector by $x^{(k)}$ at Newton iteration k , and the state at time t_n by x_{t_n} . We distinguish Hessian eigenvalues ν from the damping parameters λ .

D.1 CONDITION NUMBER CONTROL

Theorem D.1 (Condition Number Bound via Adaptive Sparsity). *Let $\mathcal{J}_{\text{true}}$ be the exact Jacobian of the full coupled system. Let $\mathcal{J}^{(k)}$ be the block-banded approximation constructed by ROPA using bandwidths $w_n^{(k)}$ and damping $\lambda_{\text{damp},n}^{(k)}$. Assume the off-diagonal couplings of $\mathcal{J}_{\text{true}}$ decay exponentially with time distance (a property of parabolic diffusion operators). Then, there exist bandwidths w_n and damping factors λ_n such that:*

$$\kappa(\mathcal{J}^{(k)}) \leq \kappa_{\text{th}}.$$

Proof. We analyze the spectrum of the preconditioned operator. The condition number is determined by the spread of eigenvalues. We control this via two mechanisms: bandwidth (truncation error) and damping (eigenvalue shifting).

1. Bandwidth and Spectral Radius Control. Let $E = \mathcal{J}_{\text{true}} - \mathcal{J}_{\text{band}}^{(k)}$ be the truncation error matrix resulting from restricting the Jacobian to bandwidth $\{w_n\}$. For diffusion processes, the coupling strength between x_{t_n} and $x_{t_{n+k}}$ decays as the diffusion kernel width relative to the time gap.

By the **Gershgorin Circle Theorem**, the eigenvalues of the approximate matrix $\mathcal{J}_{\text{band}}^{(k)}$ are contained in the union of discs centered at diagonal entries, with radii equal to the sum of absolute off-diagonal entries. Increasing w_n includes more off-diagonal mass into the matrix, effectively reducing the "leakage" mass $\|E\|_\infty$ outside the band. ROPA’s adaptive rule increases w_n when residuals are high (a proxy for strong coupling). This ensures that the truncation error $\|E\|_2$ is kept below a threshold δ , keeping the spectrum of $\mathcal{J}^{(k)}$ close to the well-conditioned regime of the true operator.

2. Damping and Eigenvalue Shifting. Even with zero truncation error, the local Jacobian block J_n may be ill-conditioned due to high curvature ν_{max} . The damping operation $\mathcal{J}_\lambda = \mathcal{J} + \lambda I$ shifts the spectrum:

$$\kappa(\mathcal{J}_\lambda) = \frac{\lambda + \nu_{\text{max}}}{\lambda + \nu_{\text{min}}}. \quad (45)$$

To enforce $\kappa \leq \kappa_{\text{th}}$, we require:

$$\lambda \geq \frac{\nu_{\text{max}} - \kappa_{\text{th}}\nu_{\text{min}}}{\kappa_{\text{th}} - 1}. \quad (46)$$

ROPA’s trust-region mechanism (checking gain ratios) implicitly finds this λ . When κ is large, the standard Newton step fails to reduce residuals, causing the gain ratio to drop and triggering an increase in λ until the condition above is satisfied. Thus, $\kappa(\mathcal{J}^{(k)})$ is deterministically bounded. \square

D.2 CONVERGENCE ANALYSIS

Theorem D.2 (Local Convergence with LML Correction). *Under the bounding conditions of Theorem D.1, and assuming the LML correction is applied when alignment γ is high, the ROPA iterations converge linearly with a small contraction factor $\rho \ll 1$ (approaching superlinear) to a solution x^* on the manifold \mathcal{M} .*

Proof. Consider the error propagation $e_{k+1} = x^{(k+1)} - x^*$. The approximate Newton update is:

$$x^{(k+1)} = x^{(k)} - (\mathcal{J}^{(k)})^{-1} \mathcal{R}(x^{(k)}).$$

Standard perturbation theory for Newton methods gives the error bound:

$$\|e_{k+1}\| \leq \underbrace{\|\mathcal{J}^{-1}(\mathcal{J} - \mathcal{J}^{(k)})\|}_{\text{Approximation Error}} \|e_k\| + \underbrace{C\|e_k\|^2}_{\text{Newton Quadratic Term}}. \quad (47)$$

Convergence requires the linear coefficient (contraction factor) to be < 1 . The term $(\mathcal{J} - \mathcal{J}^{(k)})$ represents the error in the Jacobian approximation. In high-curvature regions, this error is dominated by the stiffest eigenvector v_{\max} corresponding to ν_{\max} . The **LML Correction** (Eq. 12) explicitly constructs a rank-1 approximation of this inverse Hessian component:

$$H_{\text{LML}}^{-1} \approx (\mathcal{J}_{\text{stiff}})^{-1}.$$

By substituting this correction into the update rule when alignment is detected, ROPA effectively "preconditions" the stiffest direction, rendering the term $\|\mathcal{J}^{-1}(\mathcal{J} - \mathcal{J}^{(k)})\| \approx 0$ along the normal vector of \mathcal{M} . For tangent directions, the adaptive bandwidth ensures the error is small. Thus, the contraction factor ρ is minimized, ensuring robust convergence $e_{k+1} \leq \rho e_k$ even in stiff regimes where standard parallel solvers diverge. \square

D.3 COMPLEXITY ANALYSIS

Theorem D.3 (Expected Linear Complexity). *The expected computational cost per Newton step of ROPA is $\mathcal{O}(N)$, where N is the number of time steps.*

Proof. The complexity is dominated by the linear solve of the block-banded system. For a block-banded matrix of size $N \times N$ (block size d) with bandwidth w , the Cholesky or LU factorization cost is $\text{Cost}(w) \approx N \cdot d \cdot (w \cdot d)^2 = \mathcal{O}(Nw^2)$.

The bandwidth w_n is adaptive. From **Assumption 2.1 (Anisotropy Index)****, the manifold \mathcal{M} exhibits high curvature (requiring large w_{\max}) only on a measurable subset $\mathcal{M}_{\text{curv}}$. Let $p_{\text{stiff}} = \mu(\mathcal{M}_{\text{curv}})/\mu(\mathcal{M})$ be the probability of the trajectory traversing a high-curvature region. The expected bandwidth is:

$$\mathbb{E}[w] = p_{\text{stiff}} \cdot w_{\max} + (1 - p_{\text{stiff}}) \cdot w_{\text{base}}. \quad (48)$$

Since w_{\max} is a small constant (typically $8 \sim 16$) independent of N , the expected bandwidth is $\mathcal{O}(1)$. Therefore, the expected total cost is:

$$\mathbb{E}[\text{Cost}] = \sum_k \mathcal{O}(N\mathbb{E}[w]^2) = \mathcal{O}(N). \quad (49)$$

This confirms that ROPA scales linearly with sequence length, preserving the efficiency advantage of parallel sampling. \square