Bayesian Computation with Generative Diffusion Models via Multilevel Monte Carlo

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Outline

- Motivation
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- Multilevel Monte Carlo (MLMC)
- Numerical Results and Discussion

Motivation: The Promise of Diffusion Models

- **Diffusion Models:** Class of stochastic samplers that rely on neural networks to generate samples from a posterior distribution.
- High-quality sampling: DMs provide state-of-the-art sample quality for complex distributions.
- **Flexible conditioning**: Handle diverse inverse problems (denoising, inpainting, super-resolution) through unified framework.
- Well-defined probability model enables proper UQ.

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The Computational Challenge

- Large-scale UQ: Reliable uncertainty estimates may require 10⁴-10⁶ samples, especially when variability is significant.
- **Slow sampling**: Typical DM requires 10^2 - 10^3 neural function evaluations (NFEs) per sample.
- Prohibitive cost: Full UQ analysis can require 10⁷-10⁹ NFEs.

Motivation: Why Diffusion Models for UQ?

The need for speed

- Practical deployment in time-sensitive applications (medical imaging, real-time systems).
- Large-scale uncertainty studies (global sensitivity analysis, robust optimization).
- Hyperparameter tuning (prior selection, likelihood calibration).

Solutions:

- Reduce cost per NFE (pruning, quantization).
- Reduce number of NFE's per sample (distillation).
- Reduce number of NFE's per statistic computation.

Bayesian framework

- Aim of a Machine Learning algorithm: Estimate $x \in \mathbb{R}^n$, $n \gg 1$, give observed data y.
- Bayesian framework: $Y \sim \mathcal{P}(\mathcal{A}(x))$, conditional on X = x.
- Sampling from the conditional density of X, $p(\cdot|y) \propto \mathcal{L}(y|\cdot)\pi(\cdot)$, for likelihood \mathcal{L} and prior π .
- Our goal is to quantify the uncertainty that such a probability distribution entails.

Denoising Diffusion Probabilistic Models

- Finite sequence of noising kernels, or a continuous-time SDE.
- Build sequence of states $\{X_i\}_{i=0}^T$ with $X_0 = X$.
- Sample from the joint density of $(X_0, ... X_T)$ given Y = y; joint density admits the desired conditional density as marginal.
- Write

$$p_{0:T}(x_0,...,x_T|y) = \left(\prod_{t=1}^T \widehat{X}_{t-1}(x_{t-1}|x_t,y)\right) p_T(x_T|y),$$

for **reverse transition kernels**, $(\widehat{\lambda}_t)_{t=0}^{T-1}$, and a *pre-determined* posterior density $p_T(\cdot|y)$ of the final state X_T .

• For example, $p_T(\cdot | y) \in \{\phi_n(\cdot), \delta_v(\cdot), \phi_n(\cdot; (\mathbb{I}_n - \mathbb{M})y, \mathbb{M})\}.$

DDPMs: Forward transition kernels, K

Gaussian, Markovian forward process:

$$K_t(x|x_{t-1}) = \phi_n\left(x; \sqrt{\frac{\gamma_t}{\gamma_{t-1}}} x_{t-1}, \left(1 - \frac{\gamma_t}{\gamma_{t-1}}\right) \mathbb{I}_n\right),$$

for $x \in \mathbb{R}^n$ and where $1 = \gamma_0 > \gamma_1 > \ldots > \gamma_T > 0$.

Hence

$$K_{t;0}(x|x_0) = \phi_n(x; \sqrt{\gamma_t}x_0, (1-\gamma_t)\mathbb{I}_n), \quad 1 \leq t \leq T,$$

and we define the score function, for any $x \in \mathbb{R}^n$,

$$\nabla \log K_{t;0}(x_t|x_0) = (\sqrt{\gamma_t}x_0 - x_t)/(1 - \gamma_t),$$

and we solve for x_0

$$x_0(x_t; y, t) = \frac{1}{\sqrt{\gamma_t}}(x_t + (1 - \gamma_t)\nabla \log K_{t;0}(x_t|x_0))$$

DDPMs: Reverse Kernel, X

By conditioning on the initial state and assuming intermediate states are independent of the observation Y,

$$\widehat{\lambda}_{t-1}(x|x_t,y) = \int_{\mathbb{R}^n} \lambda_{t-1;0,t}(x|x_0,x_t) \widehat{\lambda}_{0;t}(x_0|x_t,y) dx_0.$$

We use

$$\widehat{\mathsf{A}}_{0;t}(\cdot|x_t,y)=\delta_{\widehat{\mathsf{x}}_0}(\cdot)$$

where

$$\widehat{x}_0(x_t; y, t) = \frac{1}{\sqrt{\gamma_t}}(x_t + (1 - \gamma_t)\nabla \log K_{t;0}(x_t|x_0))$$

Finally,

$$\lambda_{t-1;0,t}(x|x_0,x_t) = \phi_n(\cdot,\mu_{t-1;0,t},\sigma_{t-1;0,t}^2\mathbb{I}_n)$$

for some known mean and variance.

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where

$$\widehat{x}_0(x_t; y, t, \theta) = \frac{1}{\sqrt{\gamma_t}}(x_t + (1 - \gamma_t) \quad s_\theta(x_t, y, t))$$

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DDPMs: Four Sources of Errors

$$\begin{split} x_{t-1} &= \sqrt{\frac{\gamma_{t-1}}{\gamma_t}} x_t - \sqrt{\frac{\gamma_t}{\gamma_{t-1}}} \left(\Gamma_{t-1} \ - \frac{\gamma_{t-1}}{\gamma_t} \Gamma_t \right) s_\theta(x_t, y, t) \\ &+ \sqrt{\frac{\Gamma_{t-1}}{\Gamma_t} \left(1 - \frac{\gamma_t}{\gamma_{t-1}} \right)} \xi_t, \qquad \text{where } \Gamma_t = 1 - \gamma_t \end{split}$$

- **Model error**: From imperfect score network training based on a finite training-set.
- Finite-time error: We have to choose $\gamma_T > 0$ (to ensure γ_{t-1}/γ_t is small), hence X_T will not be exactly Gaussian (not fully diffused).
- Truncation error: The posterior $p(\cdot | y)$ has smaller support than $p_t(\cdot | y)$, leading to blow up in s_θ as $t \to 0$; need to stop early or have a Gaussian approximation independent of the score.
- **Discretization error** (our focus today): We can skip M steps, sampling directly x_{t-M} given x_t .

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Monte Carlo Estimation

- Goal: Estimate E[f(X)], where $X \sim p(\cdot)$.
- Monte Carlo estimator:

$$\mathsf{E}[f(X)] \approx \mathsf{E}[f(\widehat{X})] \approx \frac{1}{N} \sum_{i=1}^{N} \widehat{x}^{(i)},$$

- Variance $\frac{1}{N}$ Var $[f(\widehat{X})]$, Bias $|E[f(X)] E[f(\widehat{X})]|$.
- For RMSE ε : Number of samples grow as $O(\varepsilon^{-2})$ to reduce variance. Cost per sample grows as ε decreases to decrease bias.

Multilevel Monte Carlo: Basic Identity

- Use a hierarchy of approximations: $\widehat{X}_0, \widehat{X}_1, \dots, \widehat{X}_L$ with increasing cost and accuracy (each adding M intermediate steps compared to the previous approximation).
- Telescoping sum:

$$\mathsf{E}[f(\widehat{X}_{L})] = \mathsf{E}[f(\widehat{X}_{0})] + \sum_{\ell=1}^{L} \mathsf{E}[f(\widehat{X}_{\ell}) - f(\widehat{X}_{\ell-1})]$$

Define level estimators:

$$Y_\ell = rac{1}{N_\ell} \sum_{i=1}^{N_\ell} \left(f(\widehat{x}_\ell^{(i)}) - f(\widehat{x}_{\ell-1}^{(i)})
ight)$$
 with coupled samples

Total estimator:

$$Y = \sum_{\ell=0}^{L} Y_{\ell}$$
 with $Y_0 = \frac{1}{N_0} \sum_{i=1}^{N_0} f(x_0^{(i)})$

MLMC Efficiency and Complexity

 $\begin{array}{ccc} \bullet & \mathsf{Bias:} & |\mathsf{E}[\,f(\widehat{X}_\ell) - f(\widehat{X}_{\ell-1})\,]| \sim & M^{-\alpha\ell} \\ \\ \mathsf{Variance:} & \mathsf{Var}[\,f(\widehat{X}_\ell) - f(\widehat{X}_{\ell-1})\,] =: \, V_\ell \sim & M^{-\beta\ell} \\ \\ \mathsf{Sampling cost:} & C_\ell \sim & M^\ell \end{array}$

ullet Optimal total cost: Choosing N_ℓ to minimize cost, yields

$$C_{\mathsf{MLMC}} \lesssim \quad \varepsilon^{-2} \left(\sum_{\ell=0}^{L} \sqrt{V_{\ell} C_{\ell}} \right)^{2}$$

$$\lesssim \begin{cases} \varepsilon^{-2} V_{0} C_{0} & V_{\ell} C_{\ell} \to 0 \\ \varepsilon^{-2} L^{2} & V_{\ell} C_{\ell} \approx \mathrm{const} \\ \varepsilon^{-2} V_{L} C_{L} & V_{\ell} C_{\ell} \to \infty \end{cases}$$

Compare with standard MC at finest level:

$$C_{\rm MC} \lesssim \varepsilon^{-2} V_0 C_L \Rightarrow {\rm MLMC}$$
 much cheaper if $V_\ell C_\ell \downarrow$

Need for Exponential Integrators

• Diffusion models discretise stiff SDEs in reverse:

$$dX_t = \left(A_t X_t - B_t^2 \nabla \log K_{t;0}(x_t | x_0)\right) dt + B_t dW_t$$

- As $t \to T$, we have $\gamma_t \to 0$, hence A_t becomes unbounded, leading to instability.
- Exponential integrators mitigate instability due to the linear terms:

$$\begin{aligned} x_{t-M}^r &= & e^{-\int_t^{t-M} A_\tau d\tau} x_t \\ &- s_\theta(x_t, y, t) \int_t^{t-M} e^{\int_s^{t-M} A_\tau d\tau} B_s^2 \, \mathrm{d}s \\ &+ \int_t^{t-M} e^{\int_s^{t-M} A_\tau d\tau} B_s \, \mathrm{d}W_s \end{aligned}$$

• Crucially, correlating the fine and coarse paths needs to take into account previous variance coefficient.

Three Imaging Inverse Problems

left to right: truth x; observation y; posterior sample from a DM.



















- Super-resolution: y = Ax, A downsampling, ill-posed
- ② Denoising: $y = x + \eta$, high noise
- **1** Inpainting: y = Mx, partial masking

Use $f(x) = x^2$ to estimate marginal second moment (modelling pixel-wise uncertainty).

Convergence Rates

• Empirically fit α and β :

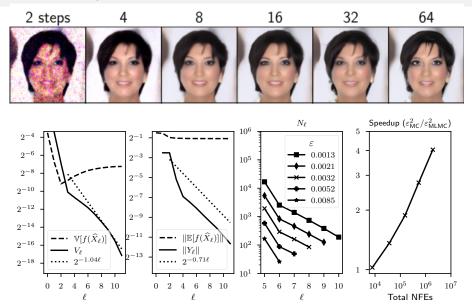
$$\left| \mathsf{E}[f(X) - f(\widehat{X}_{\ell})] \right| \sim M^{-\alpha \ell}$$

and

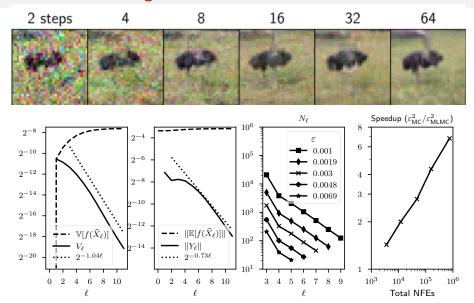
$$\mathsf{Var}[f(\widehat{X}_\ell) - f(\widehat{X}_{\ell-1})] \sim M^{-\beta\ell}$$

• Expected: $\alpha = 1$ and $\beta = 2$; corresponding to rates from Milstein scheme.

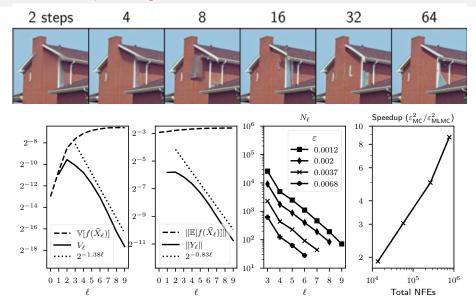
Results: Super-resolution



Results: Denoising



Results: Inpainting



Concluding Remarks

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- MLMC is a powerful variance reduction tool for diffusion-based Bayesian inference, MLMC reduces NFEs by 4–9 times for fixed accuracy.
- Requires careful time discretisation and coupling.
- Need better analysis of the approximation of the score function to understand degradation of convergence rates.
- Future work
 - Combine with distillation and quantisation.
 - Multilevel training of DMs.
 - Tackle cost associated to other approximations parameters (model, finite-time and truncation errors).