# Score-based diffusion models for imaging inverse problems



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Acknowledgments:

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#### J. Fessler Score



#### Introduction

Inverse problems

Generative models

Score matching / diffusion models

#### Patch-based models

Non-overlapping patch model

Patch Diffusion Inverse Solver (PaDIS)

CT reconstruction results

3D CT reconstruction

#### Distribution shifts

Summary

Book

**Bibliography** 

### Under-determined inverse problems

► Applications: compressed sensing MRI, sparse-view CT, PET, inpainting, ...
All have *linear* forward models for data:

$$y = Ax + \varepsilon$$

y: sensor data (e.g., sinogram)

A: wide system matrix (known)

x: latent image (or image series in dynamic problems)

 $\varepsilon$ : noise with known distribution provides likelihood p(y|x)

► Maximum-likelihood estimation (physics-based fitting) is usually non-unique:

$$\hat{x} = \underset{x}{\operatorname{arg \, max}} \log p(y|x) = \underbrace{\underset{x}{\operatorname{arg \, min}} \|Ax - y\|_{2}^{2}}_{x}$$
(for gaussian noise)

▶ Minimum-norm least-squares solution is unique but usually impractical or useless:

$$\hat{\mathbf{x}} = \mathbf{A}^{+}\mathbf{y} = \mathbf{y}$$
 for inpainting problem



hand-crafted regularizers:

$$\hat{\mathbf{x}} = \arg\min_{\mathbf{x}} - \log p(\mathbf{y}|\mathbf{x}) + R(\mathbf{x}) = \arg\min_{\mathbf{x}} \frac{1}{2\sigma_{\varepsilon}^2} \|\mathbf{A}\mathbf{x} - \mathbf{y}\|_2^2 + R(\mathbf{x})$$

black-box data-driven supervised methods:

$$\mathbf{A}^{+}\mathbf{y} 
ightarrow \boxed{\mathsf{NN}} 
ightarrow \hat{\mathbf{x}}$$

- unrolled deep learning methods (PNP, RED, MoDL, ...)
- ▶ Bayesian methods (e.g., MAP) based on a prior p(x), lately (?) relabeled as generative models (or "genAI")

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▶ black-box data-driven supervised methods:

$$\mathbf{A}^+\mathbf{y} o \boxed{\mathsf{NN}} o \hat{\mathbf{x}}$$

- unrolled deep learning methods (PNP, RED, MoDL, ...)
- ▶ Bayesian methods (e.g., MAP) based on a prior p(x), lately (?) relabeled as generative models (or "genAI")
- ► Appeal:
  - $\circ$  PNP-like training independent of **A** or p(y|x)
  - o Strong priors for complex systems with aggressive under-sampling
  - $\circ$  Posterior sampling from p(x|y) for uncertainty quantification

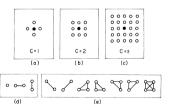
## Long history of Bayesian models for inverse problems



#### Markov random field models

$$p(\textbf{\textit{x}}) \propto \prod_c \mathrm{e}^{-\textit{U}_c(\textbf{\textit{x}}_c)}$$

(e.g.) Geman & Geman 1984 [1]



Mostly for inference?

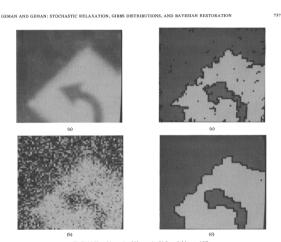


Fig. 7. (a) Blurred image (roadside scene). (b) Degraded image: Additive noise. (c) Restoration including line process; 100 iterations. (d) Restoration including line process; 1000 iterations.

### Long history of generative models and inverse problems



MRF as generators?

[2] T-PAMI 1994

#### An Empirical Study of the Simulation of Various Models Used for Images

A. J. Gray, J. W. Kay, and D. M. Titterington

Abstract- Markov random fields are typically used as priors in Bayesian image restoration methods to represent spatial information in the image. Commonly used Markov random fields are not in fact capable of representing the moderate-to-large scale clustering present in naturally occurring images and can also be time consuming to simulate,













#### Gray, Kay, Titterington [2] T-PAMI 1994

... the local properties of spatial Markov models are undoubtedly plausible descriptors of the local associations typical of many images, which is the way in which the models are often used. Nevertheless. it would be reassuring if models used as priors did in fact provide a realistic representation of our prior assumptions and if their (empirical) properties were more widely known.

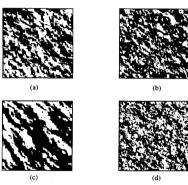


Fig. 4. Realizations of two-dimensional, one-parameter, autologistic Markov Mesh models: (a) binary, second-order model with  $\beta = \log 5$ ; (b) three-color second-order model with  $\beta = \log 5$ ; (c) binary second-order model with  $\beta = \log 3$ .

# Generative models are hot in imaging / inverse problems





Zhao, Ye, Bresler: Jan. 2023 IEEE SpMag survey paper [3]

- ► Generative adversarial network (GAN) models
- ► Variation auto-encoder (VAE) models [4]
- ► Normalizing flows [5, 6]
- Score-based diffusion models
  - o Zaccharie Ramzi et al., NeurIPS Workshop 2020 [7]
  - o Yang Song & Liyue Shen et al., NeurIPS Workshop 2021, ICLR 2022 [8, 9]
  - o Ajil Jalal et al. ... Jon Tamir, NeurlPS 2021 [10]
  - o Hyungjin Chung & Jong Chul Ye, MIA, Aug. 2022 [11]
  - o Luo et al., MRM, 2023 [12]
  - o ...
- ► Kazerouni et al. [13] have github catalog, including >20 (!) survey papers
- ightharpoonup (hopelessly incomplete lists) Common aim: model/learn prior p(x)

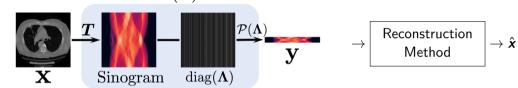


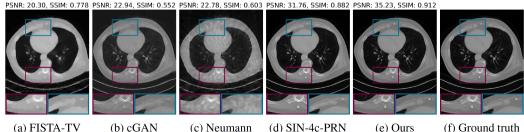


From Song & Shen et al., ICLR 2022 [9].

Trained with 47K 2D CT images. Recon 23 projection views ( $\approx$  17-fold dose reduction)

$$\boldsymbol{A} = \mathcal{P}(\boldsymbol{\Lambda})\boldsymbol{T}$$





### Challenges with Bayesian models



- 1. Learning whole-image prior models requires many high-quality training images Some applications like dynamic MRI have few *if any* realistic training samples
  - Curse of dimensionality
  - o Images live near manifolds (unsuitable for traditional density estimators)
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2



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  - GPU memory
  - o training data requirements

3.

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- 4.

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- 4. What does "uncertainty" mean if prior is misspecified?



▶ Bayesian inference methods use the posterior:

$$p(x|y) = \underbrace{p(y|x)}_{\text{physics}} \underbrace{p(x)}_{\text{prior}} / p(y)$$

- Here the prior p(x) is for quantifying (prior) probability, not necessarily for generation.
- A model for the posterior p(x|y) opens many doors:
  - Maximizing p(x|y) is maximum a posteriori (MAP) estimation
  - ▶ The conditional mean  $E[x|y] = \int x p(x|y) dx$  is the MMSE estimator
  - $\triangleright$  Sampling from the posterior p(x|y) facilitates uncertainty quantification in inference
- All these methods require the prior p(x), i.e., a prior model  $p(x; \theta)$ .

# Bayesian methods (generative models)



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- Or do they?



Sampling from a *prior*  $p(x; \theta)$  just needs its score function  $\nabla_x \log p(x; \theta)$ , using Langevin dynamics, aka stochastic gradient ascent of log-prior:

$$\mathbf{x}_{t} = \mathbf{x}_{t-1} + \alpha_{t} \underbrace{\nabla \log p(\mathbf{x}_{t-1}; \boldsymbol{\theta})}_{\text{score function}} + \beta_{t} \underbrace{\varepsilon_{t}}_{\text{t}}, \quad t = 1, \dots, T.$$

- $\circ$  Draws samples from  $p(x; \theta)$  for suitable choices of  $\{\alpha_t\}$ ,  $\{\beta_t\}$ , and (large) T [14].
- $\circ$  If  $\alpha_t = 0$  and  $\beta_t = \beta$ , then akin to (isotropic) diffusion or Brownian motion



- ► Typical distribution models:  $p(x; \theta) = \frac{1}{Z(\theta)} e^{-U(x; \theta)}$ . Goal: learn  $\theta$  from training data  $x_1, \dots, x_T$
- ▶ For IID samples  $\{x_t\}$ , one could try to learn  $\theta$  by ML estimation:

$$\hat{m{ heta}} = rg \max_{m{ heta}} p(m{x}_1, \dots, m{x}_T; m{ heta}) = rg \max_{m{ heta}} \sum_{t=1}^T \log(p(m{x}_t; m{ heta}))$$

$$= rg \max_{m{ heta}} \left( -T m{Z}(m{ heta}) + \sum_{t=1}^T -U(m{x}_t; m{ heta}) \right).$$

Typically intractable due to the partition function  $Z(\theta)$ .

# Distribution learning vs score learning



- ► Typical distribution models:  $p(\mathbf{x}; \boldsymbol{\theta}) = \frac{1}{Z(\boldsymbol{\theta})} e^{-U(\mathbf{x}; \boldsymbol{\theta})}$ . Goal: learn  $\boldsymbol{\theta}$  from training data  $\mathbf{x}_1, \dots, \mathbf{x}_T$
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Typically intractable due to the partition function  $Z(\theta)$ .

In contrast, the score function is easier to handle:

$$s(x; \theta) \triangleq \nabla_x \log p(x; \theta) = \nabla_x (-\log Z(\theta) - U(x; \theta)) = -\nabla_x U(x; \theta).$$



- ▶ Given training data  $x_1, ..., x_T$ , learn score function  $s(x; \theta) \stackrel{?}{=} \nabla_x \log p(x; \theta)$



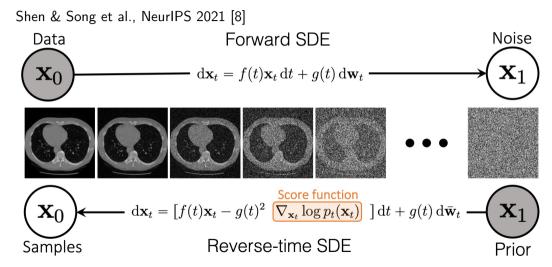
- ▶ Given training data  $x_1, ..., x_T$ , learn score function  $s(x; \theta) \stackrel{?}{=} \nabla_x \log p(x; \theta)$
- Explicit score matching (ESM) (Hyvärinen, 2005 [15])
- ► Implicit score matching (ISM)
- ▶ Denoising score matching (DSM) (Vincent, 2011 [16])
- ▶ Noise-conditional score matching (NCSM) (Song, 2019 [17, eqn. (5)]):

$$\ell(\boldsymbol{\theta}; \sigma) \triangleq \frac{1}{2} \, \mathsf{E}_{\mathsf{q}_0(\boldsymbol{x})} \bigg[ \mathsf{E}_{\boldsymbol{g}_{\sigma}(\boldsymbol{z})} \bigg[ \bigg\| \boldsymbol{s}(\boldsymbol{x} + \boldsymbol{z}; \boldsymbol{\theta}, \sigma) + \frac{\boldsymbol{z}}{\sigma^2} \bigg\|_2^2 \bigg] \bigg], \quad \mathcal{L}(\boldsymbol{\theta}; \{\sigma_I\}) = \frac{1}{L} \sum_{I=1}^L \sigma_I^2 \, \ell(\boldsymbol{\theta}; \sigma_I),$$

where  $s(x; \theta, \sigma)$  denotes a noise-conditional score network (NCSN).

- ▶  $d(x; \theta) \triangleq x + \sigma^2 s(x; \theta, \sigma)$ : equivalent image denoiser by Tweedie's formula [18]
- ▶ Recommended choice [19]:  $s(x; \theta, \sigma) \triangleq \tilde{s}(x; \theta)/\sigma$ , where  $\tilde{s}$  is unitless





### Score-based diffusion models: trade-offs





- ► No adversarial training needed
- ► High quality sample generation (if enough training data)

## Score-based diffusion models: trade-offs



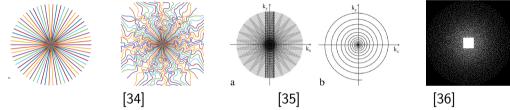
- ► No adversarial training needed
- ► High quality sample generation (if enough training data)
- ► Expensive sample generation (vs GAN models)
  - o Distillation methods [20]
  - o Consistency models [21]
  - o Geometric decomposition [22]
  - o Multi-scale [23, 24] and pyramidal [25] and coarse-to-fine [26] models
  - Faster ODE solvers [27]
  - o Warm starts [28]
  - Latent diffusion models: use VAE and diffuse in latent space [29–31].
     Used in Stable Diffusion by start-up Stability AI
  - o 3D image reconstruction using 2D models [32, 33]
- ► Learning 3D (or 3D+T) whole-image generative models is challenging (training data, GPU memory, ...)





Jan. 2023 survey paper on generative models [3] does not mention "patch" once!?

#### MRI k-space sampling:



Patch-based models have long history in inverse problems, e.g.,

- patch GAN [37–39]
- patch dictionary models [40, 41]
- non-local means, BM3D
- Wasserstein patch prior [42, 43] ...



➤ Can patch-based generative models be effective priors for inverse problems in applications with very limited training data? e.g., dynamic MRI

► Can patch-based generative models provide better robustness to distribution shifts, perhaps at the cost of reduced in-distribution performance?

► Can we use the "latest" generative models, e.g., score-based models, for patches?

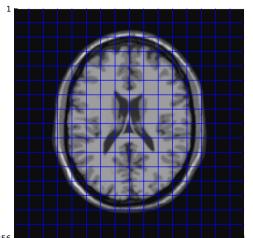
## Patch diffusion model: Simple version



#### Warm up:

simple, but less effective, approach:

- Fixed patch size
- Fixed patch grid
- No position information (Fessler, Hu, Xu, BASP 2023 [46])



256



▶ Start with MRF formulation, aka *fields of experts* model [51–53] for image **x**:

$$p(\boldsymbol{x};\boldsymbol{\theta}) = \frac{1}{Z(\boldsymbol{\theta})} e^{-\sum_{c} V_{c}(\boldsymbol{x};\boldsymbol{\theta})} = \frac{1}{Z(\boldsymbol{\theta})} \prod_{c} e^{-V_{c}(\boldsymbol{x};\boldsymbol{\theta})}.$$

- ullet heta : parameter vector that describes the prior
- $V_c$ : clique potential for the cth image patch
- $Z(\theta)$  : (intractable) partition function
- Assume (temporarily) statistical spatial stationarity (image shift invariance):

$$V_c(\mathbf{x}; \boldsymbol{\theta}) = V(\mathbf{G}_c \mathbf{x}; \boldsymbol{\theta})$$

- $oldsymbol{G}_c$  : wide binary matrix that grabs pixels of the cth patch from image  $oldsymbol{x}$
- $V(\mathbf{v}; \boldsymbol{\theta})$ : common patch clique function



Resulting log-prior:

$$\log p(\mathbf{x}; \boldsymbol{\theta}) = -\log Z(\boldsymbol{\theta}) - \sum_{c} V(\mathbf{G}_{c}\mathbf{x}; \boldsymbol{\theta})$$

► Corresponding overall *image score function* arises from *patch score function*:

$$\mathbf{s}(\mathbf{x};\boldsymbol{\theta}) \triangleq \nabla_{\mathbf{x}} \log p(\mathbf{x};\boldsymbol{\theta}) = \sum_{c} \mathbf{G}'_{c} \mathbf{s}_{V}(\mathbf{G}_{c}\mathbf{x};\boldsymbol{\theta}), \qquad \mathbf{s}_{V}(\mathbf{v};\boldsymbol{\theta}) \triangleq -\nabla_{\mathbf{v}} V(\mathbf{v};\boldsymbol{\theta}).$$

- ▶ All we must learn is the patch score function  $s_V(\mathbf{v}; \theta) : \mathbb{R}^n \to \mathbb{R}^n$ , e.g., a UNet.
- ► For non-overlapping patches:

$$\underbrace{\left\| \mathbf{s}(\mathbf{x} + \mathbf{z}; \boldsymbol{\theta}) + \mathbf{z}/\sigma^{2} \right\|_{2}^{2}}_{\text{image "denoise"}} = \left\| \sum_{c} \mathbf{G}_{c}' \mathbf{s}_{V}(\mathbf{G}_{c}(\mathbf{x} + \mathbf{z}); \boldsymbol{\theta}) + \mathbf{z}/\sigma^{2} \right\|_{2}^{2} \\
= \sum_{c} \underbrace{\left\| \mathbf{s}_{V}(\mathbf{x}_{c} + \mathbf{z}_{c}); \boldsymbol{\theta} \right) + \mathbf{z}_{c}/\sigma^{2} \right\|_{2}^{2}}_{\text{patch "denoise"}}, \quad \mathbf{z}_{c} \triangleq \mathbf{G}_{c} \mathbf{z}$$



► For training image patches  $\{v_1, \dots, v_T\}$ , apply denoising score matching (DSM) of Vincent, 2011 [16], typically for a range of noise variances  $\sigma^2$  [14]:

$$\hat{\boldsymbol{\theta}} = \arg\min_{\boldsymbol{\theta}} \frac{1}{T} \sum\nolimits_{t=1}^{T} \mathsf{E}_{\sigma \sim p(\sigma)} \Bigg[ \sigma^2 \, \mathsf{E}_{\mathbf{z} \sim \mathcal{N}(\mathbf{0}, \sigma^2 \mathbf{I}_n)} \Bigg[ \frac{1}{2} \, \bigg\| \mathbf{s}_V(\mathbf{v}_t + \mathbf{z}; \boldsymbol{\theta}, \sigma) + \frac{\mathbf{z}}{\sigma^2} \bigg\|_2^2 \Bigg] \Bigg] \, .$$

- Final patch score model is  $\mathbf{s}_V(\mathbf{v}; \hat{\boldsymbol{\theta}}, \sigma_{\min})$ .



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- Network input is just image patches, never the entire image ⇒ scales to large 2D images, 3D, 4D, etc.



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- Final patch score model is  $\mathbf{s}_V(\mathbf{v}; \hat{\boldsymbol{\theta}}, \sigma_{\min})$ .
- Network input is just image patches, never the entire image ⇒ scales to large 2D images, 3D, 4D, etc.
- ► Drawbacks:
  - Visible patch boundaries
  - o Fixed patch size slows learning
  - Suboptimal stationarity assumption (cf. vertebrae)

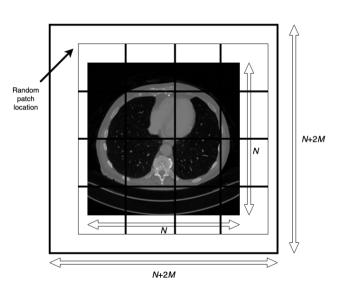
## Improved patch modeling

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- ▶ zero-pad image x
- use multiple grid locations

#### Inspirations:

- Wavelet "cycle spinning"[47, 54–57]
- $\circ$  Wang, NeurIPS 2023 [58]





- $\triangleright$   $N_1 \times N_2$ : original image size
- $ightharpoonup P_1 \times P_2$ : patch size
- $ightharpoonup K_i \triangleq 1 + |N_i/P_i|, i = 1, 2: \# non-overlapping patches for original image$
- $\triangleright$   $(N_1 + 2M_1) \times (N_2 + 2M_2)$ : padded image size;  $M_i \triangleq K_i P_i N_i$
- ▶ Product probability model:

$$p(\mathbf{x}) \triangleq \frac{1}{Z} \underbrace{\prod_{m=1}^{M_1 M_2}}_{\text{grid}} \underbrace{\left(\underbrace{p_{m,B}(\mathbf{x}_{m,B})}_{\text{border}} \underbrace{\prod_{k=1}^{K_1 K_2} p_{m,k}(\mathbf{x}_{m,k})}_{\text{patches}}\right) = \frac{1}{Z} \underbrace{\prod_{m=1}^{M_1 M_2} \prod_{k=1}^{K_1 K_2} \underbrace{e^{-V(\mathbf{x}_{m,k}; \mathbf{m}, \mathbf{k})}}_{\text{position}}}_{\text{encoding}}$$

- $\circ x_{m,B}$ : border pixels for mth shift (all zero)
- $\circ x_{m,k}$ : kth patch for mth shift



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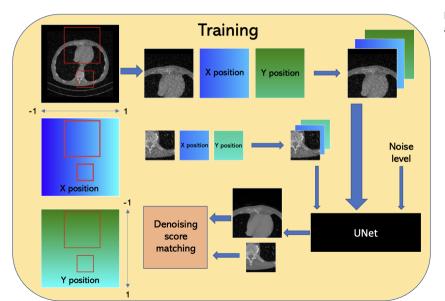
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- $\circ x_{m,B}$ : border pixels for mth shift (all zero)
- $\circ \mathbf{x}_{m,k}$ : kth patch for mth shift
- Learn position-dependent patch score function  $s(\mathbf{v}; \boldsymbol{\theta}, m, k) = -\nabla_{\mathbf{v}} V(\mathbf{v}; m, k)$

## Patch Diffusion Inverse Solver (PaDIS): Training

### J. Fessler Score





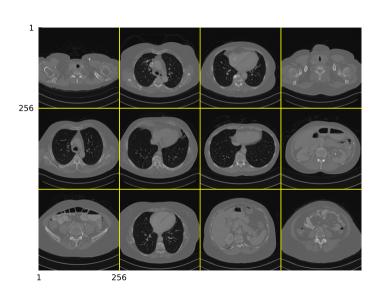
NeurIPS 2024 [60] arXiv 2406.02462

# Training images (CT)



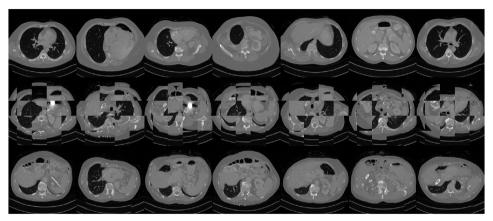
AAPM 2016 CT challenge data [61]; 10 3D volumes, rescaled to 256<sup>3</sup>

Example slices:



# Image generation (unconditional sampling from prior)





- Top: generation with a network trained on whole images (2D...)
- o Middle: patch-only version of [58] (non-overlapping patches).
- o Bottom: generation with proposed PaDIS prior.

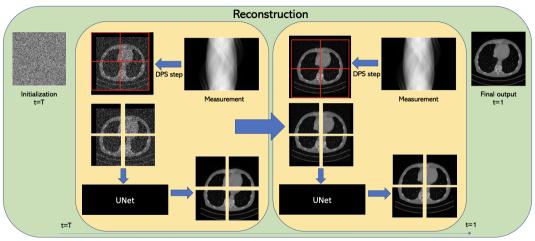


- 2 A40 GPUs using PyTorch and ADAM
- ▶ whole image model: 24 − 36 hours
- ightharpoonup patch-based model: pprox 12 hours

## Patch Diffusion Inverse Solver (PaDIS): Reconstruction







Diffusion posterior sampling (DPS) (Chung et al., ICLR 2023 [62]) with Langevin dynamics, modified to use patch score with random grid shifts.

### PaDIS algorithm (modified from DPS)



J. Fessler

Score

Input: y, A, T,  $\sigma_1 < \sigma_2 < \ldots < \sigma_T$ ,  $\epsilon > 0$ ,  $\{\zeta_t > 0\}$ ,  $P_1, P_2, M_1, M_2$ , trained noise-conditional, position-encoded patch denoiser  $d(\cdot; \theta_*, m, k, \sigma)$  Initialize random image  $x \sim \mathcal{N}(\mathbf{0}, \sigma_T^2 \mathbf{I})$ 

for t = T : 1 do

Randomly select grid integer  $m \in \{1, ..., M_1 M_2\}$  for  $k = 1 : (K_1 K_2)$  do (parallelizable)

Extract patch  $x_{m,k}$ 

Denoise patch:  $d_{m,k} \triangleq d(\mathbf{x}_{m,k}; \boldsymbol{\theta}_*, m, k, \sigma_t)$  end for

Combine denoised patches to get denoised image **d**Compute image score function:  $\mathbf{s} = (\mathbf{d} - \mathbf{x})/\sigma^2$ 

Compute image score function:  $\mathbf{s} = (\mathbf{d} - \mathbf{x})/\sigma_t^2$ Data term:  $\mathbf{x} := \mathbf{x} - \zeta_t \nabla_{\mathbf{x}} || \mathbf{A} \mathbf{d}(\mathbf{x}) - \mathbf{y}||_2^2$ 

Sample  $\mathbf{z} \sim \mathcal{N}(\mathbf{0}, \sigma_t^2 \mathbf{I})$ Step size  $\alpha_t \triangleq \epsilon \, \sigma_t^2$ 

Langevin update:  $\mathbf{x} := \mathbf{x} + \frac{\alpha_t}{2}\mathbf{s} + \sqrt{\alpha_t}\mathbf{z}$  end for

31 / 64



#### Default setup:

- 9 of 10 volumes for training ⇒ 2304 slices
- 25 slices of 10th volume for testing
- 512 element parallel-beam CT detector
- A from Operator Discretization Library (ODL)
- $56 \times 56$  patch size
- U-Net of Karras 2022 [59]
- Step size  $\zeta_t = \zeta/\|{m A}{m d}({m x}_t) {m y}\|_2$
- 1000 neural function evaluations (NFEs) [59]

### Quantitative results on three different inverse problems



Method	CT, 20 Views		CT, 8	Views	Deblu	ırring	Superresolution	
ivietnoa	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑
Baseline	24.93	0.595	21.39	0.415	24.54	0.688	25.86	0.739
ADMM-TV	26.82	0.724	23.09	0.555	28.22	0.792	25.66	0.745
PnP-ADMM [63]	26.86	0.607	22.39	0.489	28.82	0.818	26.61	0.785
PnP-RED [64]	27.99	0.622	23.08	0.441	29.91	0.867	26.36	0.766
Whole image diffusion	32.84	0.835	25.74	0.706	30.19	0.853	29.17	0.827
Langevin dynamics [17]	33.03	0.846	27.03	0.689	30.60	0.867	26.83	0.744
Predictor-corrector [11]	32.35	0.820	23.65	0.546	28.42	0.724	26.97	0.685
VE-DDNM [65]	31.98	0.861	27.71	0.759	-	-	26.01	0.727
Patch Averaging [50]	33.35	0.850	28.43	0.765	29.41	0.847	27.67	0.802
Patch Stitching	32.87	0.837	26.71	0.710	29.69	0.849	27.50	0.780
PaDIS (Ours)	33.57	0.854	29.48	0.767	30.80	0.870	29.47	0.846

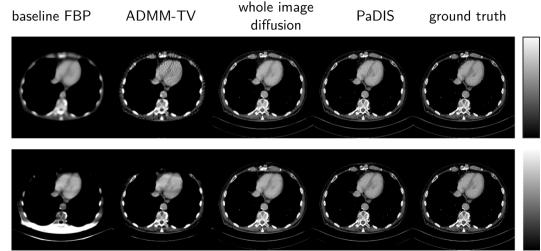
(Averages across all test images.)

# More inverse problem experiments



Method	CT, 60	Views	CT, Far	n Beam	Heavy Deblurring		
Method	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑	
Baseline	25.89	0.746	20.07	0.521	21.14	0.569	
ADMM-TV	30.93	0.833	25.78	0.719	26.03	0.724	
Whole image diffusion	35.83	0.894	26.89	0.835	28.35	0.808	
PaDIS (Ours)	39.28	0.941	29.91	0.932	28.91	0.818	





Top: 60 view CT Bottom: fan-beam CT





#### Patchsize

Ρ	PSNR↑	SSIM↑
8	32.57	0.844
16	32.57	0.829
32	32.72	0.853
56	33.57	0.854
96	33.36	0.854
256	32.84	0.835

### Positional encoding

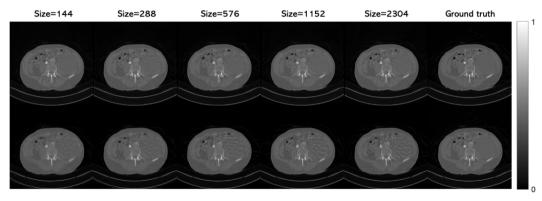
	PSNR↑	SSIM↑
no position enc.	23.25	0.459
no position+init	24.51	0.518
with position enc.	33.57	0.854

# Effect of training dataset size on CT reconstruction



Dataset	Pato	ches	Whole image		
size	56 ×	< <b>56</b>	$256 \times 256$		
	PSNR↑ SSIM↑		PSNR↑	SSIM↑	
144	32.28	0.841	29.12	0.804	
288	32.43	0.837	31.09	0.829	
576	33.03	0.846	31.81	0.835	
1152	33.01	0.849	31.36	0.834	
2304	33.57	0.854	32.84	0.835	





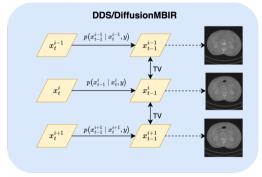
Top: PaDIS

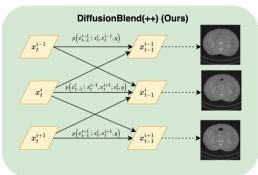
Bottom: whole image diffusion model

Challenge 2: Data dimensions & scaling to 3D (and 4D)



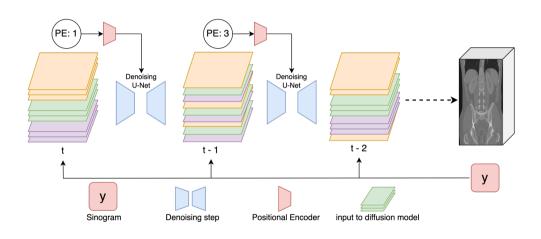
### arXiv 2406.10211 (NeurIPS 2024) [66]





## DiffusionBlend models groups of slices





## 3D prior models for X-ray CT



J. Fessler

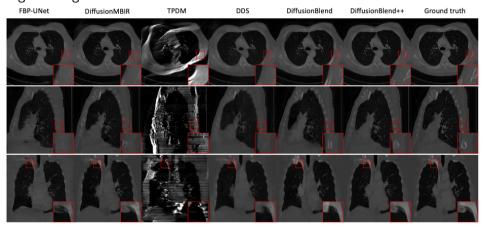
Score

Method	Distribution Model
DiffusionMBIR (2D) [32]	$\frac{1}{Z} \prod_{i=1}^{H} p(\mathbf{x}[:,:,i])$
TPDM (⊥ 2D) [33]	$rac{1}{Z} \left(\prod_{i=1}^{N} q_{ heta}({m{x}}[:,:,i])^{lpha} ight) \left(\prod_{j=1}^{N} q_{\phi}({m{x}}[j,:,:])^{eta} ight)$
DiffusionBlend	$\frac{1}{Z} \prod_{i=1}^{H} p(\mathbf{x}[:,:,i]   \mathbf{x}[:,:,i-j:i-1], \mathbf{x}[:,:,i+1:i+j])$
${\sf DiffusionBlend}{+}{+}$	$\frac{1}{Z}\prod_{i=1}^{r}p(\mathbf{x}[:,:,\mathcal{S}_{i}])$

### DiffusionBlend for 3D limited-angle CT



90° angular range



Improved quality both qualitatively and quantitively with strong prior.

# 3D limited-angle CT results



AAPM Datas							LIDC Dataset					
Method	hod Axial		Sagittal		Coronal		Axial		Sagittal		Coronal	
	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑
FBP	16.36	0.643	16.36	0.524	15.62	0.531	18.79	0.672	19.84	0.675	20.01	0.676
FBP-UNet	27.38	0.910	27.81	0.918	28.44	0.930	29.42	0.885	29.50	0.884	29.54	0.887
DiffusionMBIR	25.98	0.872	27.14	0.877	27.74	0.903	30.52	0.906	30.57	0.906	30.68	0.907
TPDM	-	-	-	-	-	-	14.44	0.141	14.06	0.141	14.54	0.313
DDS 2D	28.05	0.916	27.99	0.916	28.82	0.922	27.92	0.843	27.89	0.835	27.96	0.842
DDS	28.20	0.918	28.17	0.926	29.03	0.934	28.12	0.865	28.06	0.869	28.13	0.879
DiffusionBlend (Ours)	35.38	0.971	35.85	0.972	37.62	0.972	30.43	0.917	31.24	0.920	31.02	0.924
DiffusionBlend++ (Ours)	35.86	0.975	36.03	0.976	<u>37.45</u>	0.976	34.33	0.957	34.48	0.957	34.64	0.956
[66 Table 4]							•					

[66, Table 4]





Test-time latent x far from training distribution:

$$\mathbf{y} = \mathbf{A}\mathbf{x} + \mathbf{\varepsilon}, \quad \mathbf{x} \sim \tilde{p}(\cdot) \neq p(\cdot)$$

Non-Bayes approach Abandon training via self-supervision, e.g., deep image prior (DIP) [67]:

$$\hat{\mathbf{x}} = f_{\hat{\boldsymbol{\theta}}}(\mathbf{z}), \qquad \hat{\boldsymbol{\theta}} = \arg\min_{\mathbf{z}} \|\mathbf{y} - \mathbf{A}f_{\boldsymbol{\theta}}(\mathbf{z})\|_2^2, \qquad \mathbf{z} \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$$

Neural network  $f_{\theta}(\cdot)$  acts as implicit regularizer.

DIP is prone to overfitting of noisy measurements [67]; remedies such as early stopping, regularization, network initialization [68–70].

# Distribution shifts & test-time adaptation



➤ Self-supervised (whole-image) diffusion models [71, 72] "Deep diffusion image prior" (DDIP) or "steerable conditional diffusion:"

$$L(\boldsymbol{\theta}) = \|\boldsymbol{y} - \boldsymbol{A} \operatorname{CG}(\hat{\boldsymbol{x}}_{0|t}(\boldsymbol{x}_t; \boldsymbol{\theta}))\|_2^2$$

$$\operatorname{CG}(\hat{\boldsymbol{x}}_{0|t}) \triangleq \arg\min_{\boldsymbol{x}} \frac{\gamma}{2} \|\boldsymbol{y} - \boldsymbol{A}\boldsymbol{x}\|_2^2 + \frac{1}{2} \|\boldsymbol{x} - \hat{\boldsymbol{x}}_{0|t}\|_2^2$$

Conjugate gradient (CG) descent is used to enforce data fidelity. Still requires early stopping to avoid over-fitting.



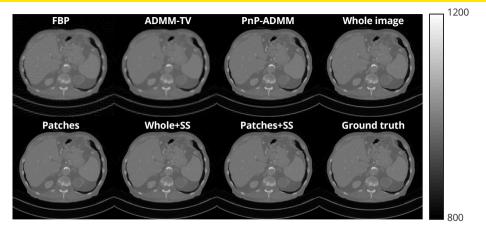
▶ Patch-based test-time adaptation [73, 74] arXiv 2410.11730 (IEEE T-CI, in-press) Test-time loss for diffusion model adaptation:

$$L(\theta) = \left\| \mathbf{y} - \mathbf{A} \sum_{c} \mathbf{G}_{c}' D_{\theta}(\mathbf{G}_{c} \mathbf{x}_{t}, c | \mathbf{y}) \right\|_{2}^{2}$$

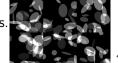
Patch-based denoiser for diffusion model

$$D_{\theta}(\mathbf{x}) = \sum_{c} \mathbf{G}'_{c} D_{\theta}(\mathbf{G}_{c}\mathbf{x}, c),$$





No in-distribution training data. Pre-trained with random ellipses. Results of 60-view CT reconstruction using self supervised (SS) loss.



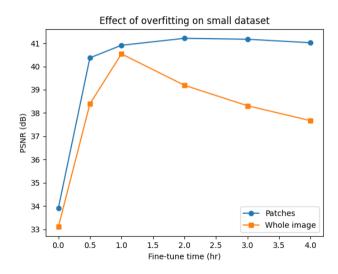
# Patch-based test-time adaptation III

Method	CT, 20 Views		CT, 60 Views		Deblurring		Superresolution	
	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑
Baseline	24.93	0.613	30.15	0.784	23.93	0.666	25.42	0.724
ADMM-TV	26.81	0.750	31.14	0.862	27.58	0.773	25.22	0.729
PnP-ADMM [63]	30.20	0.838	36.75	0.932	28.98	0.815	27.29	0.796
PnP-RED [64]	27.12	0.682	32.68	0.876	28.37	0.793	27.73	0.809
Whole image	28.11	0.800	33.10	0.911	25.85	0.742	25.65	0.742
Patches [60]	27.44	0.719	33.97	0.934	26.77	0.782	26.12	0.759
Whole+SS [72]	33.19	0.861	40.47	0.957	29.50	0.831	27.07	0.701
Patches+SS (Ours)	33.77	0.874	41.45	0.969	30.34	0.860	28.10	0.827

<sup>&</sup>quot;SS" = self-supervision, aka test-time adaptation

## Patch-based test-time adaptation IV

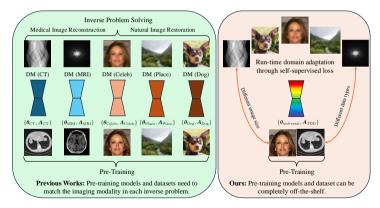




### Towards a "universal" diffusion model

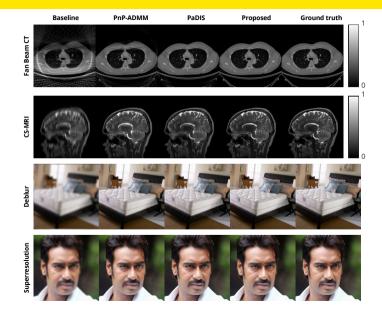


Extension to cases where # of channels at test time differs from training data, e.g., MR reconstruction (real/imag) from patch-based diffusion model pre-trained on color (RGB) natural images and grayscale CT images [75]









### SPAR results



Comparison of quantitative results on four different medical imaging inverse problems.

Method	PBCT, 60 Views		FBCT, 40 Views		$512 \times 512$ <b>CT</b>		CS-MRI, $7 \times$	
ivietnoa	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑	PSNR↑	SSIM↑
Baseline	30.15	0.784	17.86	0.381	28.33	0.700	33.94	0.894
ADMM-TV	31.14	0.862	24.20	0.628	29.36	0.788	36.74	0.924
PnP-ADMM [63]	36.75	0.932	28.86	0.747	37.48	0.910	35.77	0.907
PaDIS+FC [60]	39.16	0.942	27.91	0.796	33.11	0.831	35.17	0.904
SCD [72]	41.16	0.962	21.28	0.463	-	_	-	_
Ours (SPAR)	42.72	0.972	36.11	0.918	38.81	0.929	39.15	0.949
Ideal*	42.82	0.973	36.34	0.923	38.94	0.930	39.42	0.953

<sup>\*</sup>not available in practice with a single diffusion model

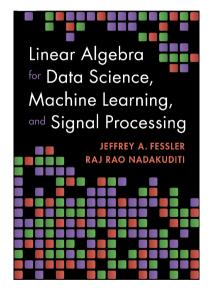
# Summary / future directions



- Challenges
  - Dearth of data
  - Dimensionality
  - Distribution shifts
- Promise
  - Generative models are promising for under-determined inverse problems
  - Learning patch score models is feasible with denoising score matching
  - For limited training data, patch-models can outperform whole-image models
- Future steps
  - ▶ Integrate invariances: amplitude scale / rotation / flip / DC offset ...
  - Explore trade-offs between generalizability and in-distribution performance
  - Extend to 3D, 3D+Time, 3D+Multicontrast

Tutorial Julia code: https://github.com/JeffFessler/ScoreMatching.jl





- Online demos: https://github.com/JeffFessler/ book-la-demo
- Topics include: low-rank matrix approximation, robust PCA, photometric stereo, video foreground/background separation, spectral clustering, matrix completion, ...
- Cambridge Univ. Press, 2024

### Resources



Talk and code available online at http://web.eecs.umich.edu/~fessler





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