

Advancing X-ray Tomography using Deep Generative Adversarial Networks (TomoGAN)

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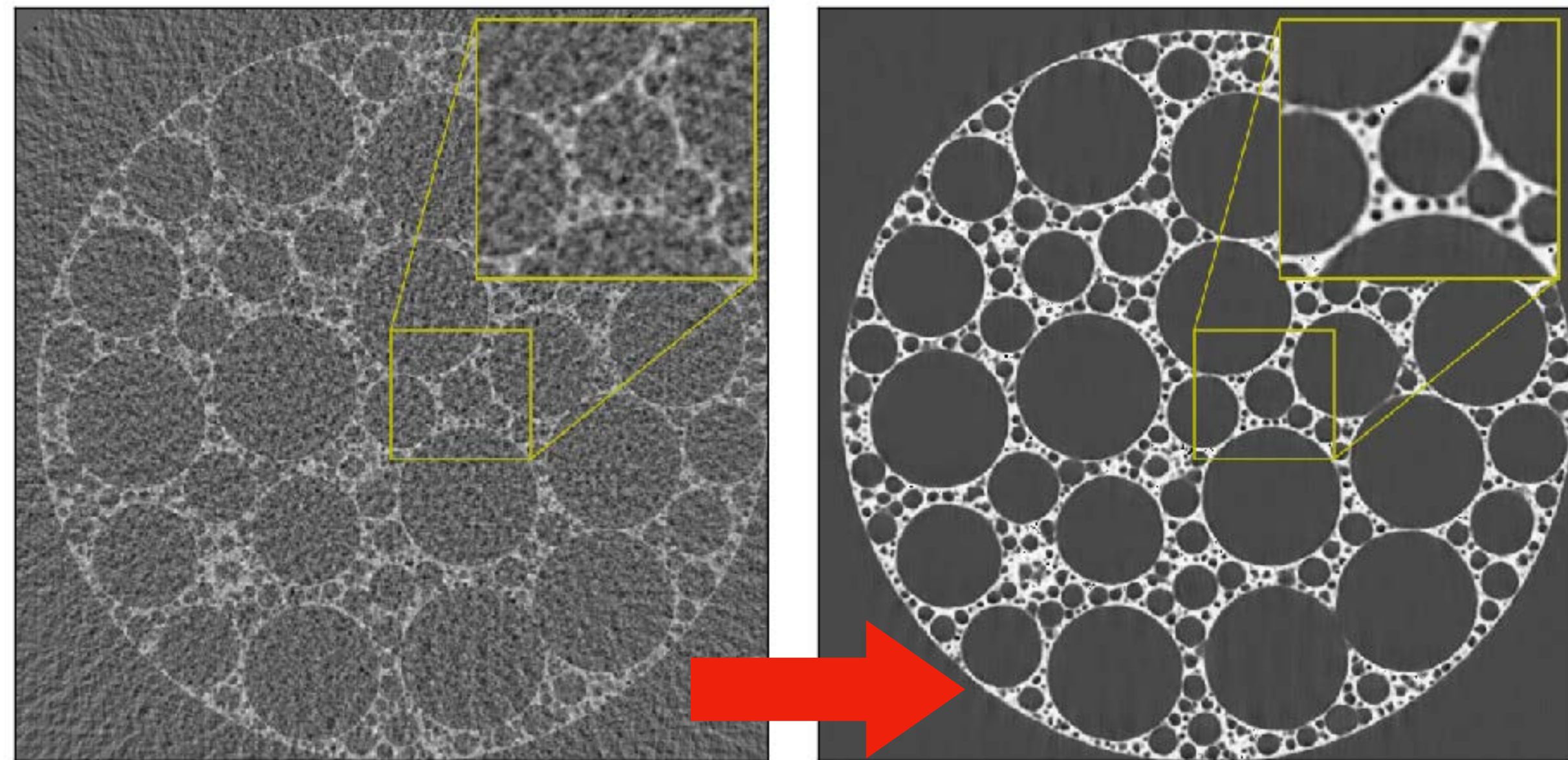
Francesco De Carlo
XSD



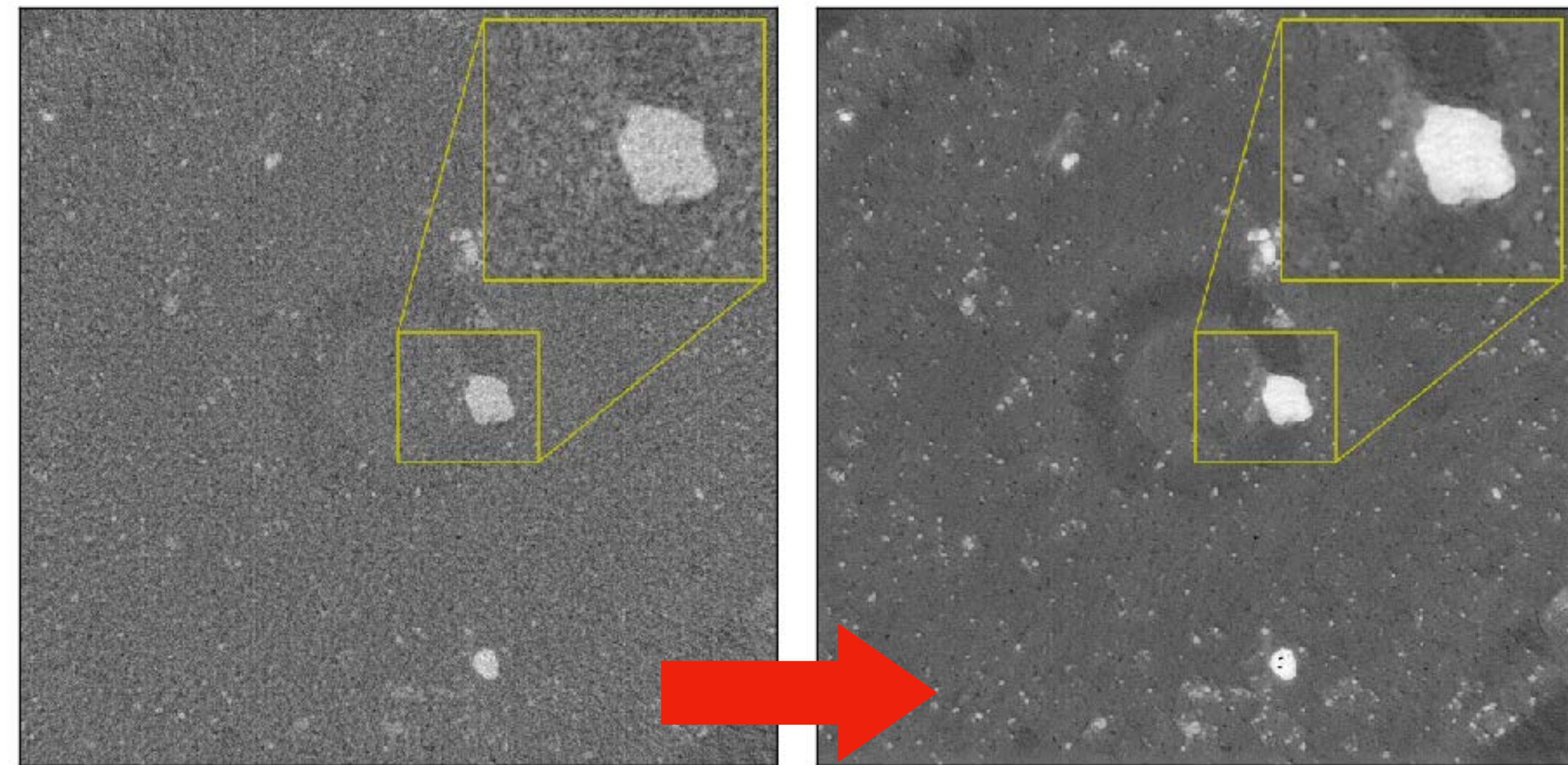
Ian Foster
DSL

Motivation

- (1) lower X-ray dosage for sensitive sample like bio-sample;
- (2) faster experiment to capture dynamic features, like in fast chemical reaction processes;
- (3) smaller dataset and less computation for [near] realtime tomography imaging.



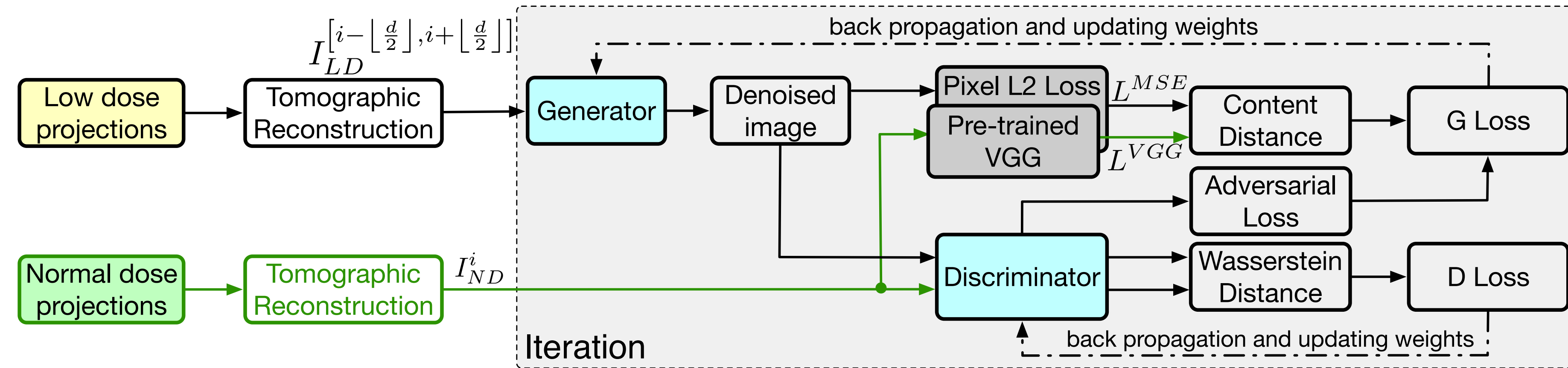
On the left, the results of conventional reconstruction, which are highly noisy. On the right, those same results after denoising with TomoGAN.



Model is trained with one shale sample imaged at APS and tested with **another** shale sample imaged at Swiss Light Source (SLS).

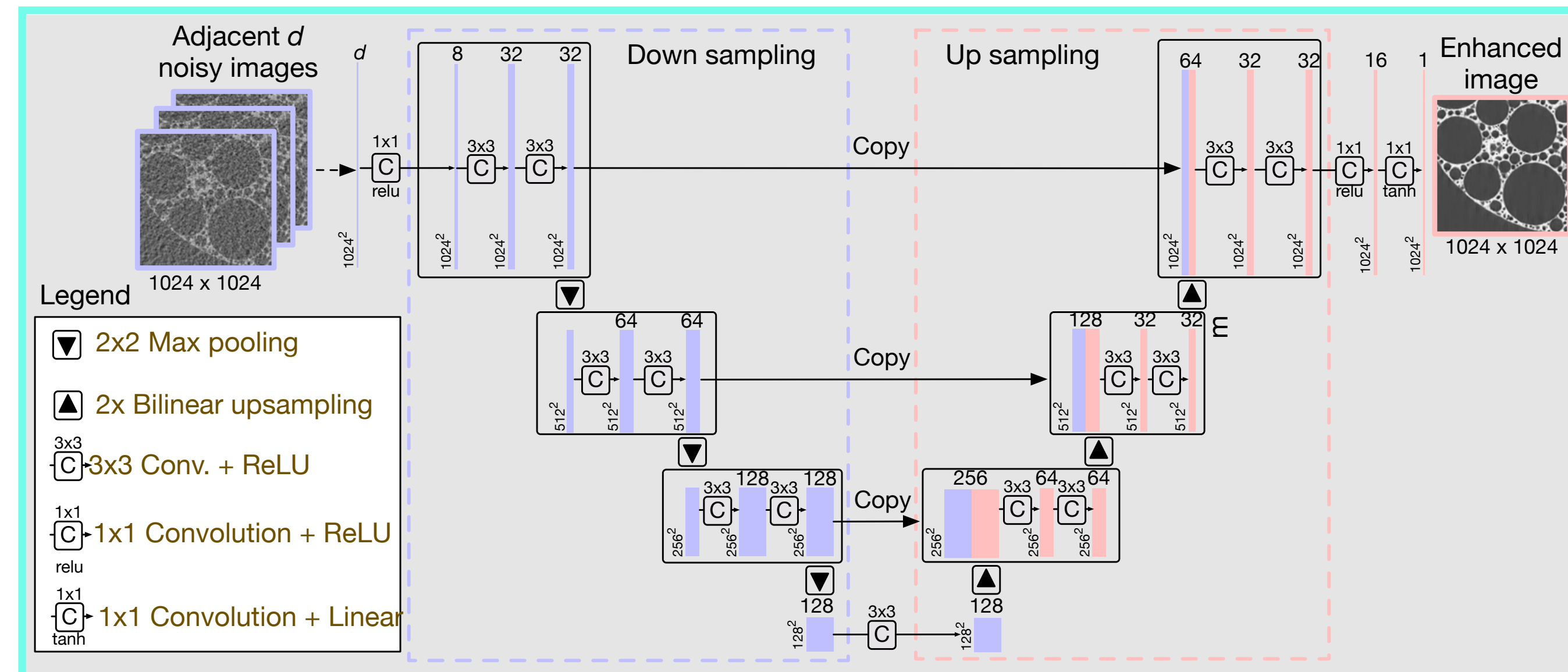
Method

A generative adversarial network (GAN) is a class of machine learning systems in which two neural networks, generator (G) and discriminator (D), contest with each other in a game (in the sense of game theory, often but not always in the form of a zero-sum game).



In our model, the discriminator's job remains unchanged, but the generator is tasked not only with fooling (indistinguishable) the discriminator but also with being near the ground truth output in an L2 sense.

The discriminator works as a helper to train the generator that we need to denoise images.



Our Generator Architecture

Training

Discriminator Wasserstein GAN [1] + gradient penalty [2]

$$L(\theta_D) = \frac{1}{m} \sum_{i=1}^m \left[D(G(I_{LD}^i)) - D(I_{ND}^i) \right] + \lambda_D \frac{1}{m} \sum_{i=1}^m \left[\left(\left\| \nabla_{\bar{I}} D(\bar{I}^i) \right\|_2 - 1 \right)^2 \right],$$

Generator Weighted average of Adversarial loss, Perceptual loss, and Pixel-wise MSE

$$\ell^G = \lambda_g \ell_{adv} + \lambda_p \ell_{mse} + \lambda_v \ell_{vgg}$$

$$\ell_{adv}(\theta_G) = -\frac{1}{m} \sum_{i=1}^m D(G(I_{LD}^i))$$

$$\ell_{vgg} = \sum_{i=1}^{W_f} \sum_{j=1}^{H_f} \left(V_{\theta_{vgg}}(I^{ND})_{i,j} - V_{\theta_{vgg}}(G_{\theta_G}(I^{LD}))_{i,j} \right)^2$$

$$\ell_{mse} = \sum_{c=1}^W \sum_{r=1}^H \left(I_{c,r}^{ND} - G_{\theta_G}(I^{LD})_{c,r} \right)^2$$

[1] Wasserstein GAN. M. Arjovsky, S. Chintala, L. Bottou. arXiv:1701.07875

[2] Improved Training of Wasserstein GANs. I. Gulrajani, F. Ahmed, M. Arjovsky, V. Dumoulin, A. Courville. arXiv:1704.00028

Experiments

Datasets

- **Three foam simulation datasets, each with 1024 slices**
- **Two shale samples imaged at both APS and SLS, totals four datasets and each with 2048 slices.**

Label	projection	reconstruction	Facility	Sample	Scan	Axis
tomo_00001	(1501, 1792, 2048)	(1792, 2048, 2048)	APS	B1	hornby	1024
tomo_00002	(1501, 1792, 2048)	(1792, 2048, 2048)	APS	N1	blakely	1029
tomo_00003	(1441, 2048, 2048)	(2048, 2048, 2048)	SLS	B1	hornby	1011
tomo_00004	(1441, 2048, 2048)	(2048, 2048, 2048)	SLS	N1	blakely	1048

Low dose cases

- **Sparse views**

Subsample the original, (i.e., normal dose) projections to 1/2, 1/4, 1/8 and 1/16 for experiments and model evaluation.

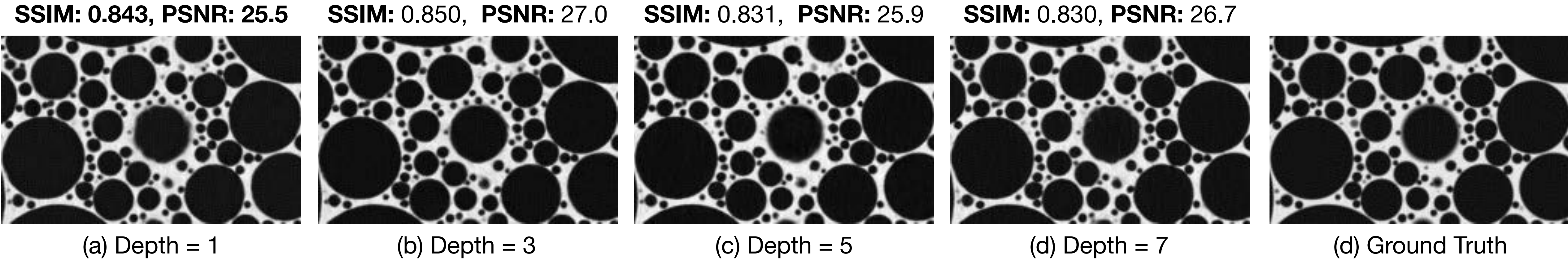
- **Short exposure time.**

For simulation datasets, we simulate x-ray projections with different photon intensities to simulate different exposure times

For experimental shale datasets, we used added noise using a Poisson distribution to simulate different exposure times.

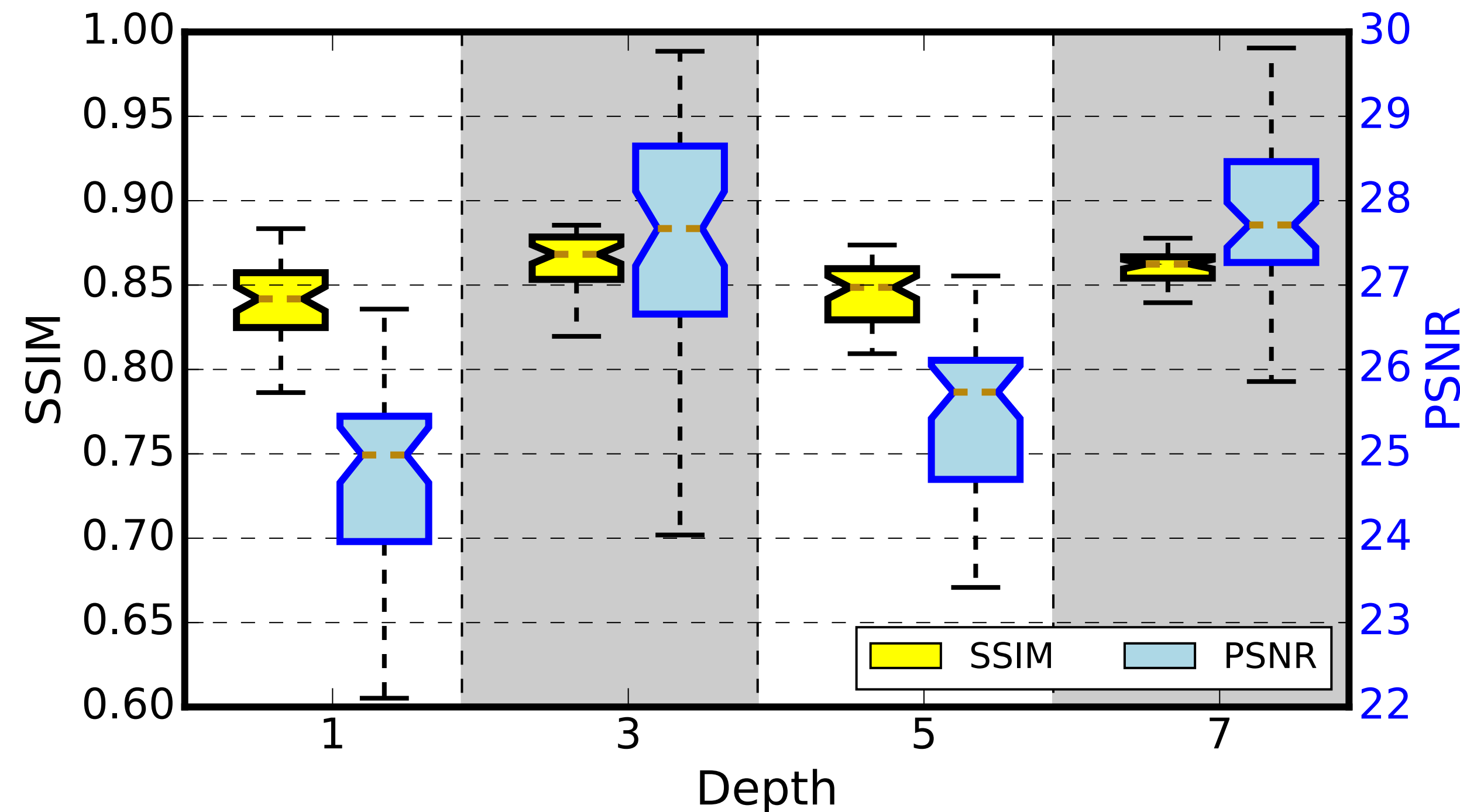
Results - Adjacent slices

Effectiveness of using adjacent slices in image enhancement



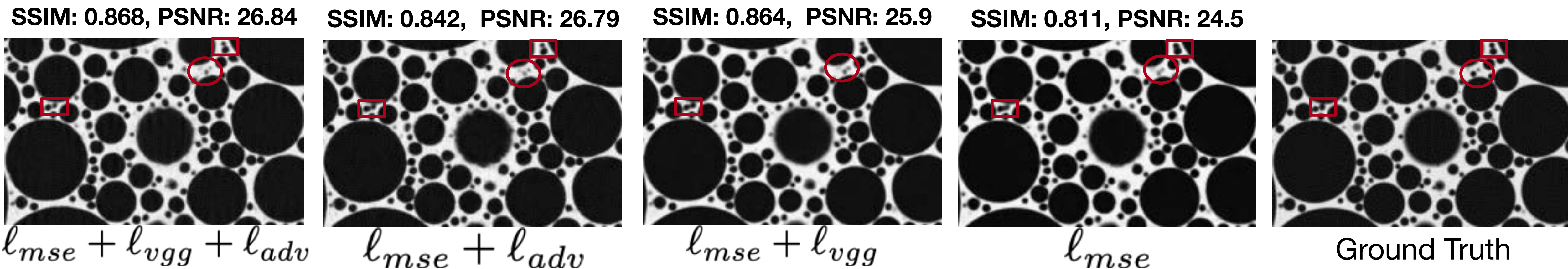
The input depth d has big influence on mode performance, and that $d=3$ gets the best quality, especially when the original feature edge is not sharp (e.g., the center circle).

We note that the best depth d depends on dataset characteristics such as feature resolution. $d=3$ may not be the best for other datasets where feature sizes change slowly across slices.



Results - Loss

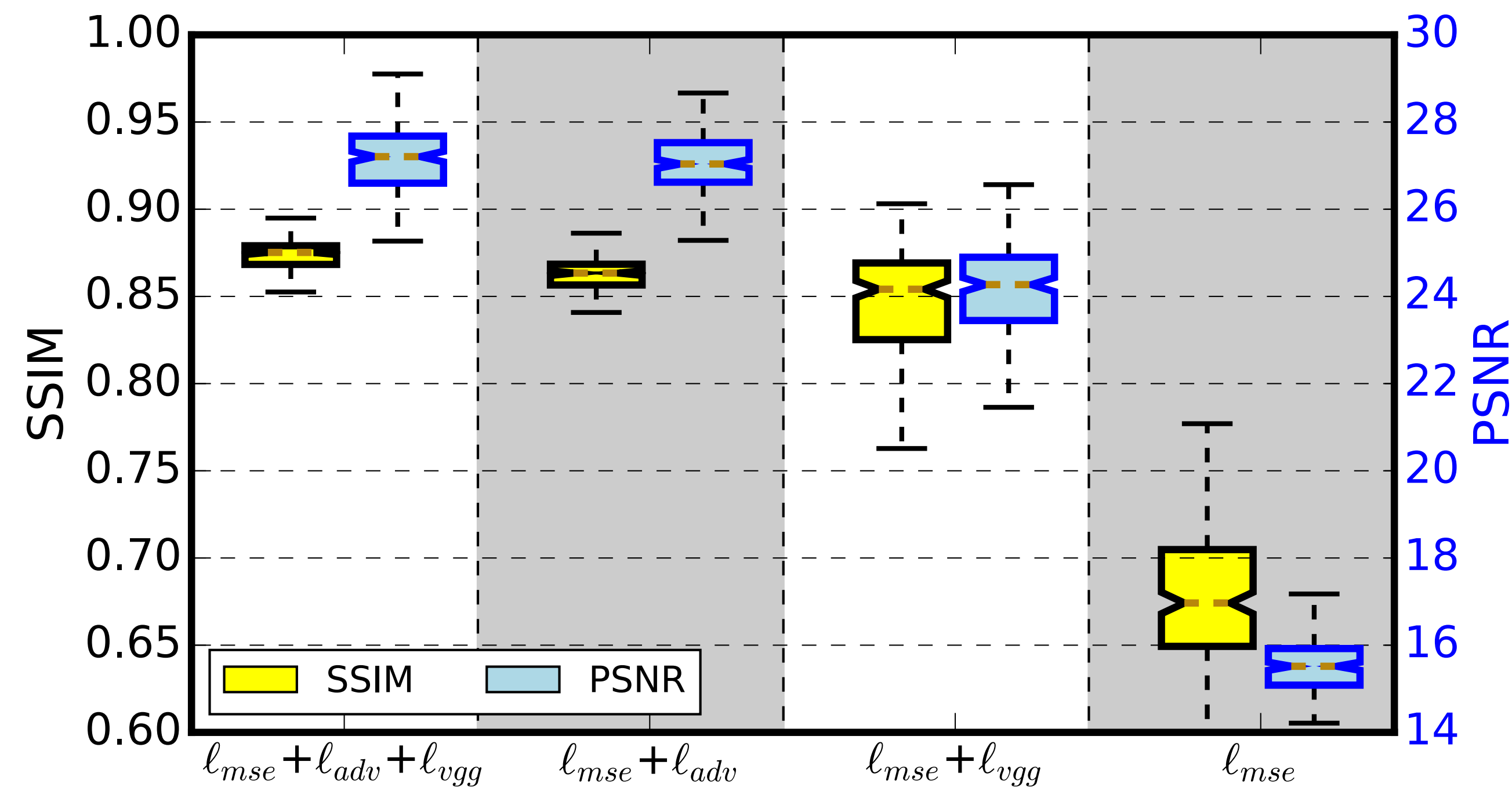
Importance of the various loss terms



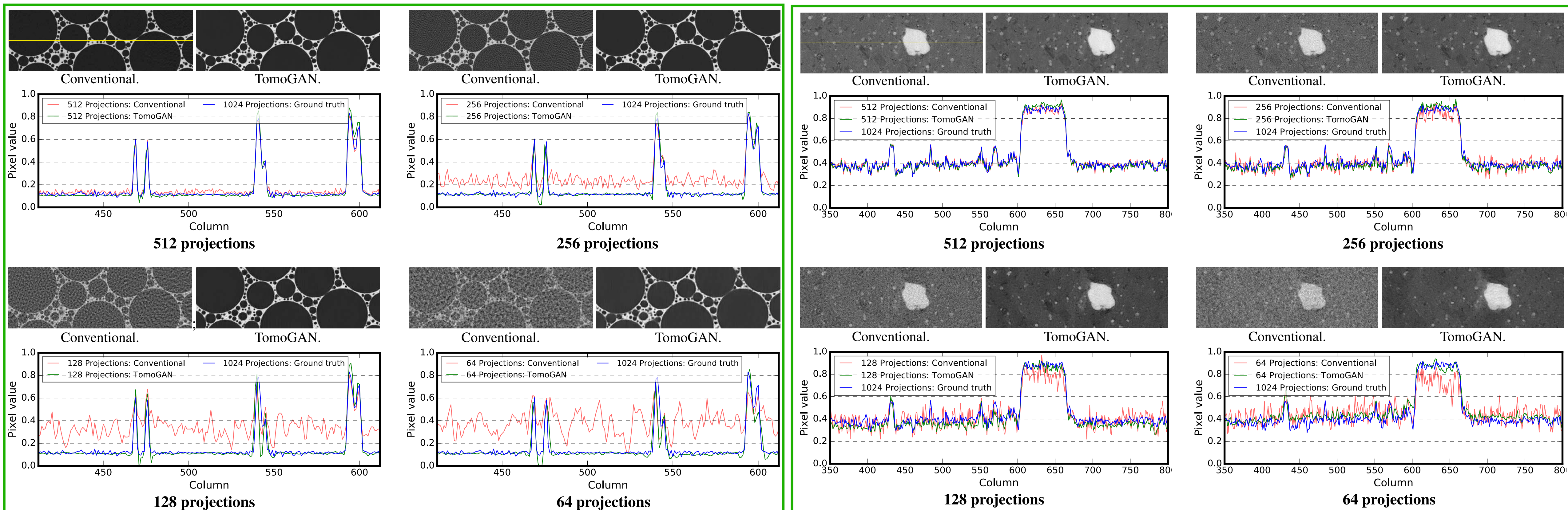
MSE is necessary to enforce correctness of low-frequency structures but MSE alone is not enough.

The adversarial and perceptual loss terms each provide considerable improvements when used in isolation.

The two together are only slightly better than adversarial loss alone.

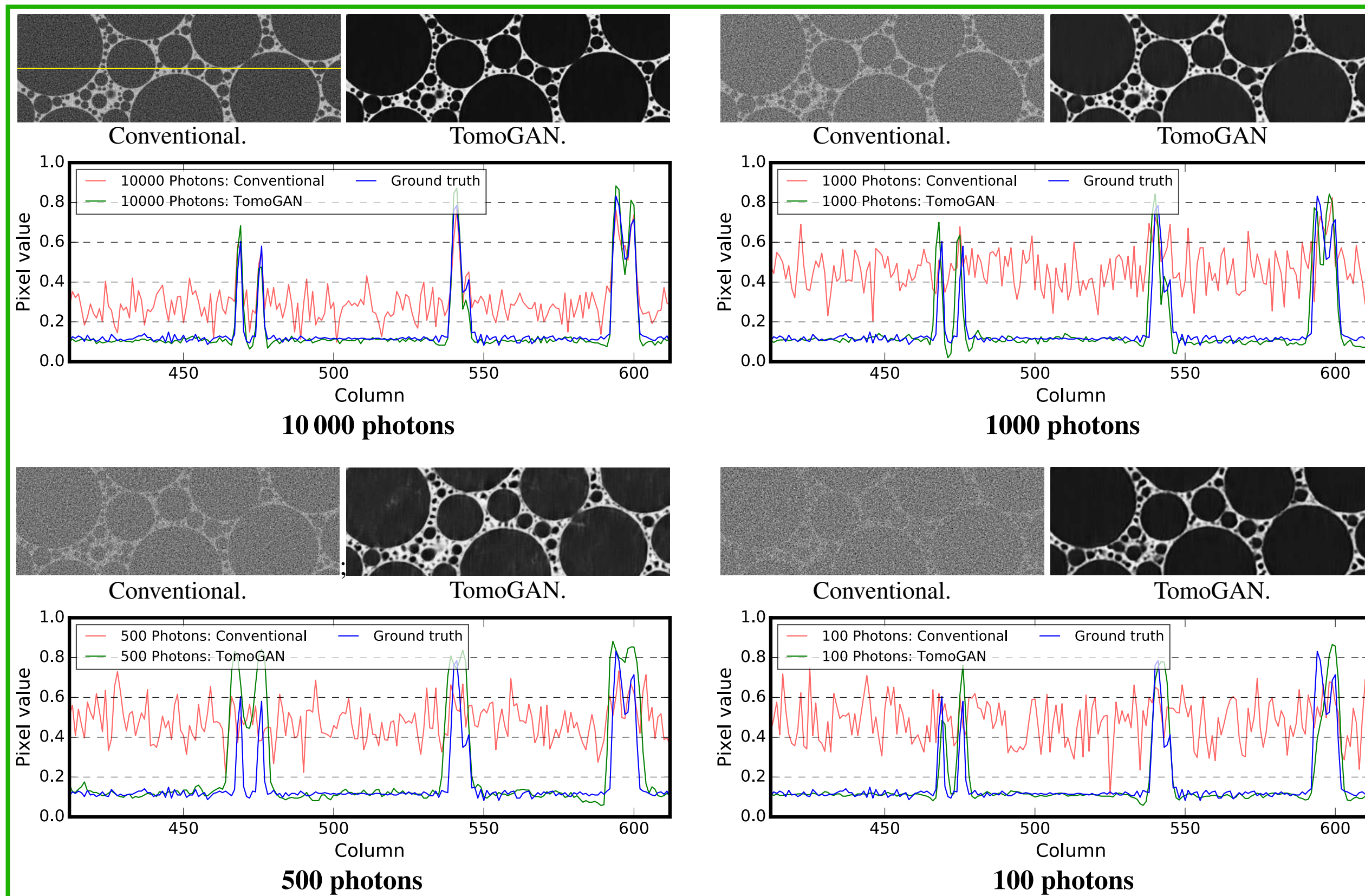


Results - Sparse views

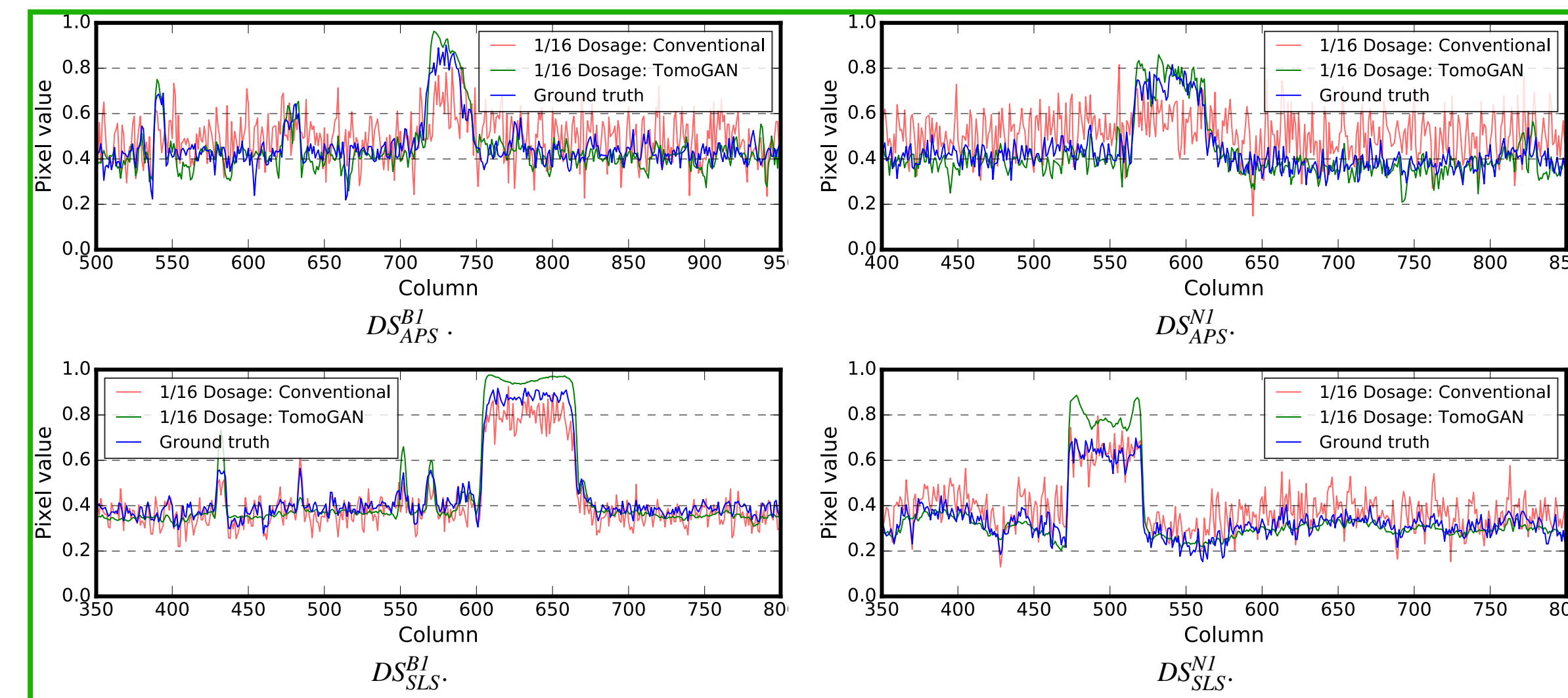


Conventional vs. TomoGAN-enhanced reconstructions of simulated (left) data and shale (right), subsampled to (512, 256, 128, 64) projections. In each group of three elements, the two images show conventional and TomoGAN reconstructions, while the plot shows conventional, TomoGAN, and ground truth values for the 200 pixels on the horizontal line in the top left image.

Results - Short exposure time



Pixel values of an arbitrarily chosen feature in each of the four experimental datasets, with projections generated by using 1/16 of the normal exposure time. Feature shapes are different for each dataset.

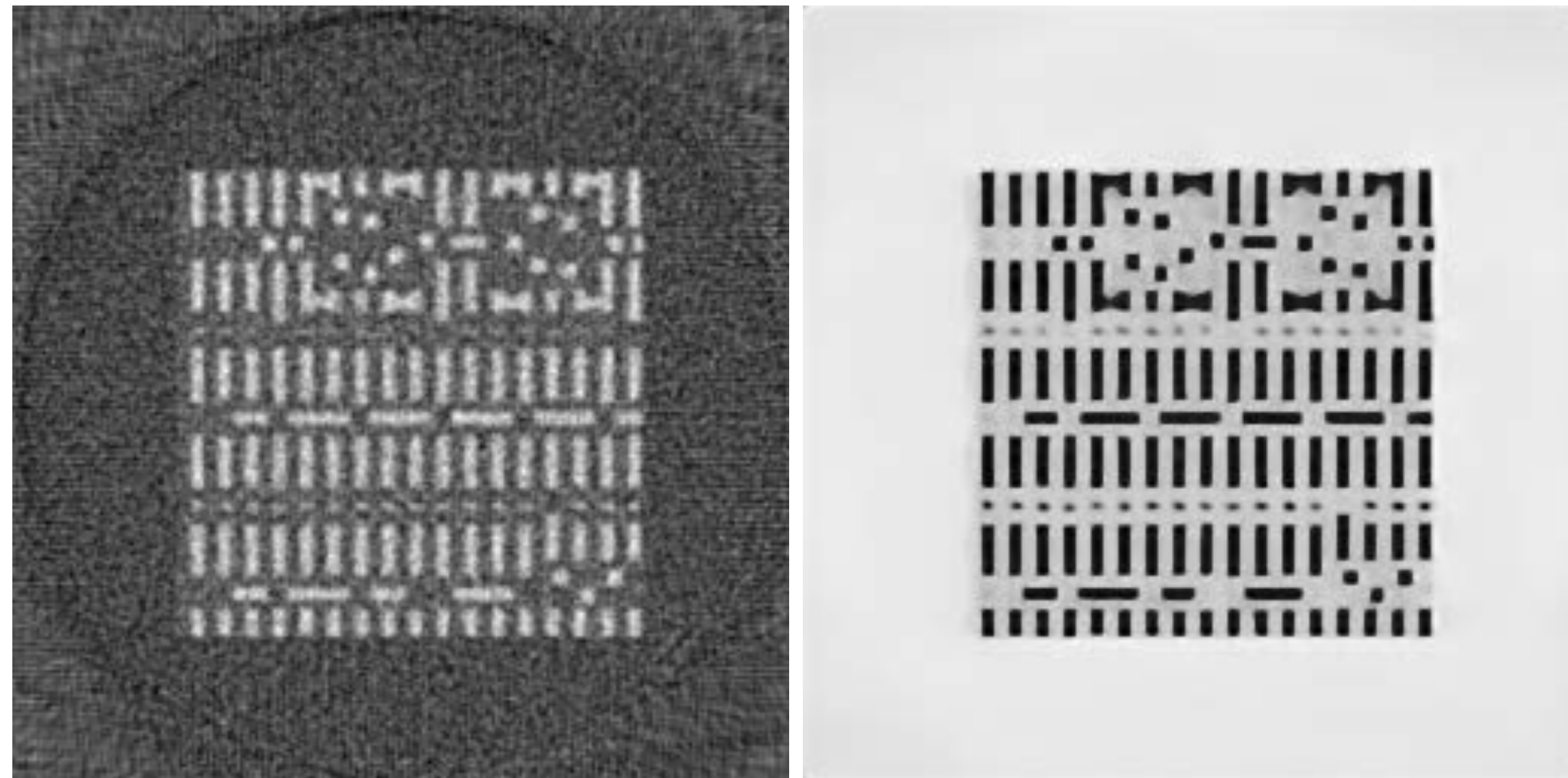


Conventional vs. TomoGAN-enhanced reconstructions of simulated data with intensity limited to 10000, 1000, 500, 100 photons per pixel.

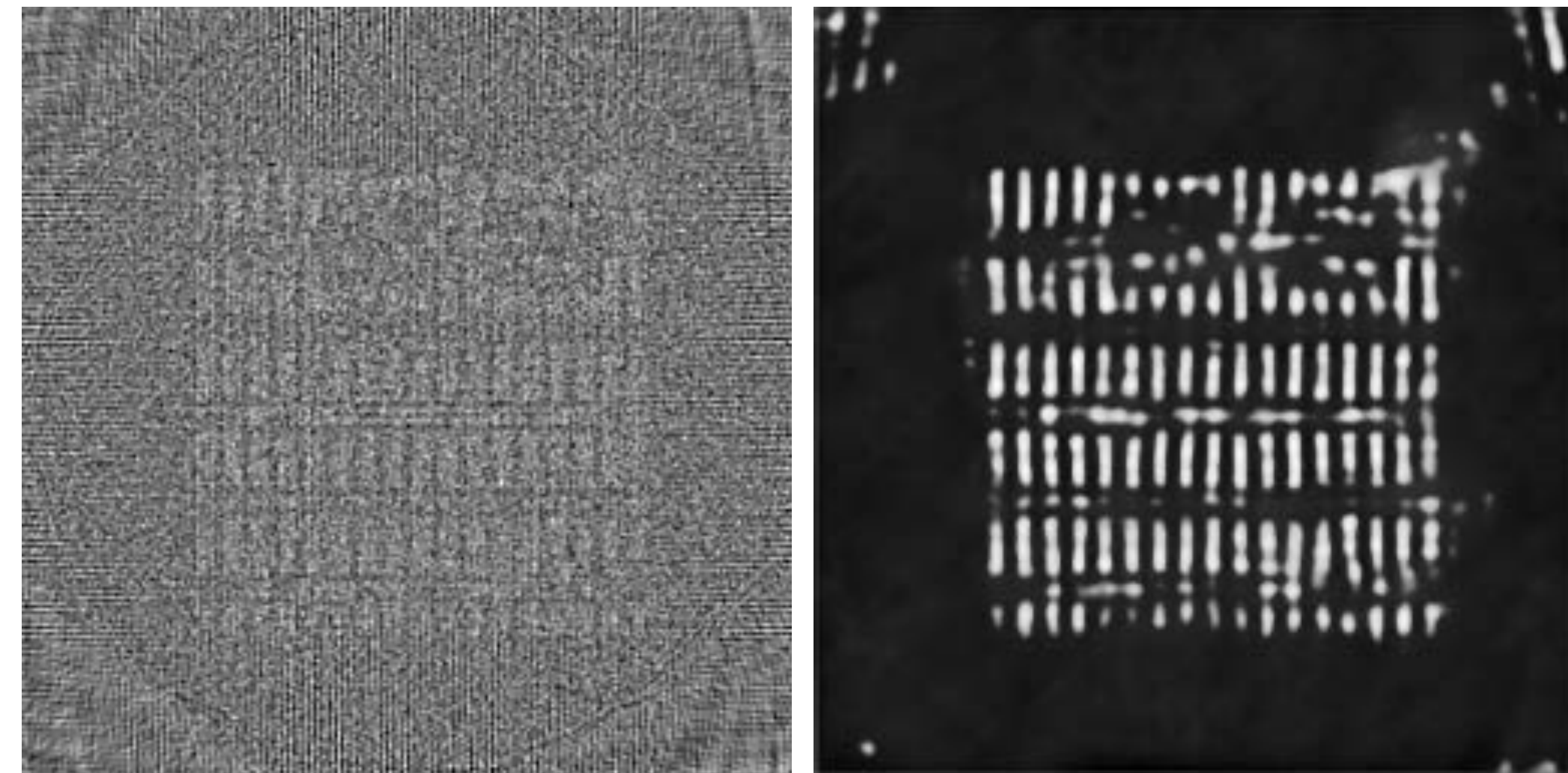
TomoGAN - Extend use case

It has been applied to the joint ptycho-tomography problem for reconstructing the complex refractive index of a 3D object.

- There is a ptychography process to reconstruct projections needed for tomography. but it is very time consuming to image the sample (month).
- Less datapoint results in noisier ptychography reconstruction and worse tomography images.
- TomoGAN here was used to enhance tomography images with less data points need to collect, i.e., faster experiment.



Delta, 0.003



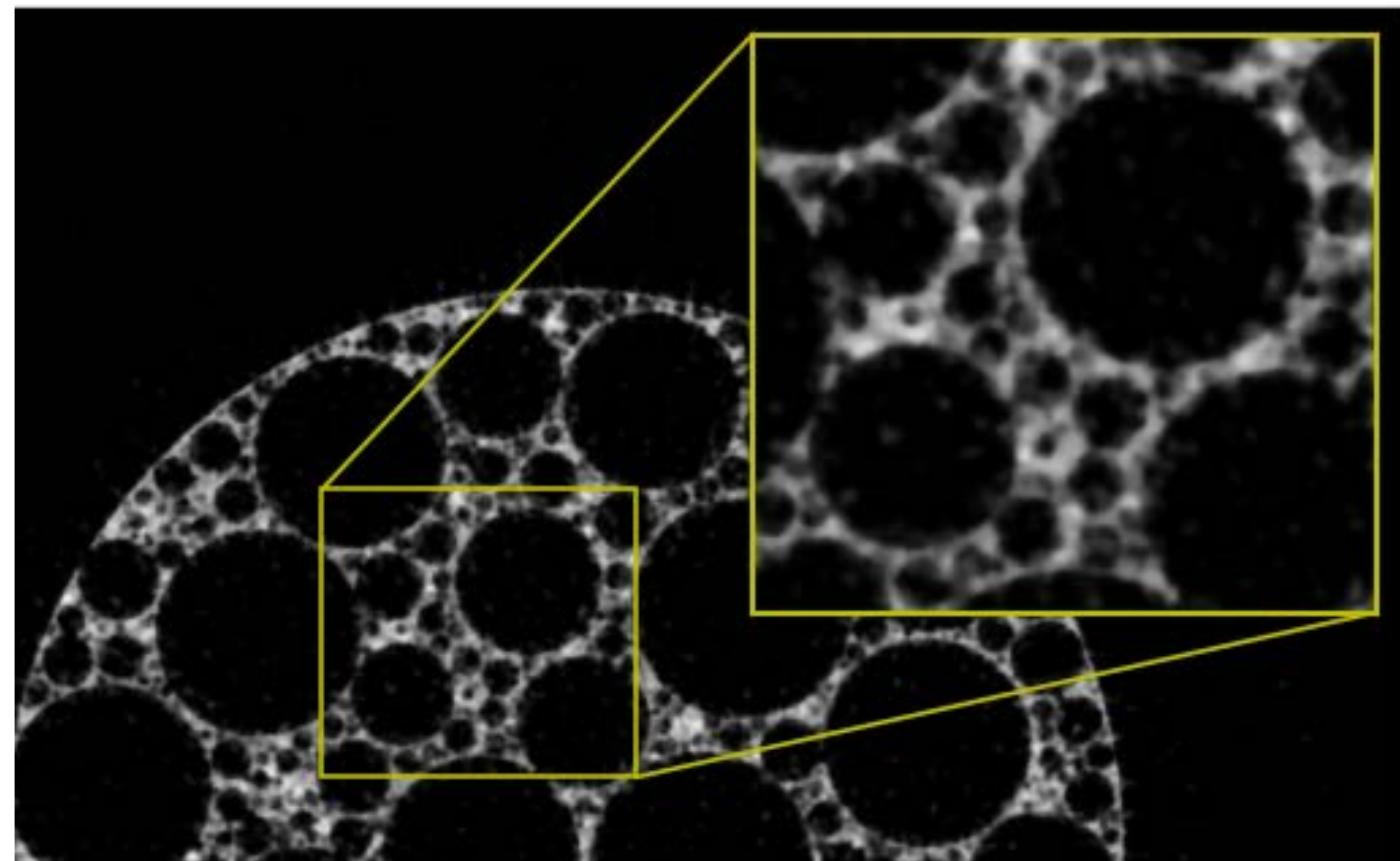
Beta, 0.03

Computational superiority

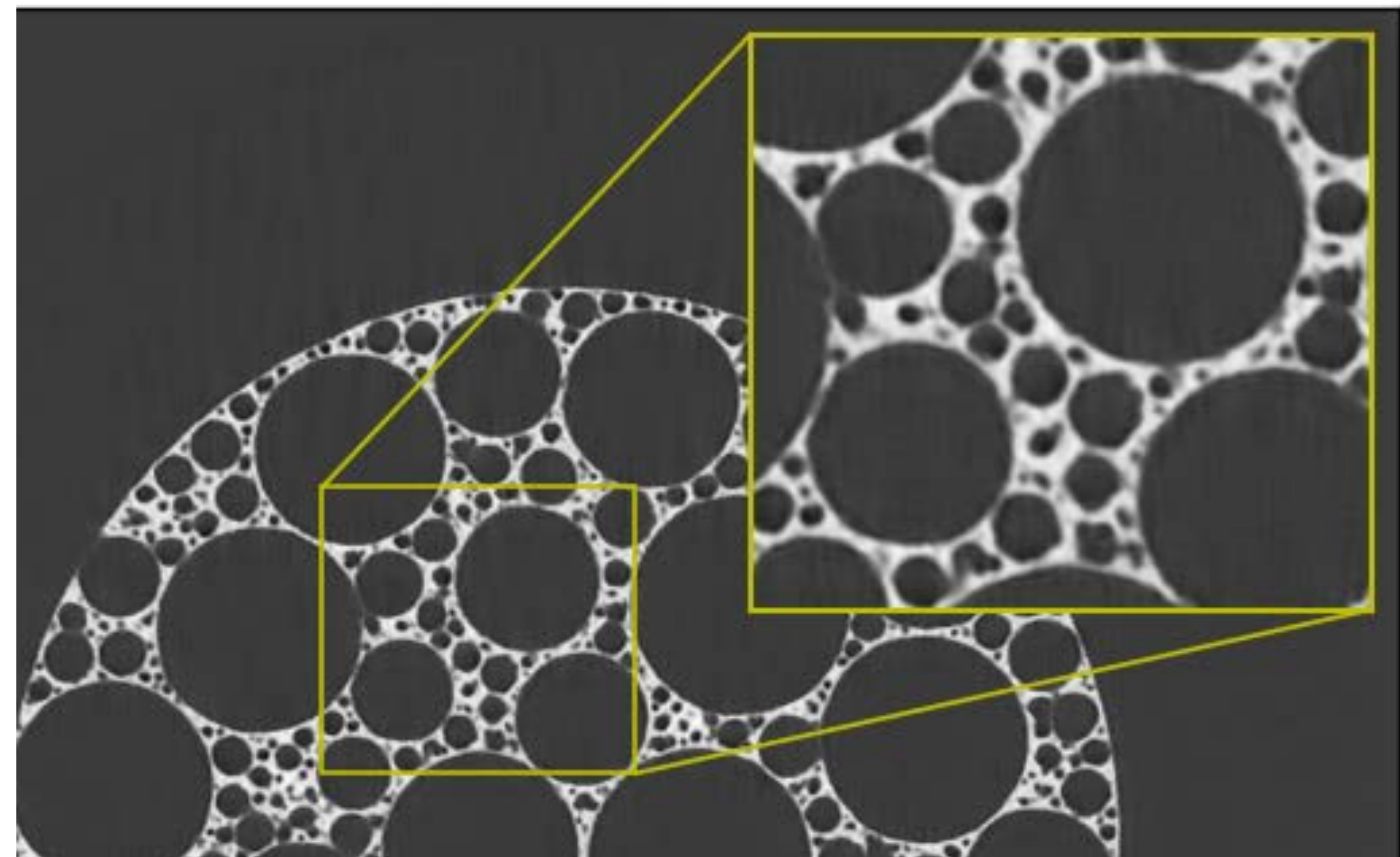
The filtered back projection (FBP) algorithm takes 40 ms to reconstruct one image (using TomoPy) and TomoGAN takes 30 ms to enhance the reconstruction, totals **70 ms** per image.

In contrast, the SIRT based solution (using TomoPy) takes **550 ms** (400 iterations), i.e., 8x faster. Times are measured using one Tesla V100 graphic card.

Moreover, iterative reconstruction does not provide better image quality than does our method.



SIRT + total variation postprocess.



Filtered back projection + TomoGAN post-process.

Make it usable

Hack and Play

open source implementation, better to have a GPU for training

```
python ./train.py -ld low-dose-img.hdf5 -nd normal-dose-img.hdf5  
python ./infer.py -ld ld-prod.hdf5
```

X as a Service

DLHub

Data and Learning Hub for Science



B. Blaiszik. arXiv:1811.11213

```
from dlhub_sdk.client import DLHubClient  
dlhub = DLHubClient()  
  
model = dlhub.get_id_by_name("tomoGAN")  
data = h5py.File("tomo_ld.hdf5", "r")["ld_img"]  
pred = dl.run(model, data)
```

Plug and Play



Thanks!