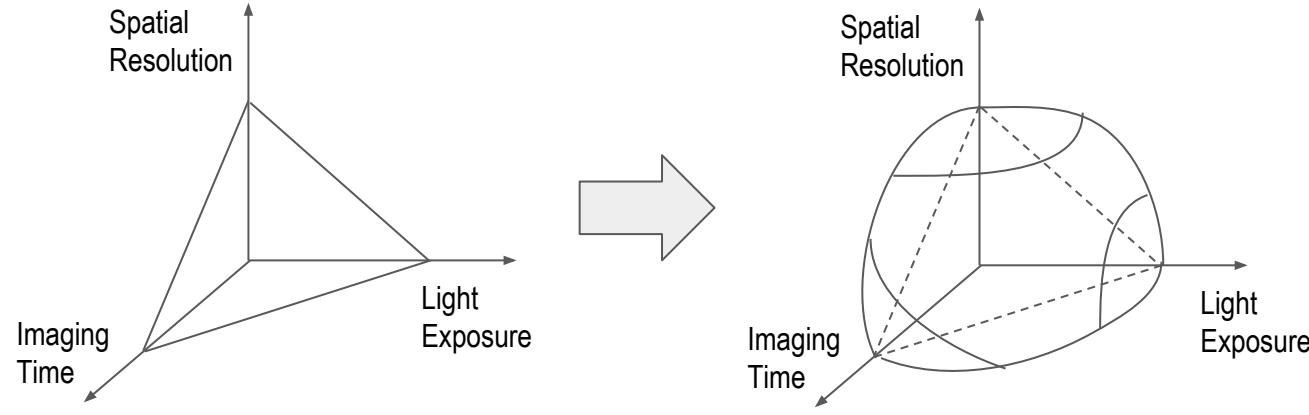


Introduction

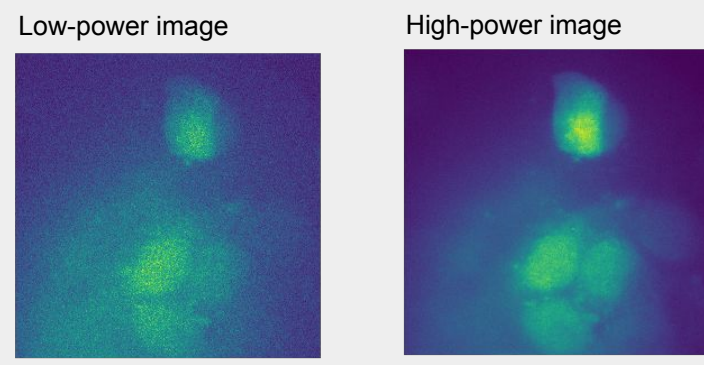
Problem statement: Advanced fluorescence microscopes use high-powered LEDs and lasers to observe cells over time. However, observational cells will eventually degrade under extended light exposure resulting in unusable, damaged cells.

Goal: Expand the tradeoff boundary between spatial resolution and exposure time using techniques from Deep Learning.



- Similar to super-resolution and image recovery tasks.
- Expand the possibilities frontier.

Motivation: Study RNA distribution and gene expression at lower power channels.



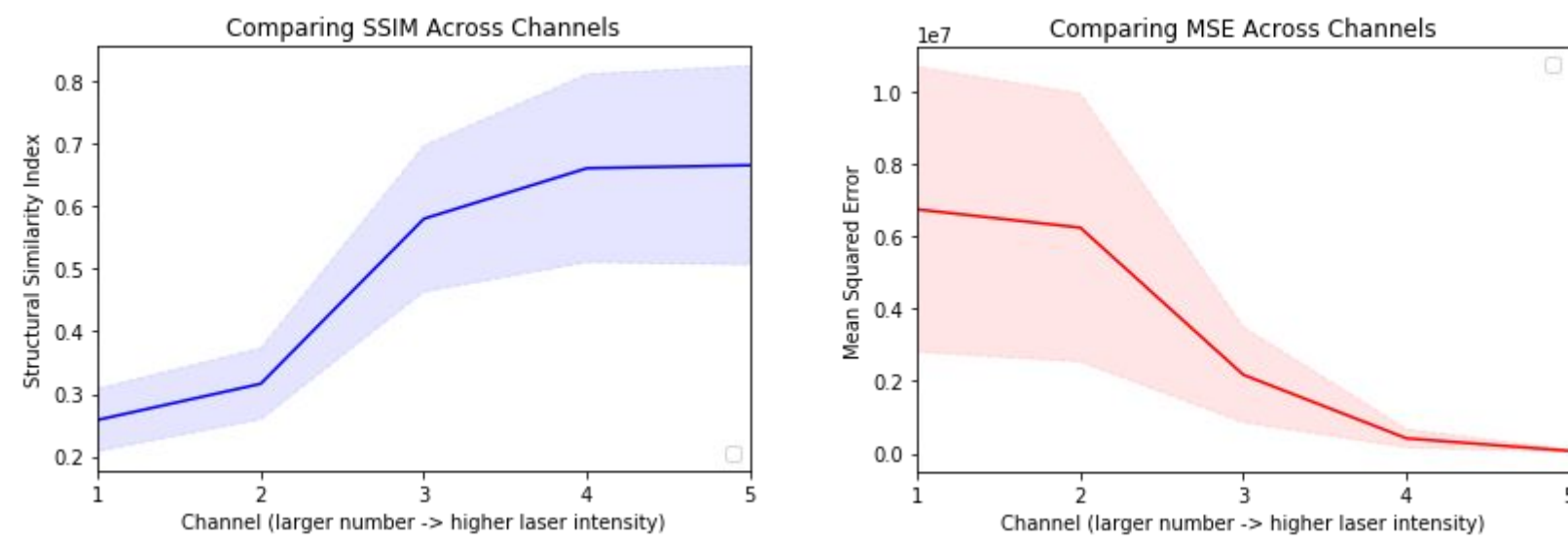
Data Overview

Data Overview: Stacks of 512 x 512 images taken by a fluorescence microscope across 7 different channels at each focal distance.

Data collected by an automated process using a fluorescence microscope on cells stained with DAPI and GFP:

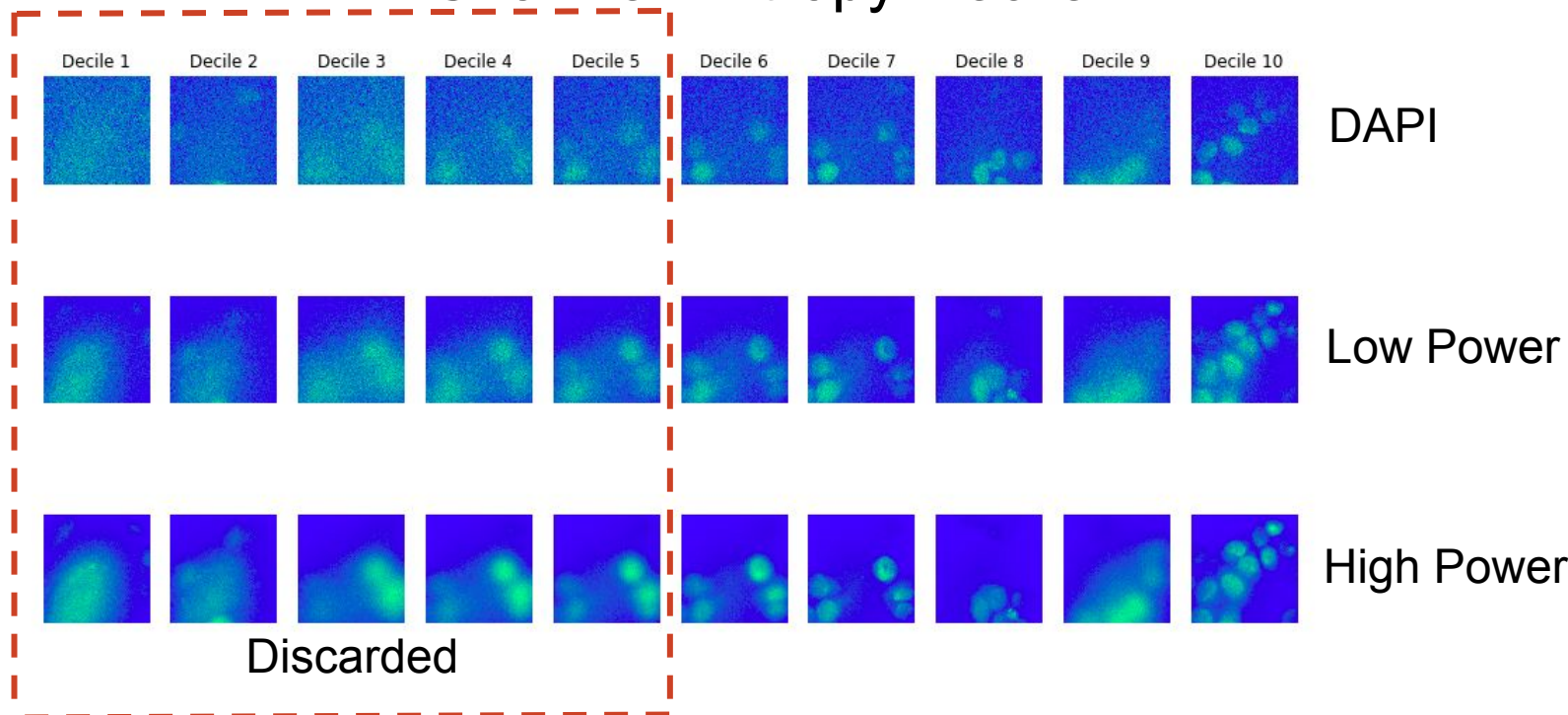
- DAPI highlights cell nuclei
- GFP tracks mRNA distribution

Each sample consists of a stack of 100 focal planes (z-slices) with 7 channels at each slice (1 for DAPI, 6 at varying power for GFP).



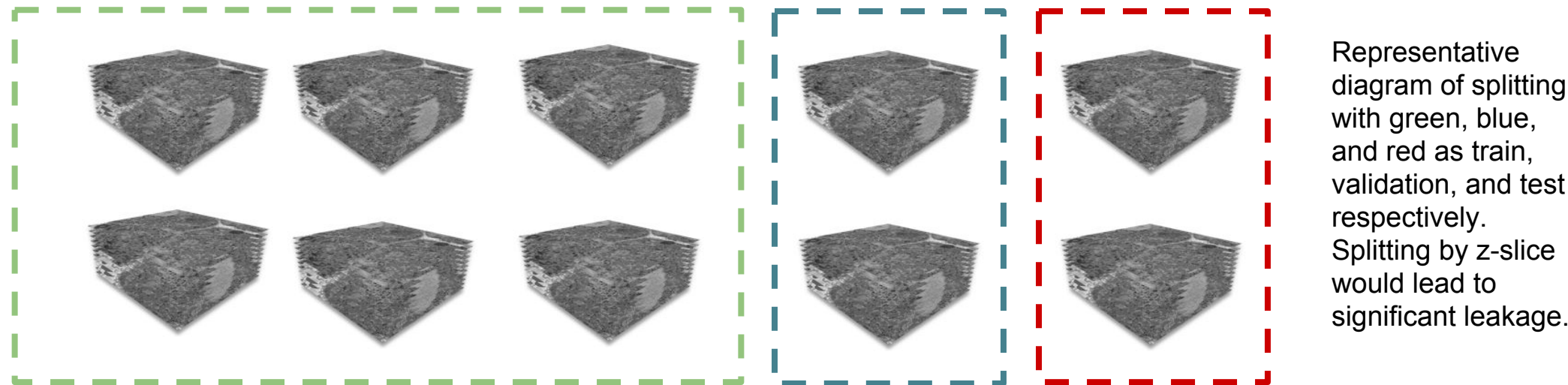
Data Selection: Due to microscope focus issues, many z-slices contain fuzzy images that are of low quality in all channels, regardless of laser intensity.

DAPI Channel Entropy Decile



We select image sets with suitable clarity using the entropy of pixel intensities for an additional DAPI stained channel.

Data Preprocessing: After selection, we split data into train/validation/test by stack because we expect adjacent z-planes to be similar enough to cause leakage issues if separated. In addition, data was augmented and normalized using standard techniques.



Evaluation Methods

MSE was generally used as the training loss (unless certain baselines had other loss functions for specific components) while evaluation was done with Structural Similarity Index (SSIM):

$$SSIM(x, y) = \frac{(2\mu_x\mu_y + c_1)(2\sigma_{xy} + c_2)}{(\mu_x^2 + \mu_y^2 + c_1)(\sigma_x^2 + \sigma_y^2 + c_2)}$$

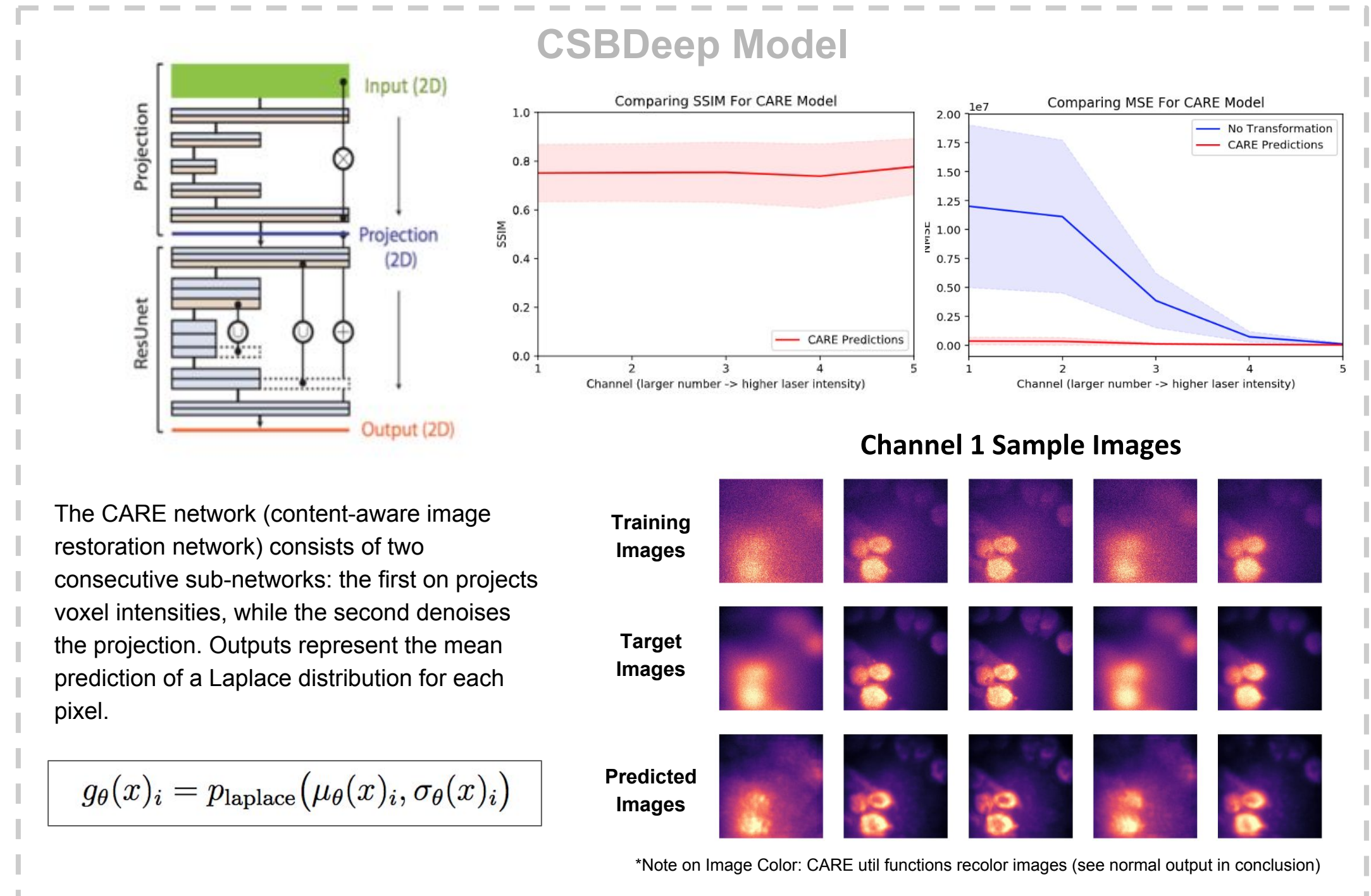
Here, μ and σ are the mean and variance (or covariance) respectively of the distribution of pixel intensity in each image. The constants are for numerical stability and generally a small value scaled with intensity range. This is computed for a small patch at a time.

- Note that evaluation methods such as SSIM and MSE are *not scale invariant* to the image data type (e.g. int16, uint32, float, etc.). Thus, we convert our data image type to float for the SSIM comparison.

- The reported SSIM values are an average of this score over corresponding patches of the two images. SSIM lies in the range [-1,1] with a score of 1 implying identical images.

Data Type	Data Range
int8	0 to 255
int16	-32768 to 32767
float	0 to 1
uint32	0 to 2 ³¹ - 1

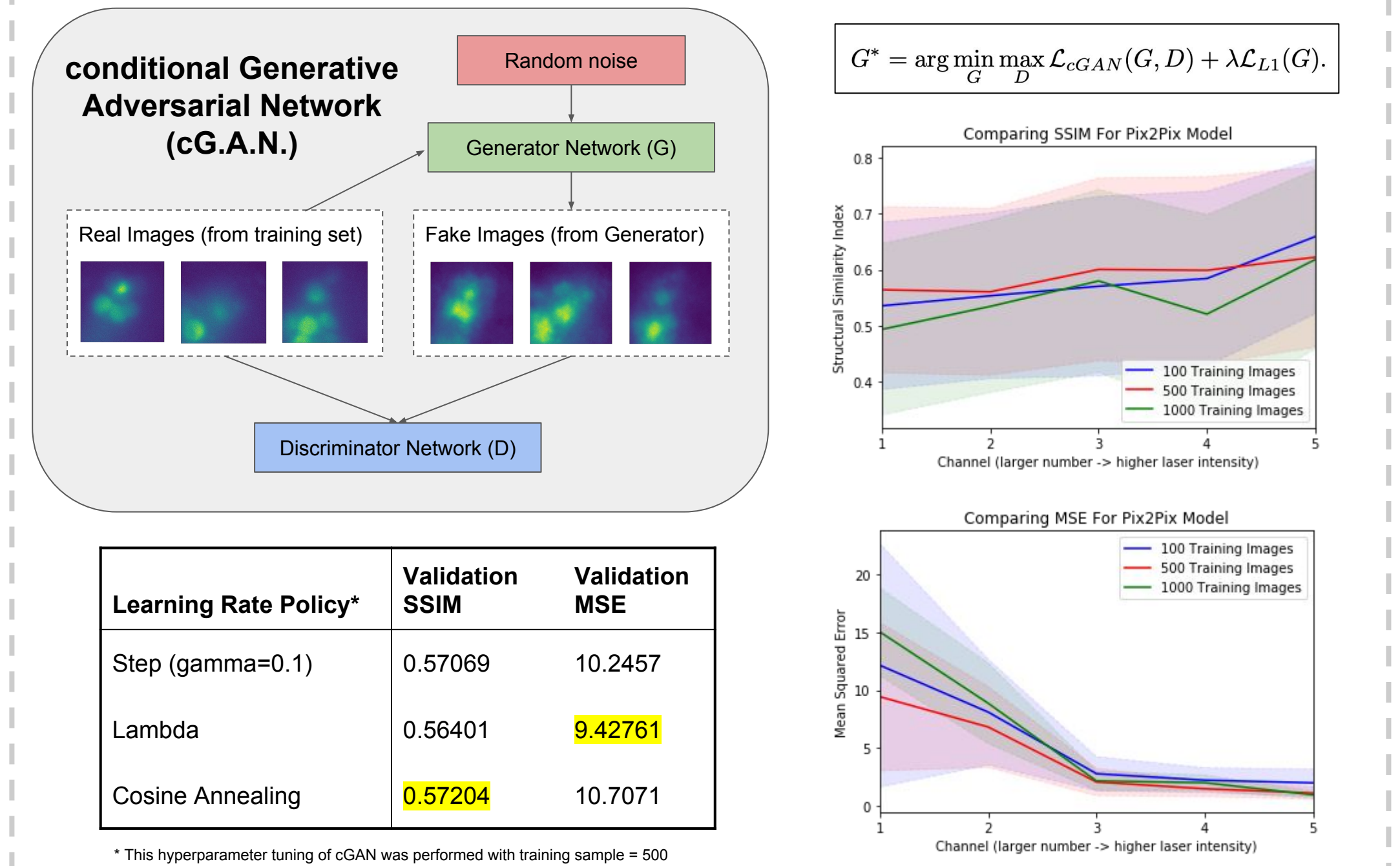
Baselines



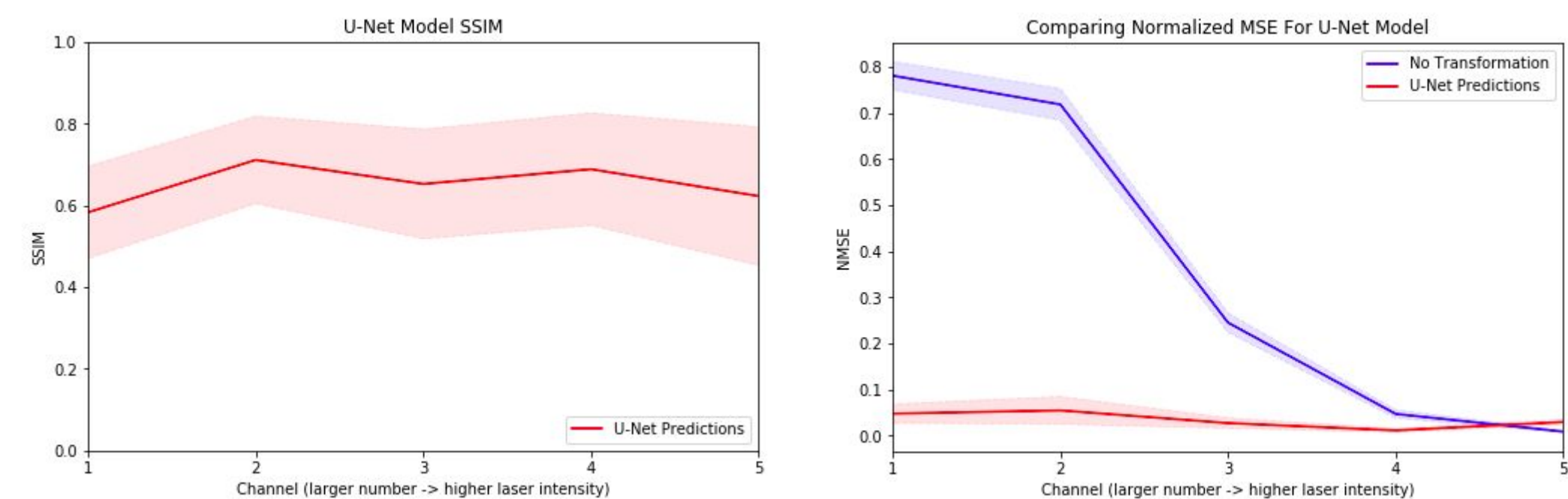
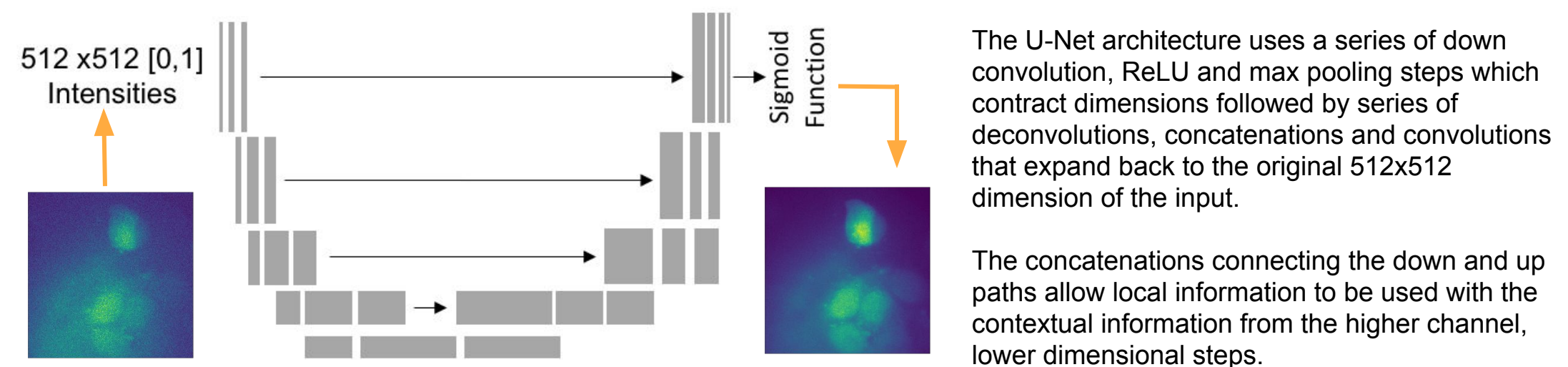
The CARE network (content-aware image restoration network) consists of two consecutive sub-networks: the first on projects voxel intensities, while the second denoises the projection. Outputs represent the mean prediction of a Laplace distribution for each pixel.

$$g_{\theta}(x)_i = p_{\text{laplace}}(\mu_{\theta}(x)_i, \sigma_{\theta}(x)_i)$$

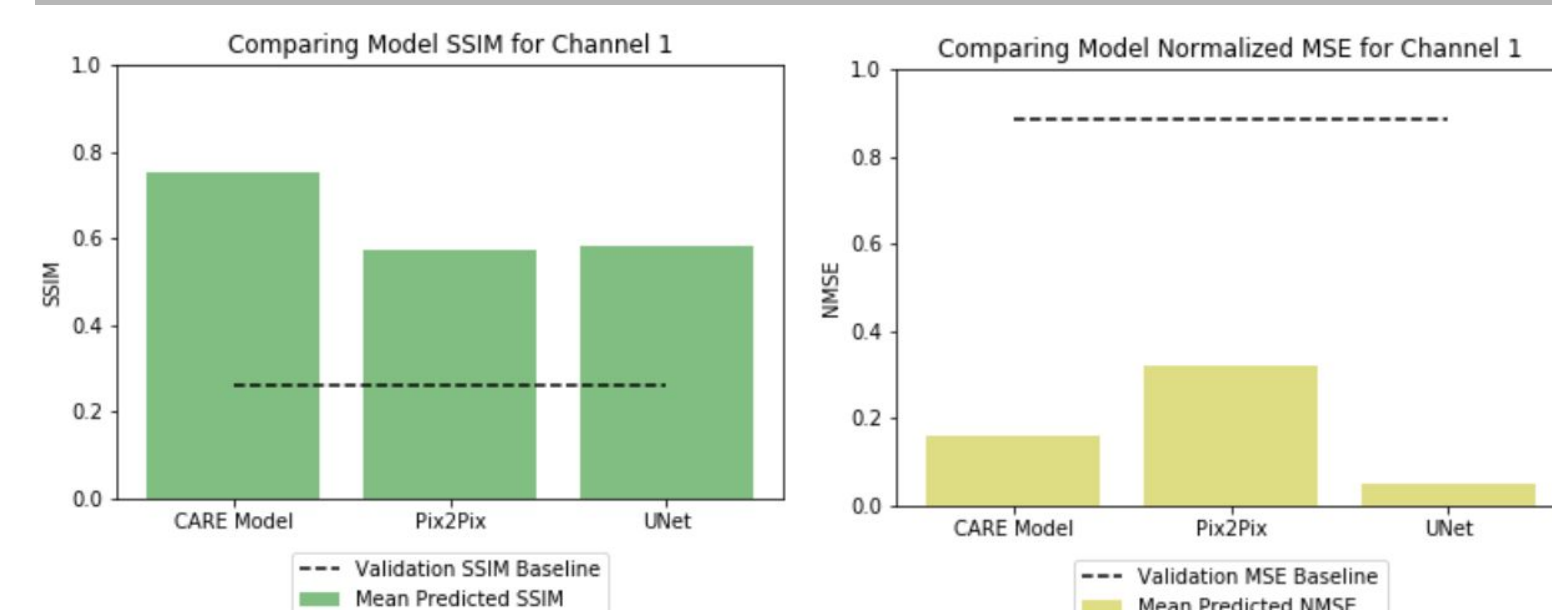
Pix2Pix Model



U-Net Model



Conclusion



For future applications, predictions based on Channel 1 (lowest power) provides the most value to end users.

Based on the mean validation set SSIM for Channel 1, the CARE Model performs best with a mean SSIM of 0.75.

