



Edge Hill University 



PAPER PRESENTATION

HistDiT: A Structure-Aware Latent Conditional Diffusion Model for High-Fidelity Virtual Staining in Histopathology

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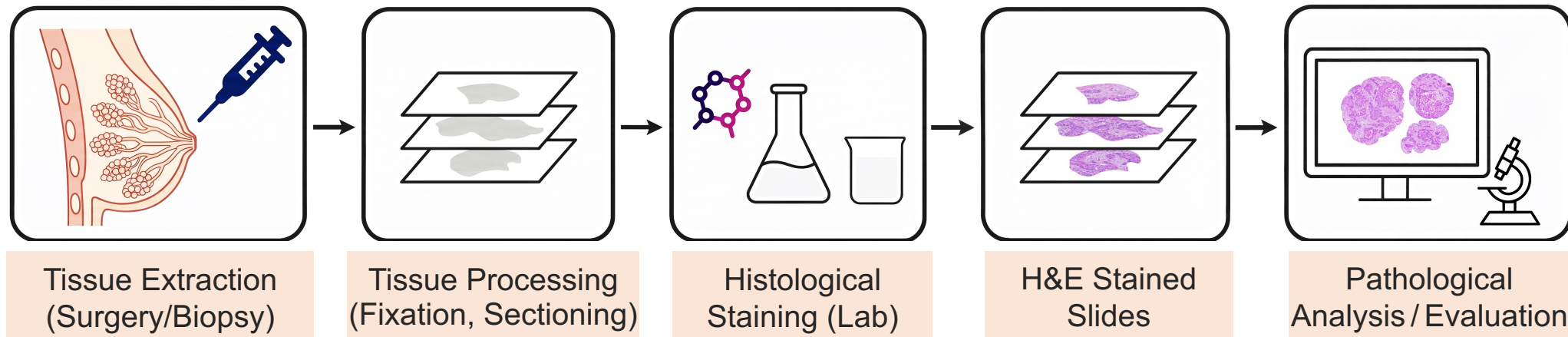
 Paper: arxiv.org/abs/2604.08305

 Code: github.com/AasimBinSaleem/HistDiT

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MOTIVATION · CLINICAL CONTEXT

- Precise treatment for **Breast Cancer** involve assessment of biomarkers.
- **Histological staining** serves as the trademark in cancer diagnosis.

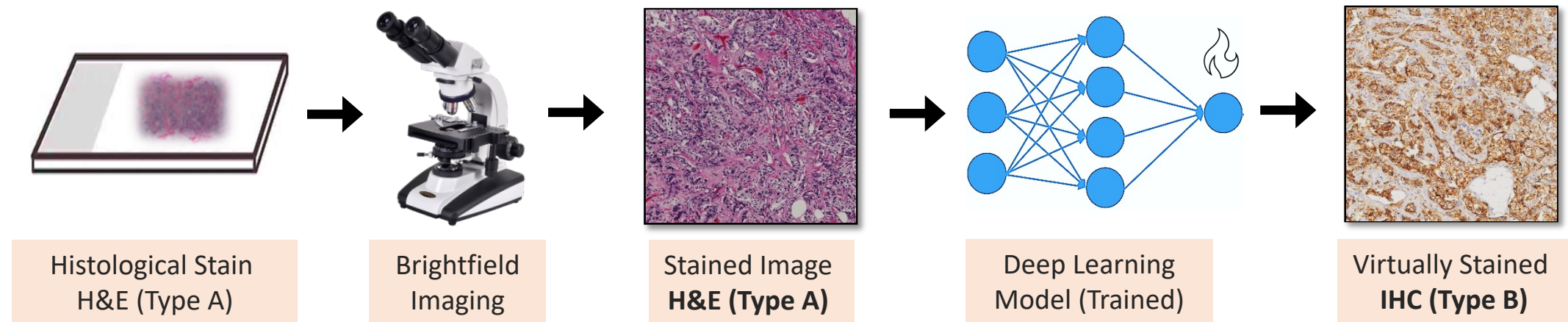


- ❖ **H&E Staining:** Standard, cheap, but lacks biomarker specificity.
- ❖ **IHC Staining:** Identifies a variety of immune biomarkers, dictates targeted therapies for early personalized treatments.

IHC is resource-intensive, time-consuming and damages tissue homeostasis.

VIRTUAL STAINING

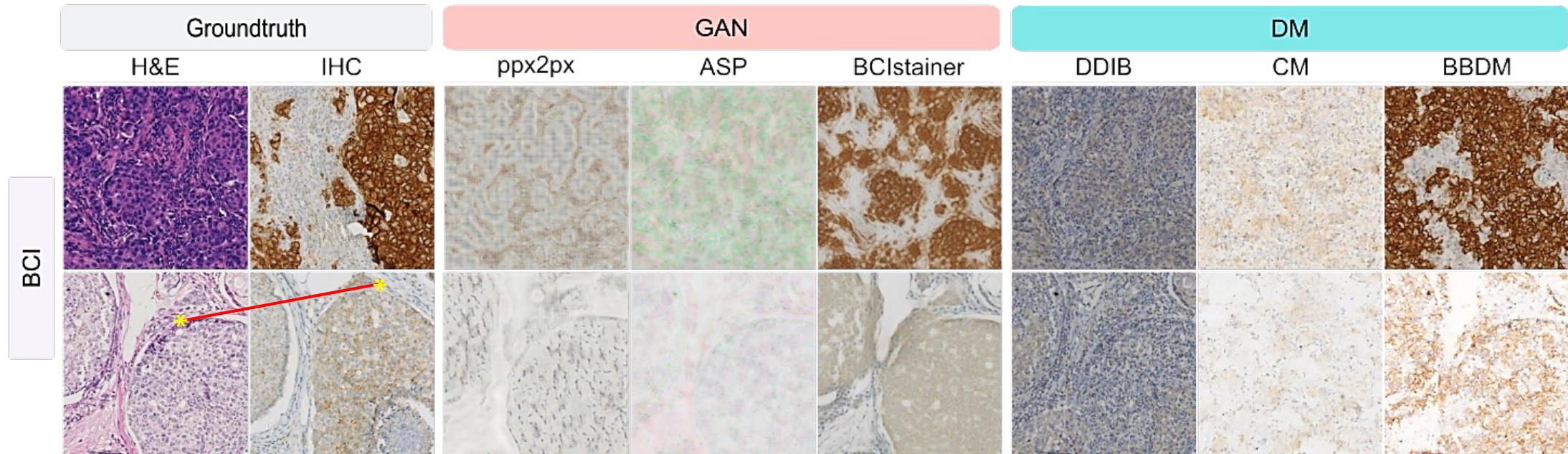
- Virtual staining uses AI to generate synthetically stained IHC samples.
- ✓ Provides instant, selective, cost-free and high-contrast diagnostics.



💡 GOAL

H&E stained to IHC stained image translation for breast cancer prognosis.

PRIOR WORK

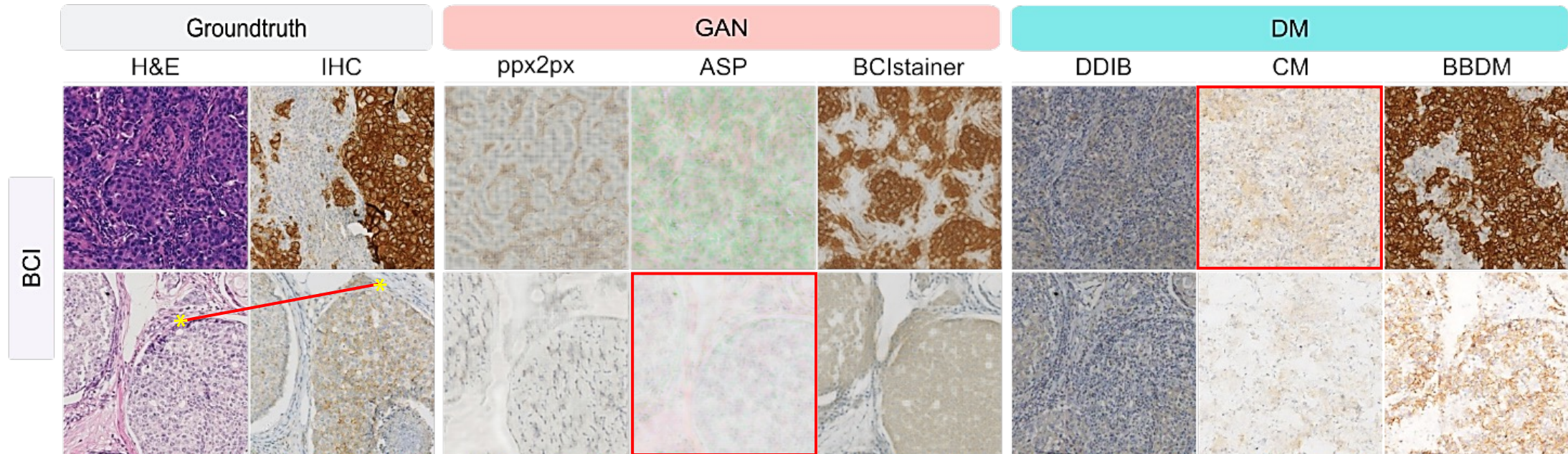


Examples for GANs and Diffusion models, trained on BCI Dataset [6]

▲ LIMITATIONS:

- **GANs:** Difficult to **train** – suffer from mode collapse issues; **Wash out** high expression regions.
- **Standard Diffusion (U-Nets):** Structure and staining **trade-offs**.
Struggle with global context – **hallucinate** cells and misalign structures in complex tissues.

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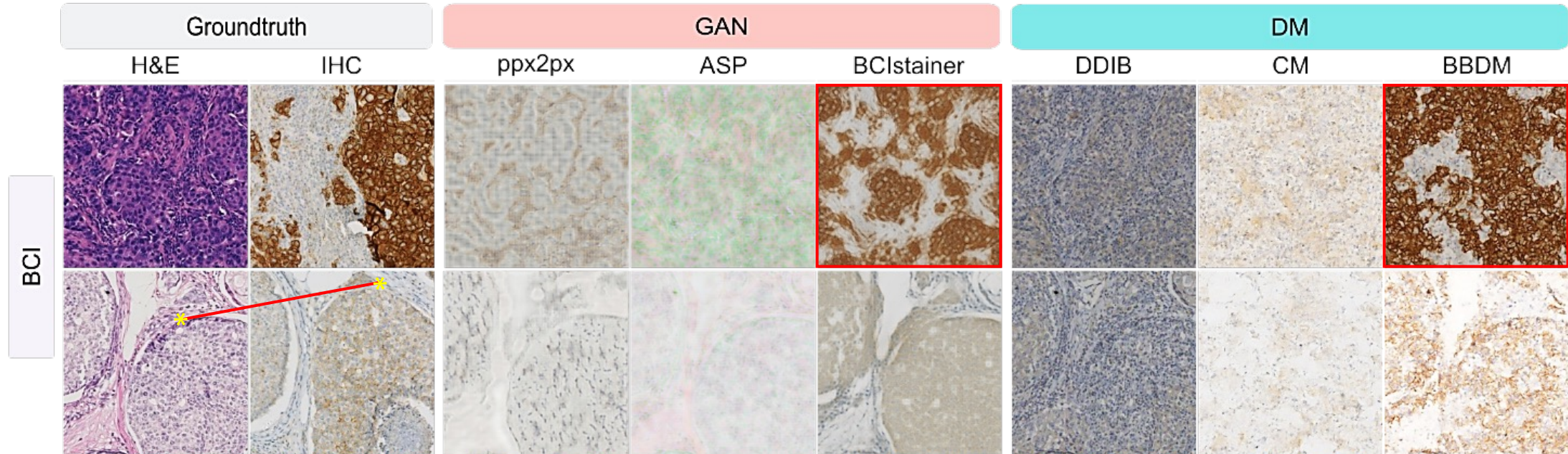


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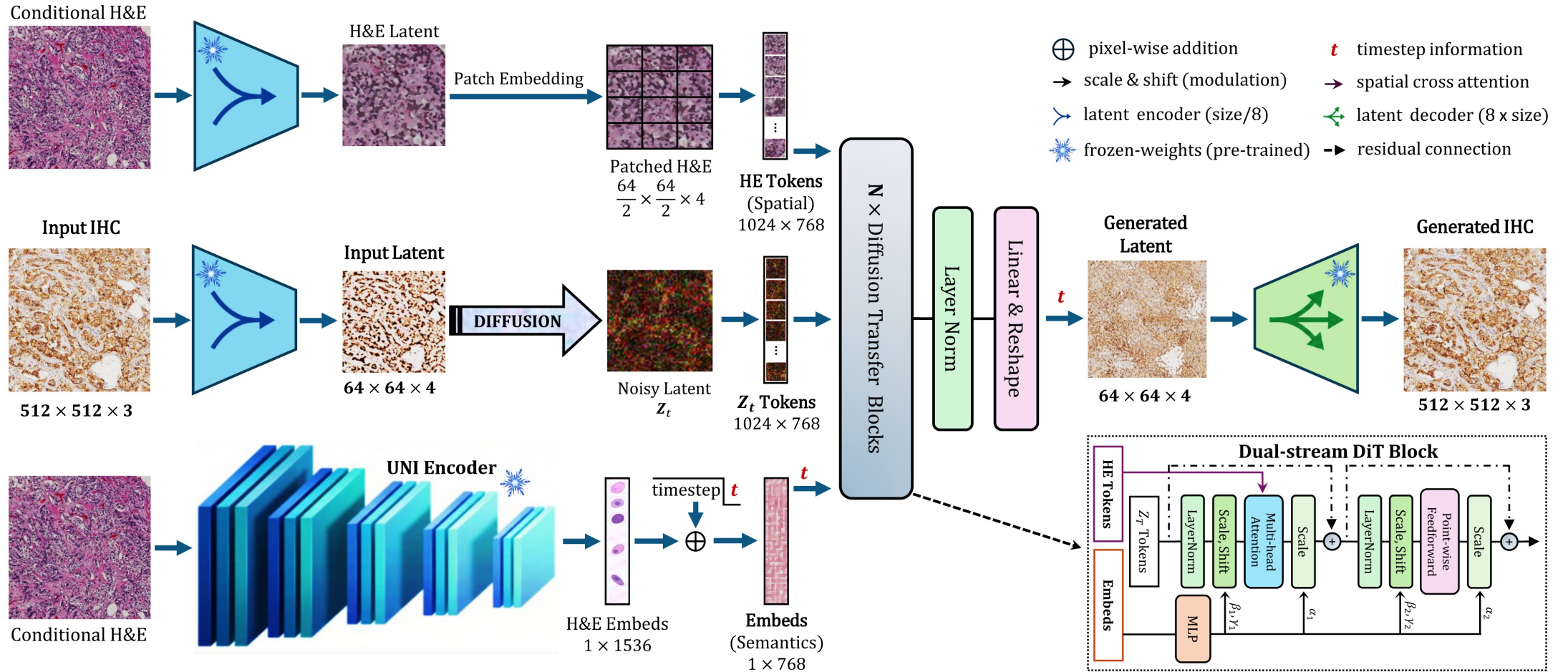


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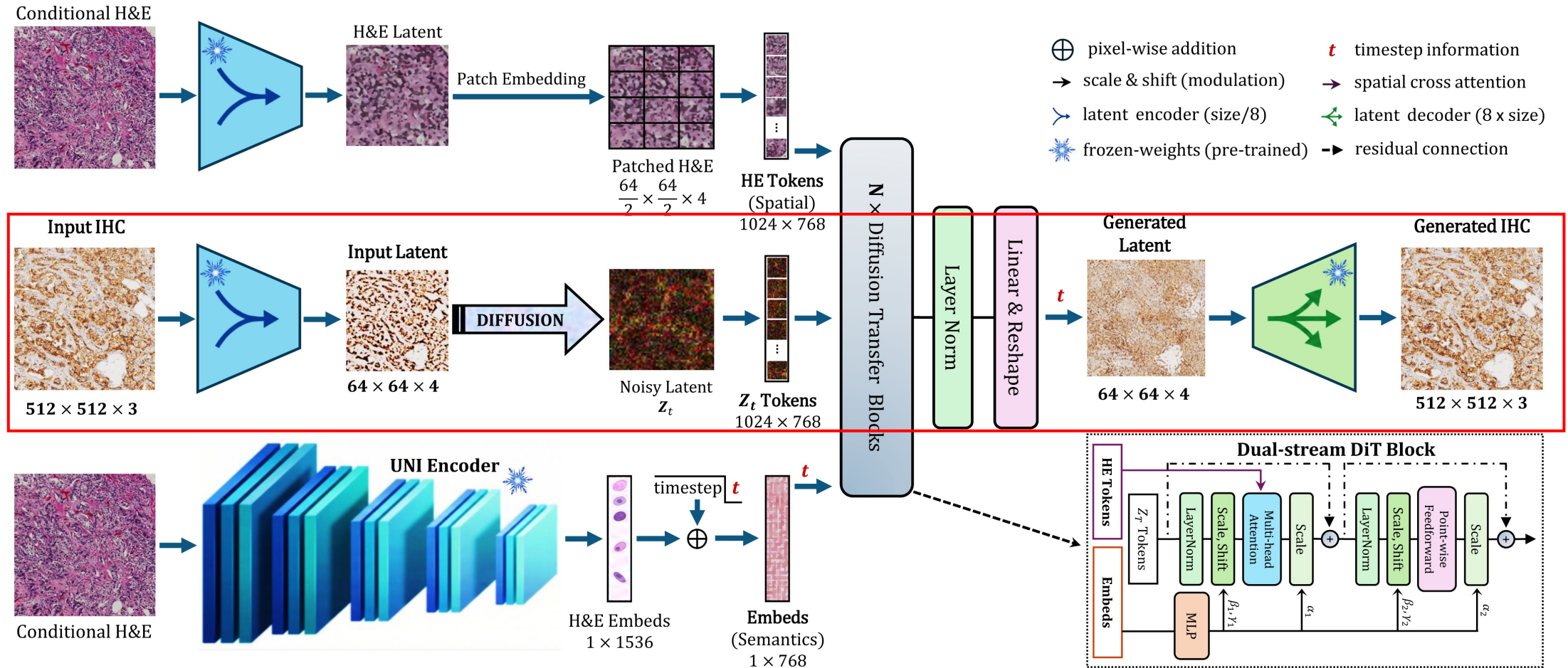
PROPOSED VIRTUAL STAINING FRAMEWORK



Training objective: $\mathcal{L}_{\text{Train}} = \alpha \cdot \mathcal{L}_{\text{MSE}} + (1 - \alpha) \cdot \mathcal{L}_{\text{L1}}$

Noise Scheduler: Standard DDPM (epsilon prediction)

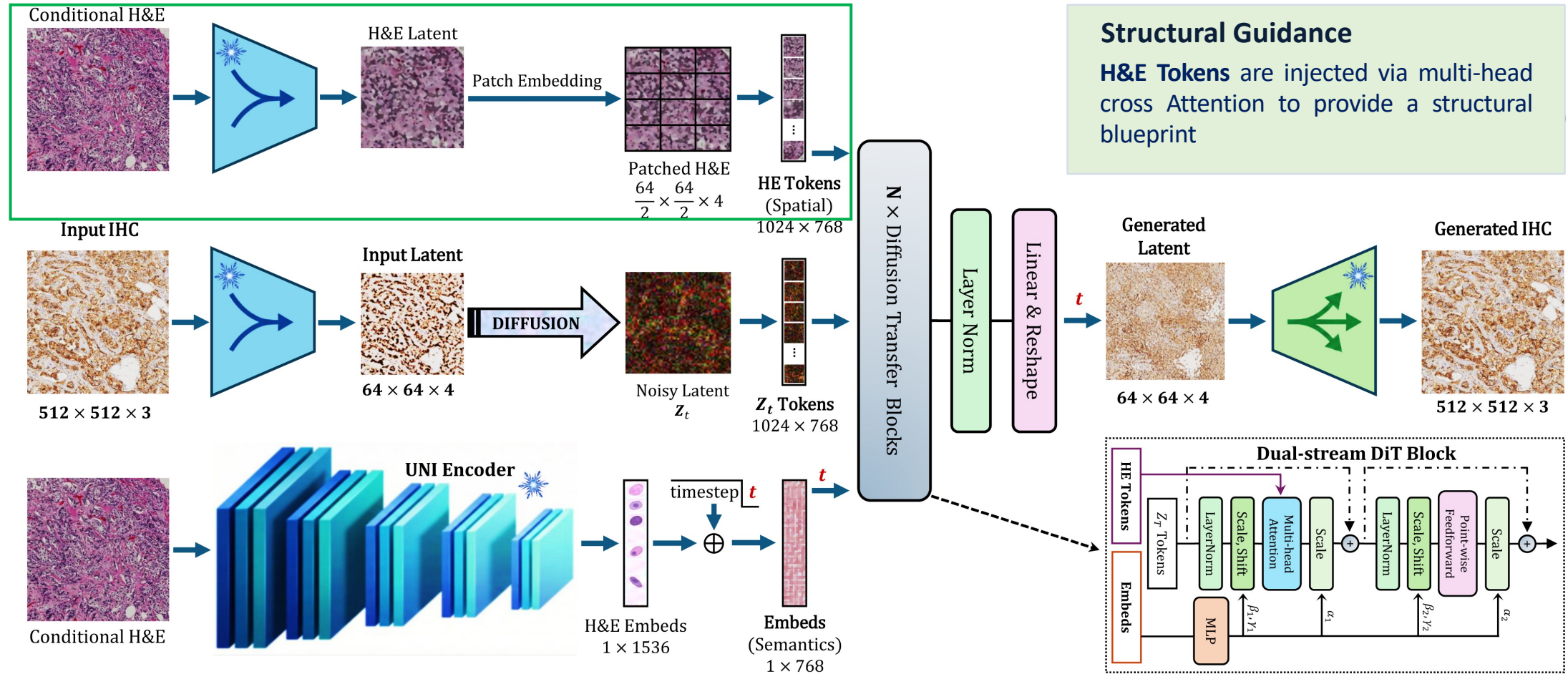
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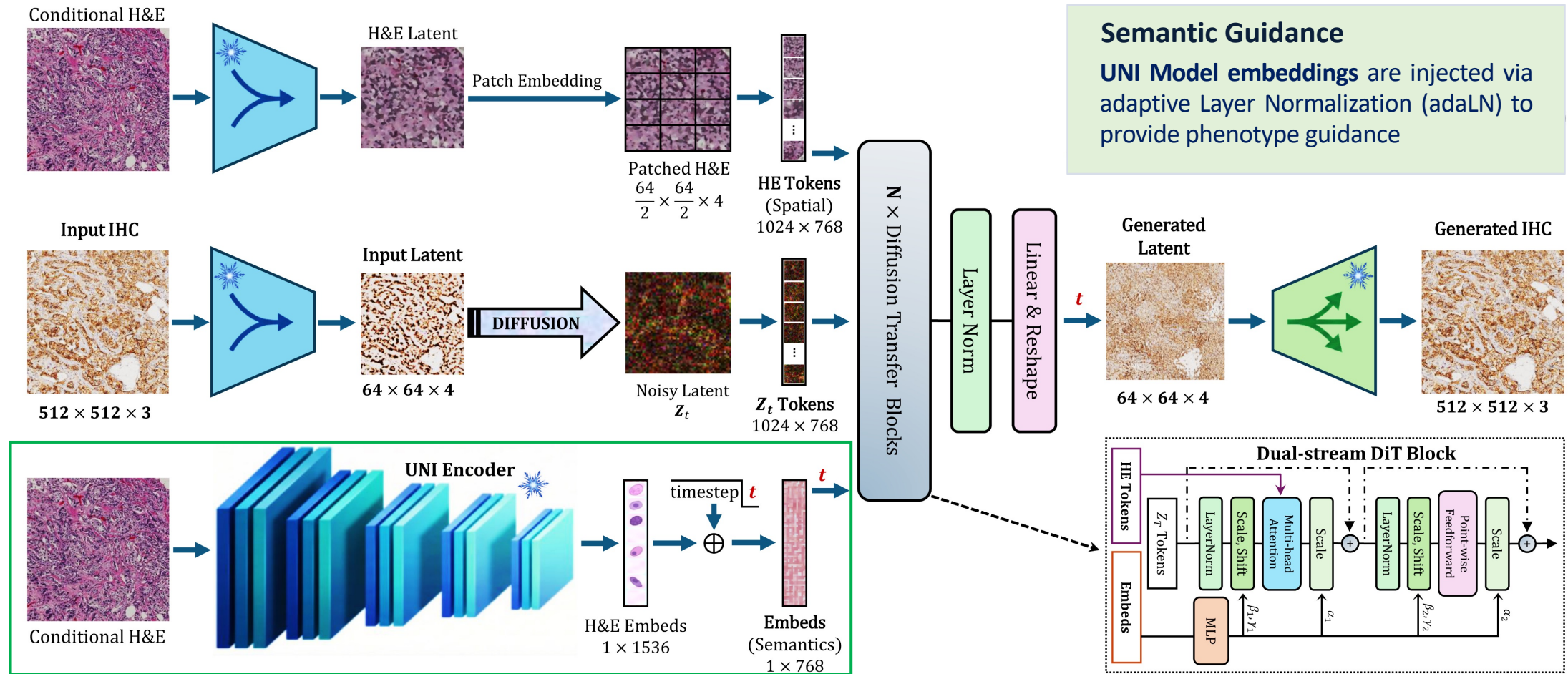
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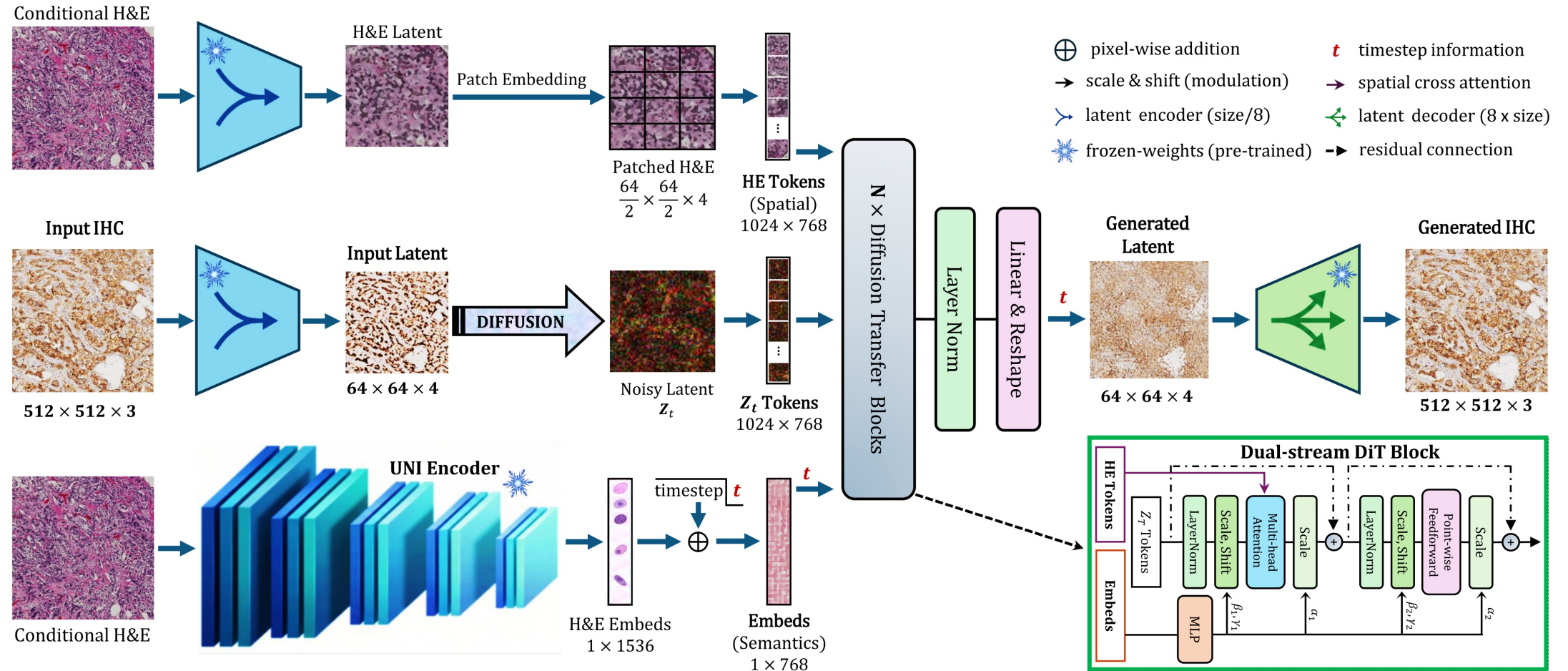
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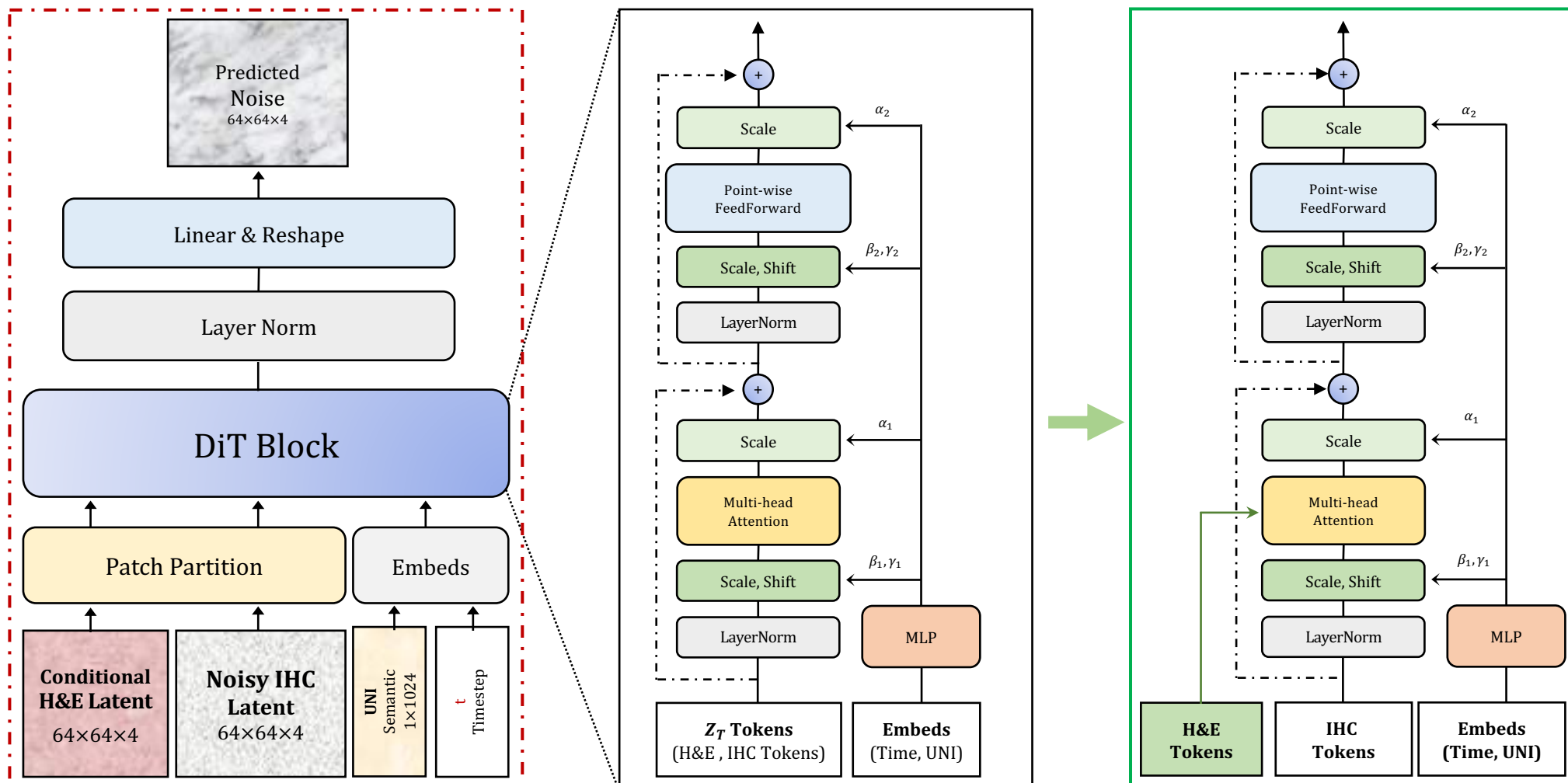
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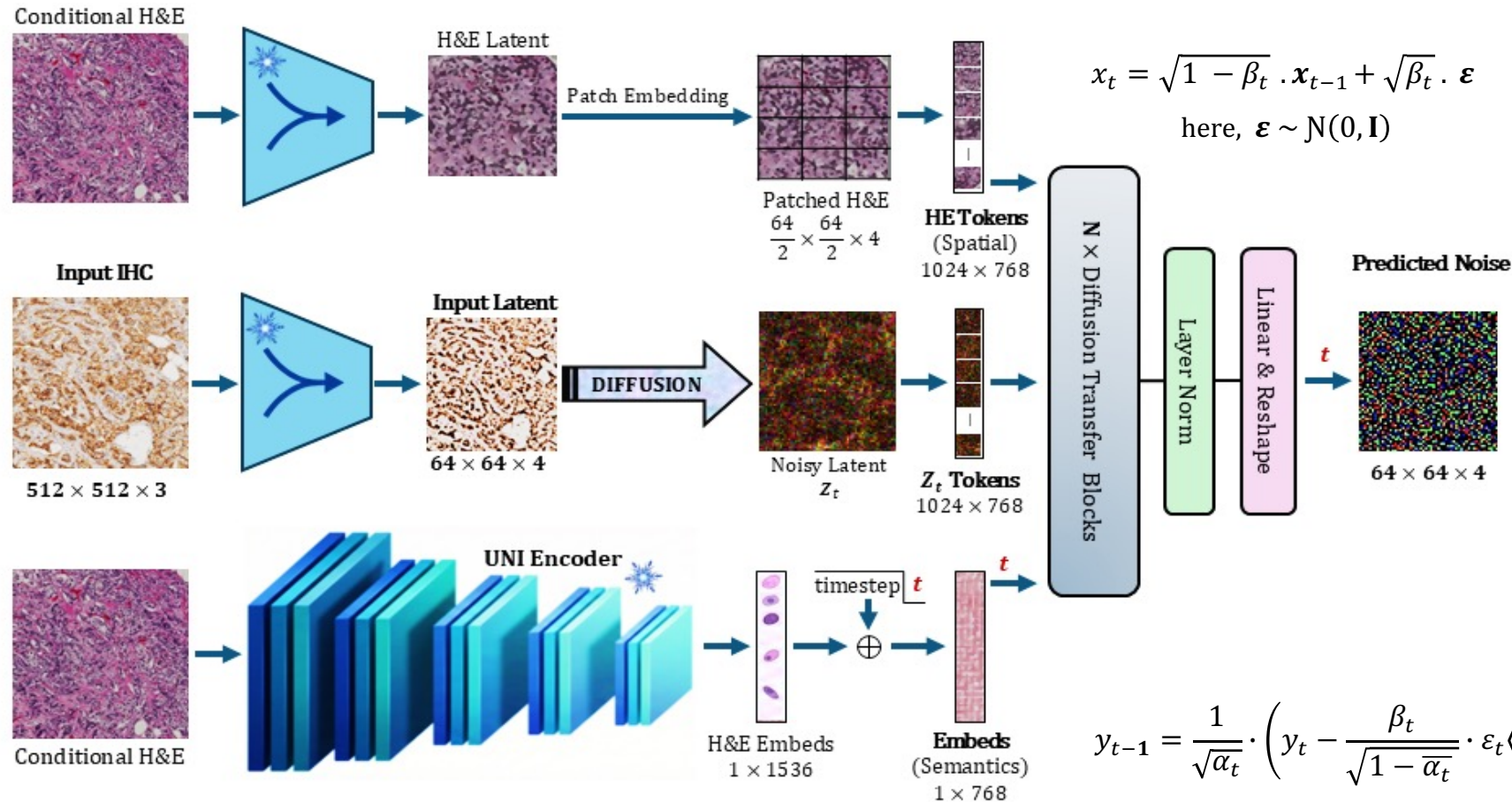
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Noise Scheduler: Standard DDPM (epsilon prediction)

DUAL-STREAM DIT BLOCK



TRAINING PIPELINE



Multi-Objective Loss

Combining an auxiliary L_1 term with the standard MSE

$$L_{Total} = 0.7 \times L_{MSE} + 0.3 \times L_{L1}$$

Predicted Noise (ϵ_t')



$\mathcal{L}_{Train}$

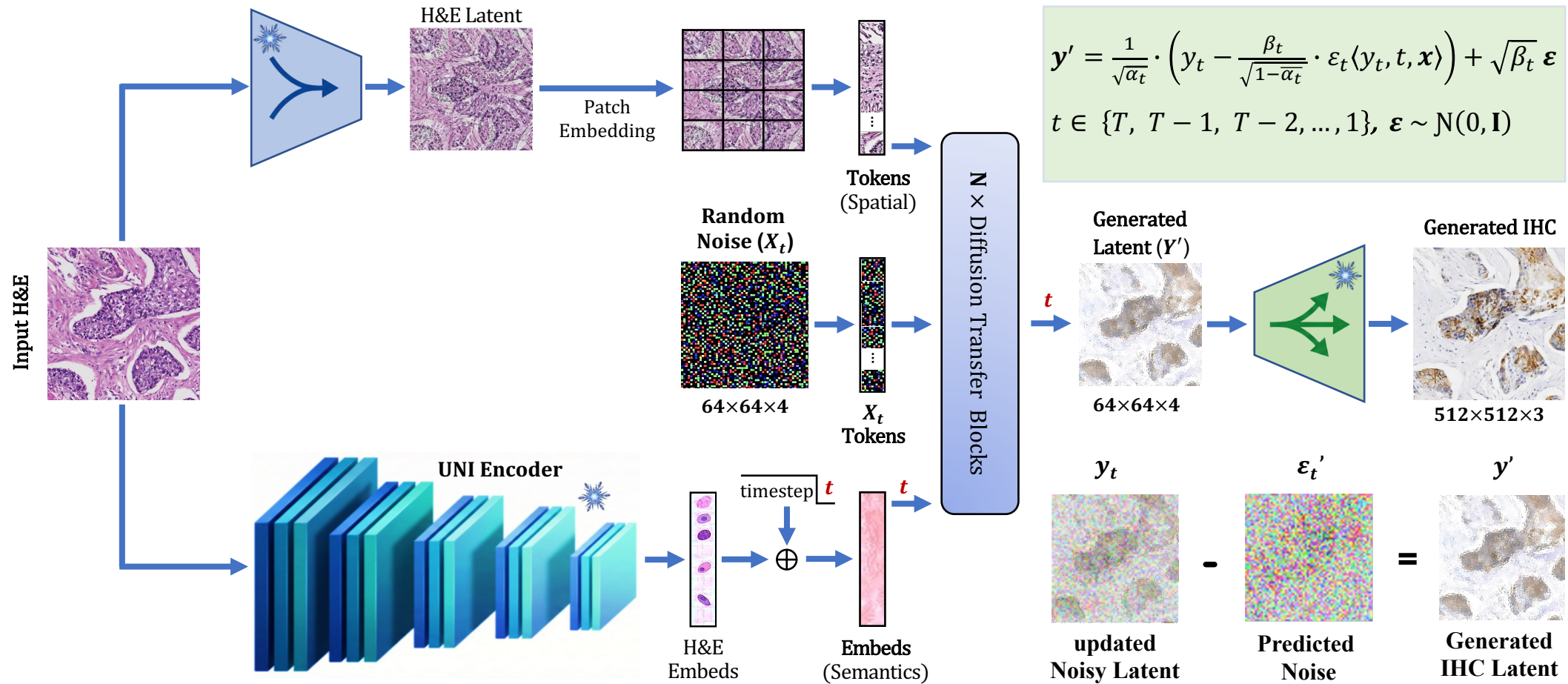


Actual Noise (ϵ_t)

Training objective: $\mathcal{L}_{Train} = \alpha \cdot \mathcal{L}_{MSE} + (1 - \alpha) \cdot \mathcal{L}_{L1}$

Noise Scheduler: Standard DDPM (epsilon prediction)

THE INFERENCE



Noise Scheduler: Standard DDPM (scaled linear)

Prediction Type: Epsilon

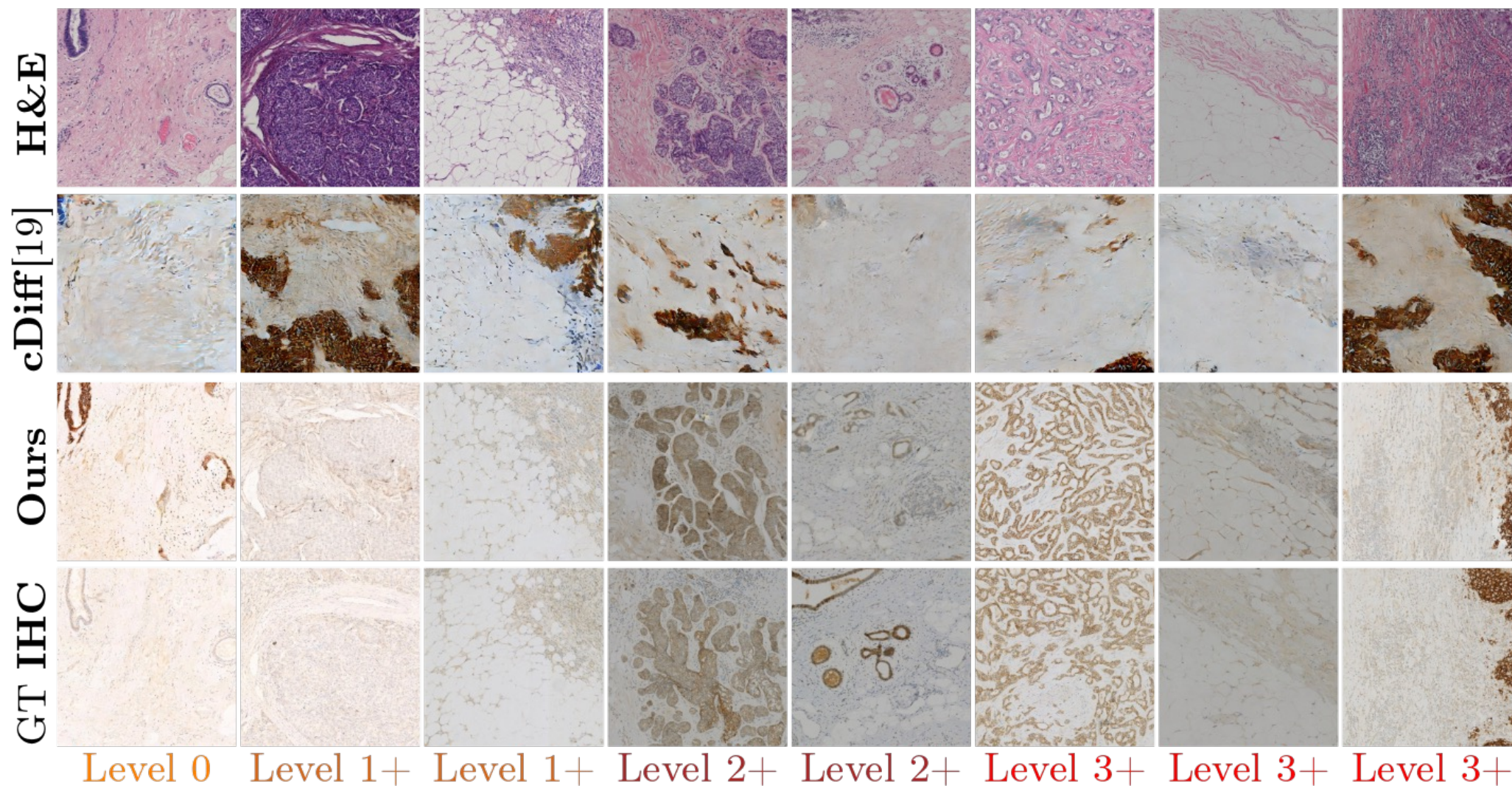
Sampling Steps: 1000

QUANTITATIVE COMARISON ON BCI DATASET

Method/Model	MSE ↓	PSNR(dB) ↑	SSIM ↑	SCM ↑	LPIPS ↓	FID ↓	
Cycle GAN [4] [Ⓔ]	1892.00	15.98	0.372	0.447	0.624	188.33	GANs
Pix2Pix [5] [Ⓔ]	1560.26	17.08	0.303	0.391	0.769	<u>160.39</u>	
Pyramid Pix2Pix [6] [Ⓔ]	<u>1348.54</u>	19.61	0.397	0.473	<u>0.466</u>	167.40	
ASP [7] [Ⓔ]	---	---	0.5032	---	---	---	
Syn Diff [8] [Ⓔ]	---	14.28±2.52	0.32±0.1	---	---	---	Diffusion
PST-Diff [9] [Ⓔ]	---	16.75±4.20	0.38±0.1	---	---	---	
Conditional Diff. [19] [Ⓔ]	3011.36	15.76	0.405	<u>0.494</u>	0.591	194.82	
Star-Diff (No Visual) [Ⓔ]	---	<u>21.30±0.01</u>	0.5301	---	---	---	
HistDiST [21] [Ⓔ]	---	---	<u>0.4693</u>	---	---	---	OUR
Proposed [HistDiT] [Ⓔ]	891.53	21.43	0.4769	0.540	0.412	49.15	
%age Improvement	33.89%	0.66%	5.2%	9.31%	11.59%	69.4%	

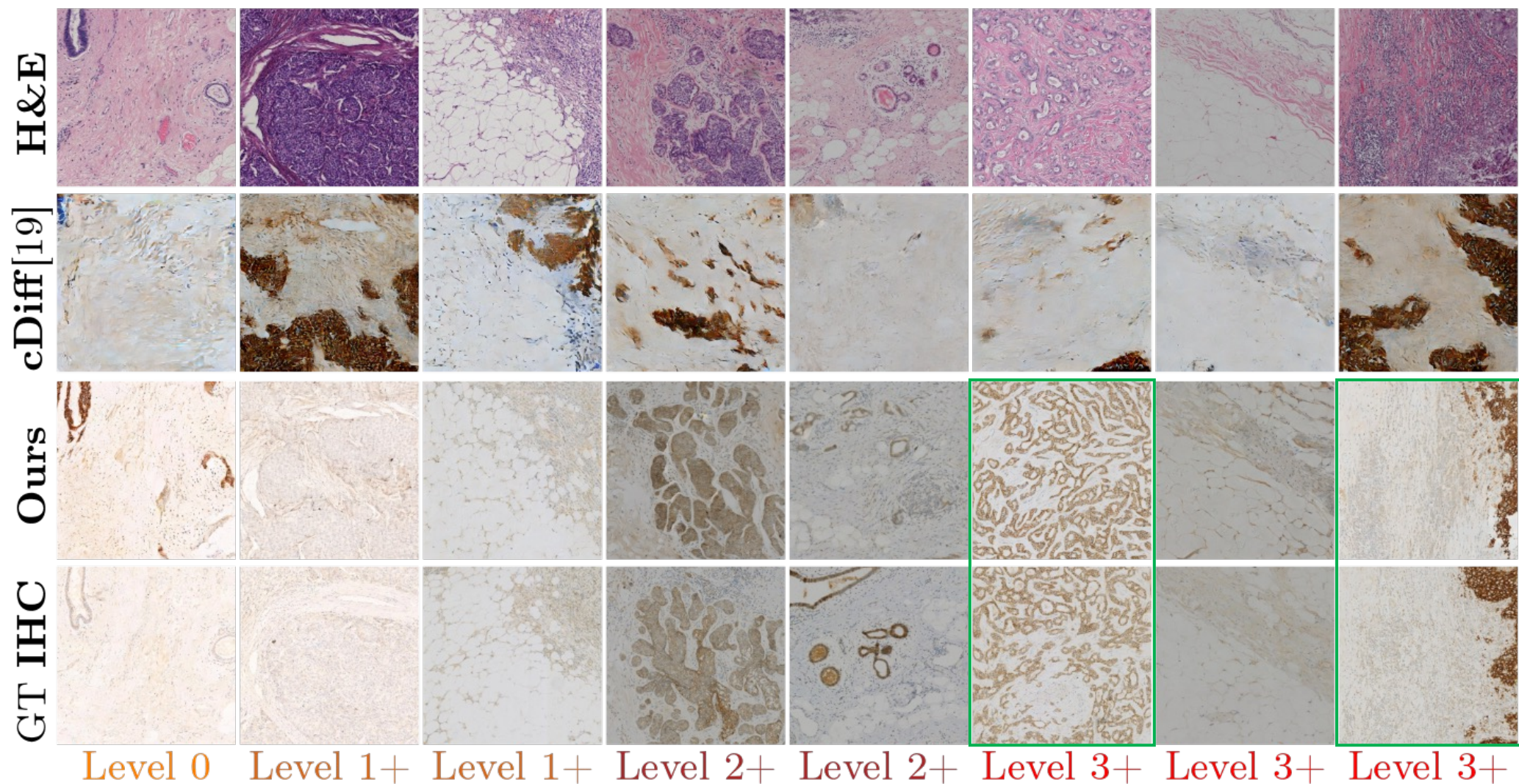
💡 Structural Correlation Metric (**SCM**) mitigates mathematical bias in standard SSIM towards brightfield microscopy.

VISUAL COMARISON ON THE BCI DATASET



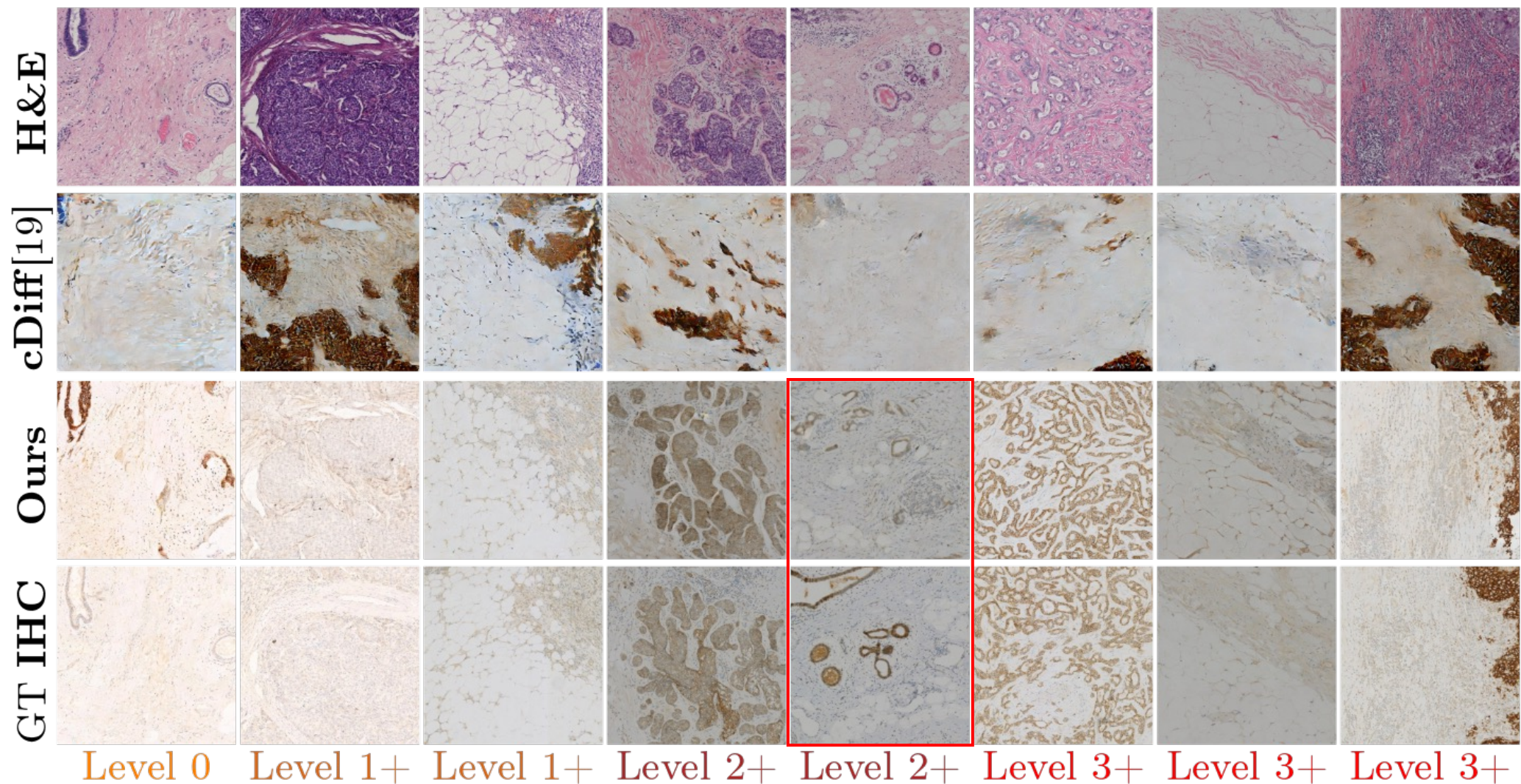
HER-2 Biomarker Grading

VISUAL COMARISON ON THE BCI DATASET



HER-2 Biomarker Grading

VISUAL COMARISON ON THE BCI DATASET



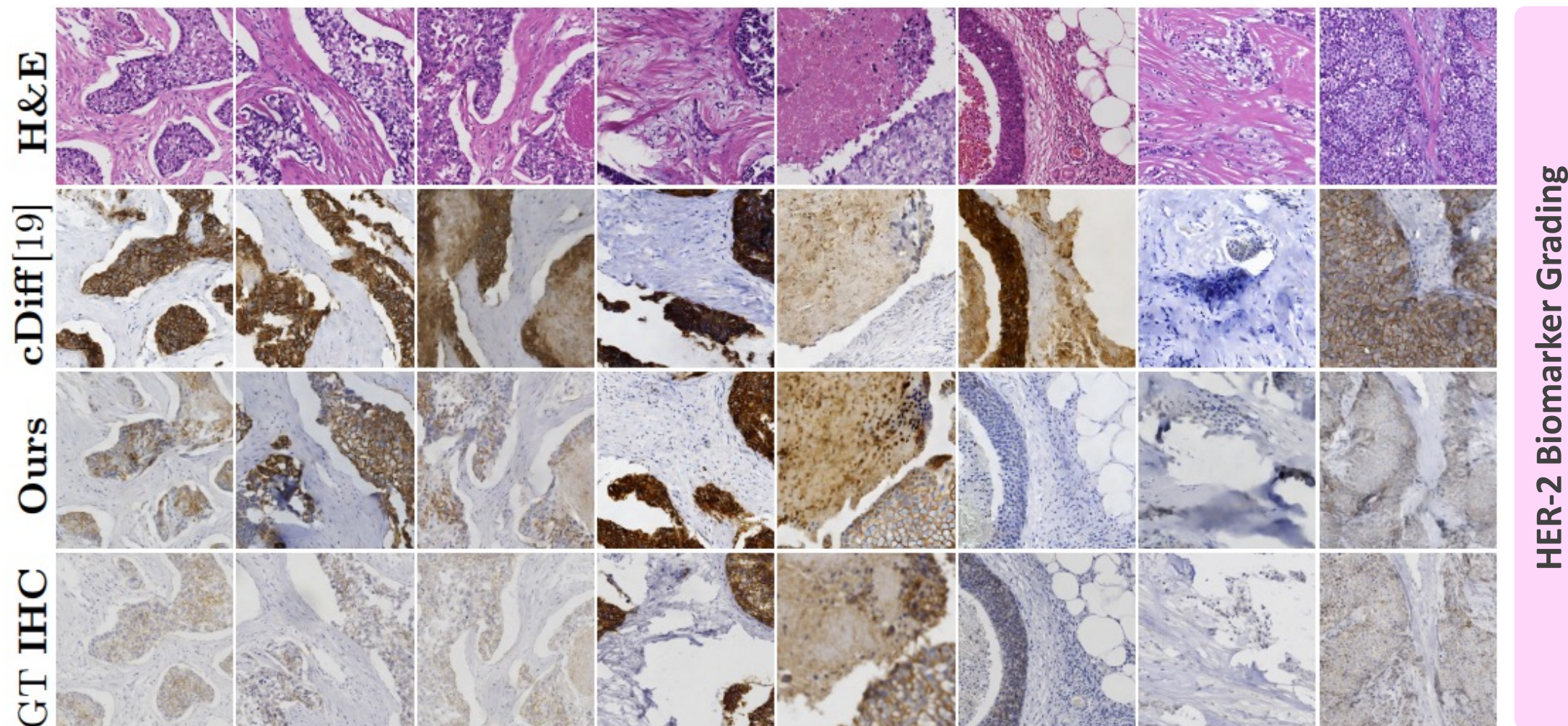
HER-2 Biomarker Grading

QUANTITATIVE COMARISON ON MIST DATASET

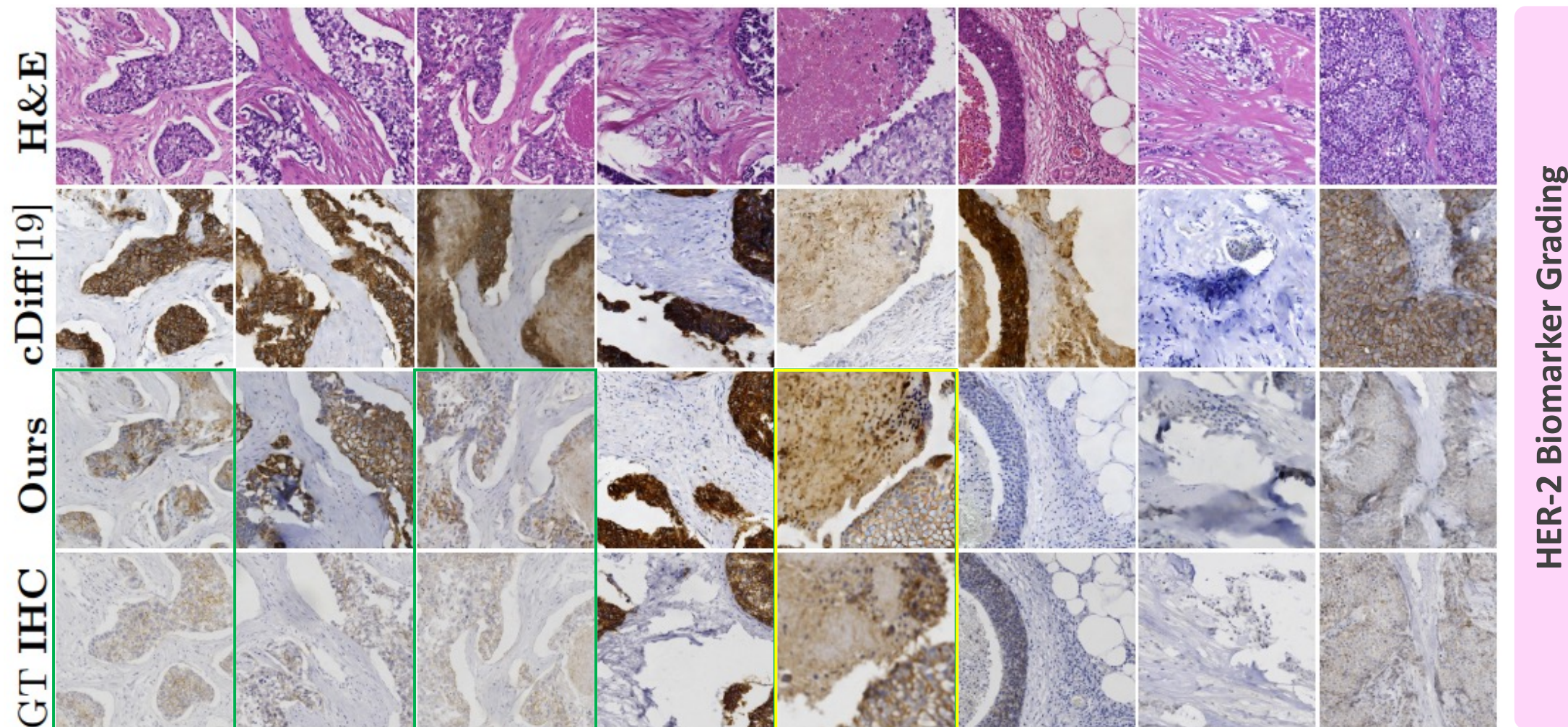
Method/Model	MSE ↓	PSNR(dB) ↑	SSIM ↑	SCM ↑	LPIPS ↓	FID ↓	
Cycle GAN [4] [Ⓔ]	4190.52	11.91	0.174	0.209	0.553	125.7	GANs
Pix2Pix [5] [Ⓔ]	3485.20	12.74	0.150	0.194	0.614	128.1	
Pyramid Pix2Pix [6] [Ⓔ]	<u>2986.44</u>	<u>14.24</u>	0.165	0.214	<u>0.543</u>	107.4	
ASP [7] [Ⓔ]	6252.98	11.13	0.168	<u>0.254</u>	0.564	82.6	
Conditional Diff. [19] [Ⓔ]	---	---	0.1810	---	---	---	DMs
HistDiST [21] [Ⓔ]	---	---	0.1945	---	---	54.28	
PixCell (no LoRA) [38] [Ⓔ]	---	---	0.1880	---	---	67.68	
Proposed [HistDiT] [Ⓔ]	3396.8	14.26	0.211	0.302	0.489	59.30	OUR
%age Improvement	13.74%	0.14%	2.5%	18.9%	9.94%	9.25%	

💡 Standard **SSIM** has a mathematical bias towards brightfield microscopy, where luminance masks structural errors. Therefore, we suggest the Structural Correlation Metric (**SCM**) for more honest comparison.

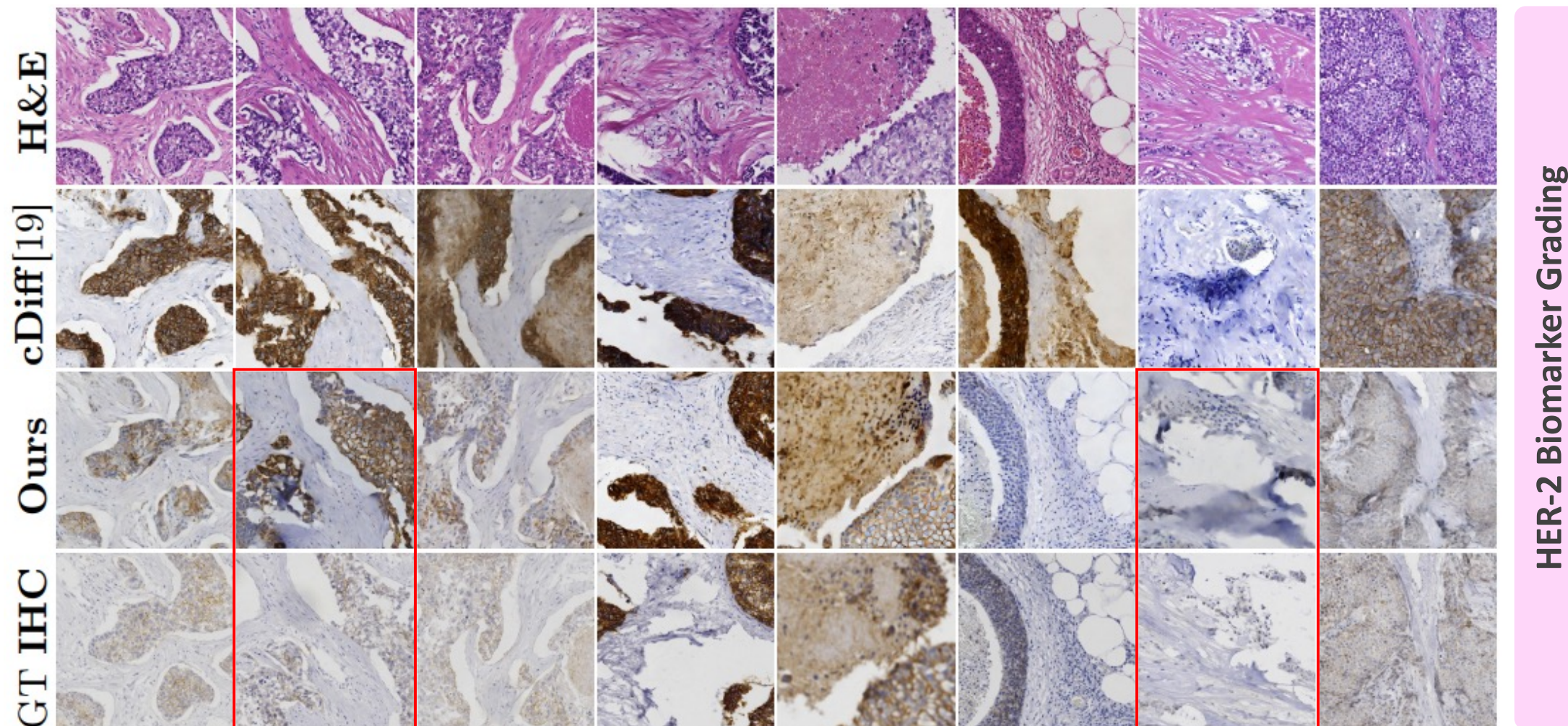
VISUAL COMARISON ON MIST DATASET



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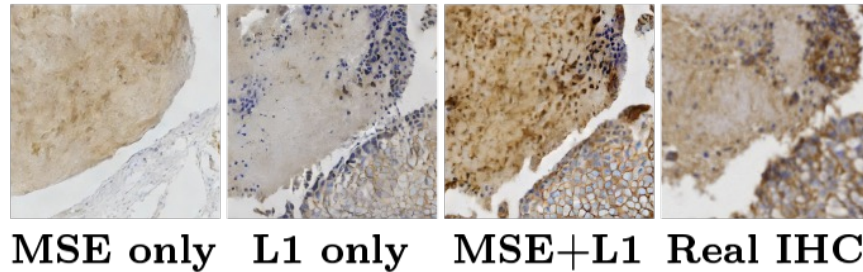
ABLATION STUDIES

Model Configuration	PSNR(dB)↑	SSIM↑	SCM↑	LPIPS↓	FID↓
Pyramid Pix2Pix [6] ^ρ	19.61	0.397	0.473	0.466	167.40
HistDiT (Semantics only) ^z	16.39	0.423	0.492	0.553	81.51
HistDiT (Spatial only with Cross-Attention) ^z	18.58	0.416	0.560	0.452	62.38
HistDiT (Spatial [Concatenation], Semantics) ^z	20.63	0.449	0.521	0.438	56.80
HistDiT (Spatial [Cross-Attention], Semantics) ^z	21.43	0.477	0.540	0.412	49.15

Impact of
Dual-stream
Conditioning

Table Comparison for different architectural choices on BCI Dataset. Best are **bold**.

Method/Approach	PSNR↑	SSIM↑	FID↓
MSE only	16.86	0.438	67.88
L_1 only	19.44	0.450	93.31
$0.9MSE + 0.1L_1$	20.44	0.443	56.18
$0.7MSE + 0.3L_1$	21.43	0.477	49.15



Impact of
Multi-objective
Training Loss

Fig. Ablation Study on Objective Functions: (Left) Quantitative comparison of loss components on the BCI dataset. $MSE+L_1$ maintains high metric scores. (Right) Visual samples demonstrating that the combined objective produces sharper structures (column 3) compared to the smoothing artifacts seen in MSE only (column 1).

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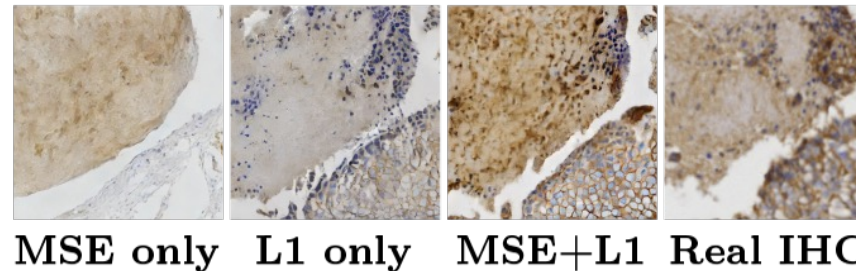


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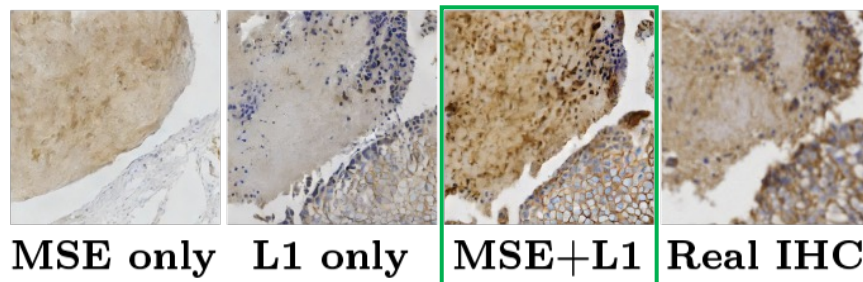


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VALIDATION & FUTURE DIRECTIONS

- ❖ **Experts** from CRMV confirmed the staining fidelity of our generated samples via blind qualitative assessment: **Indistinguishable** from real IHC.

LIMITATIONS & FUTURE WORK

- Computational overhead as main limitation.
- Domain gap for using latent encoders that offer lossy compression.
- Detailed HER-2 grading analysis, including Cohen's Kappa to be obtained.
- Stain normalization for cross-dataset evaluation.



[READ MORE](#)

Paper: arxiv.org/abs/2604.08305

Code: github.com/AasimBinSaleem/HistDiT

Email: aasimr@edgehill.ac.uk

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