Multi-StyleGAN: Towards Image-Based Simulation of Time-Lapse Live-Cell Microscopy MCCAI2021

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- study living cells, with multiple aligned channels.





Paper (preprint): https://arxiv.org/pdf/2106.08285.pdf **Code**: https://git.rwth-aachen.de/bcs/projects/tp/multi-stylegan **Dataset**: https://tudatalib.ulb.tu-darmstadt.de/handle/tudatalib/2880 Project Page: https://christophreich1996.github.io/multi_stylegan/ ^{*} Christoph Reich and Tim Prangemeier – both authors contributed equally

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- Dual-styled-convolutional block enables high-quality synthesis of multi-domain images.
- The simulations realistically capture spatio-temporal organisation of multiple living yeast

amples capture underlying biophysical s realistically (baselines are unrealistic). Multi-StyleGAN samples							
00	80	80	80	00	60	00	
80			30				
50	80	80	80	80	200		
? •			*				
t ₂	t ₀	t ₁	t ₂	t ₀	t ₁	t ₂	
outperforms the baselines significantly.							
	FID ↓			FV	D ↓		
	BF		GFP	BF		GFP	
	33.37		207.84	4.4	6	30.16	
s. [3]	200.54		224.79	45.63		35.22	
S.	76.03		298.75	5 14.75		31.48	
ataset well and transitions smoothly supplementary video).							



