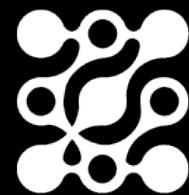


Image Translation

Shalin Mehta
Computational Microscopy Platform



**CHAN ZUCKERBERG
Biohub San Francisco**

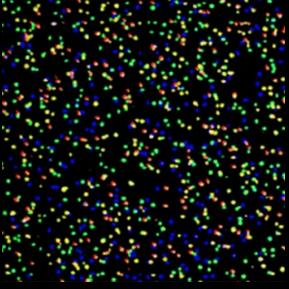


@mattersOfLight

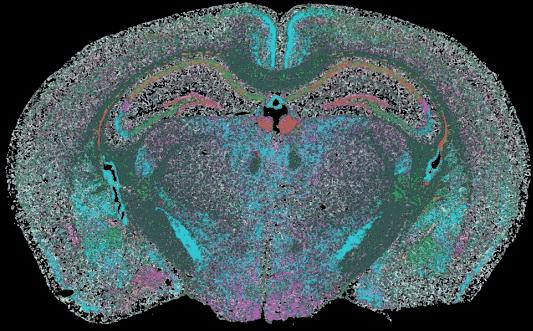
www.czbiohub.org/comp-micro/

DL@MBL
Aug 25, 2025

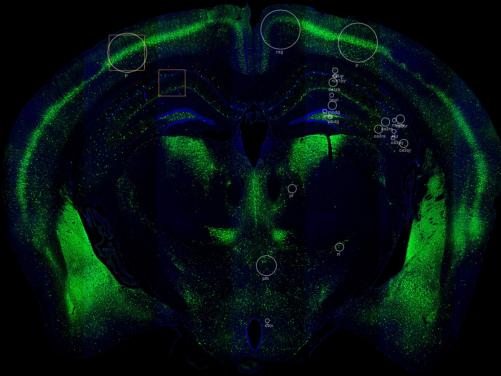
Imaging: foundational modality for biology



genome



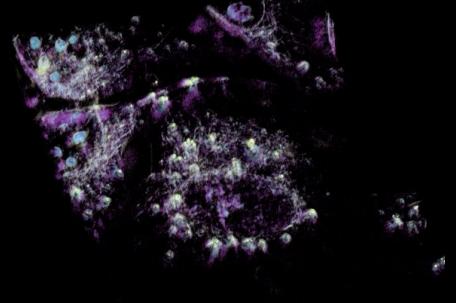
transcriptome



proteome

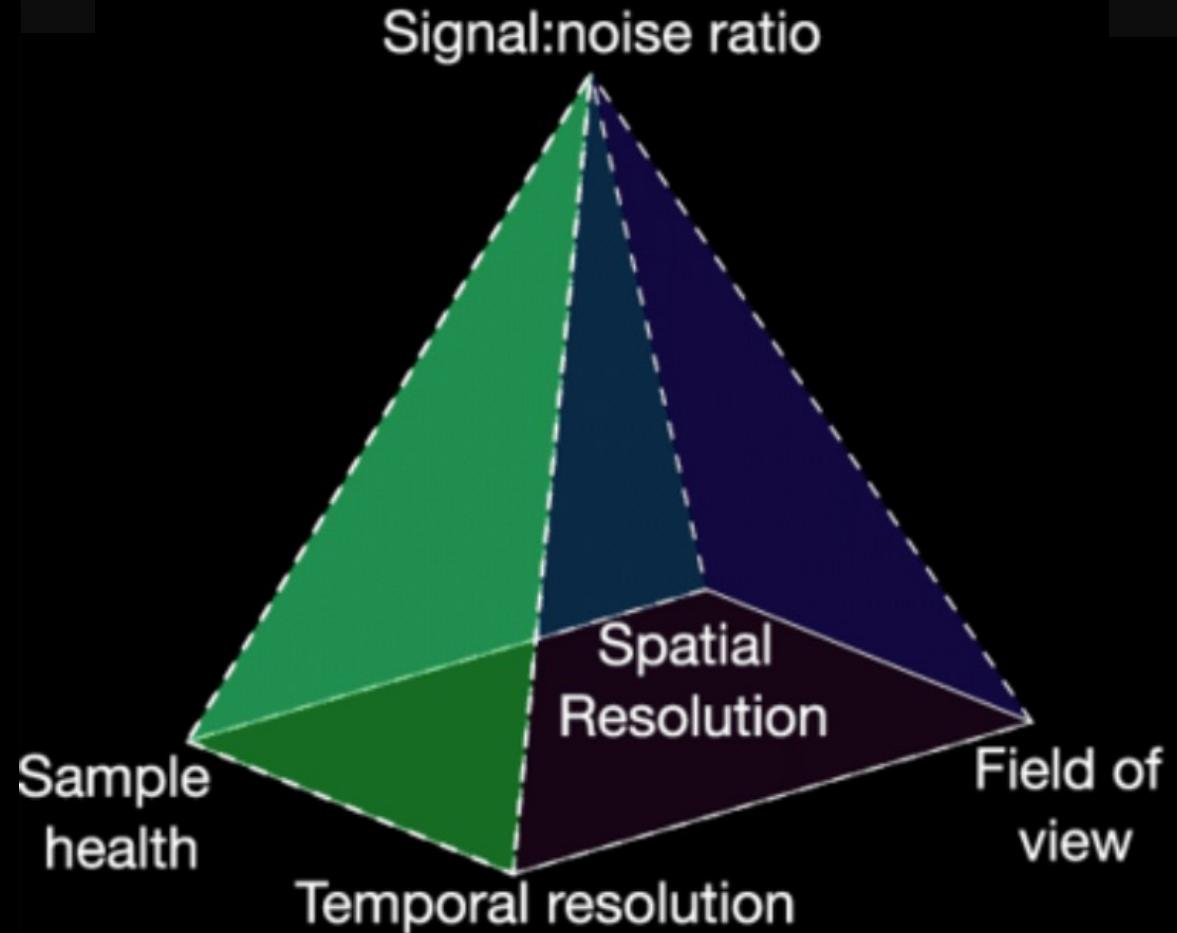


structure

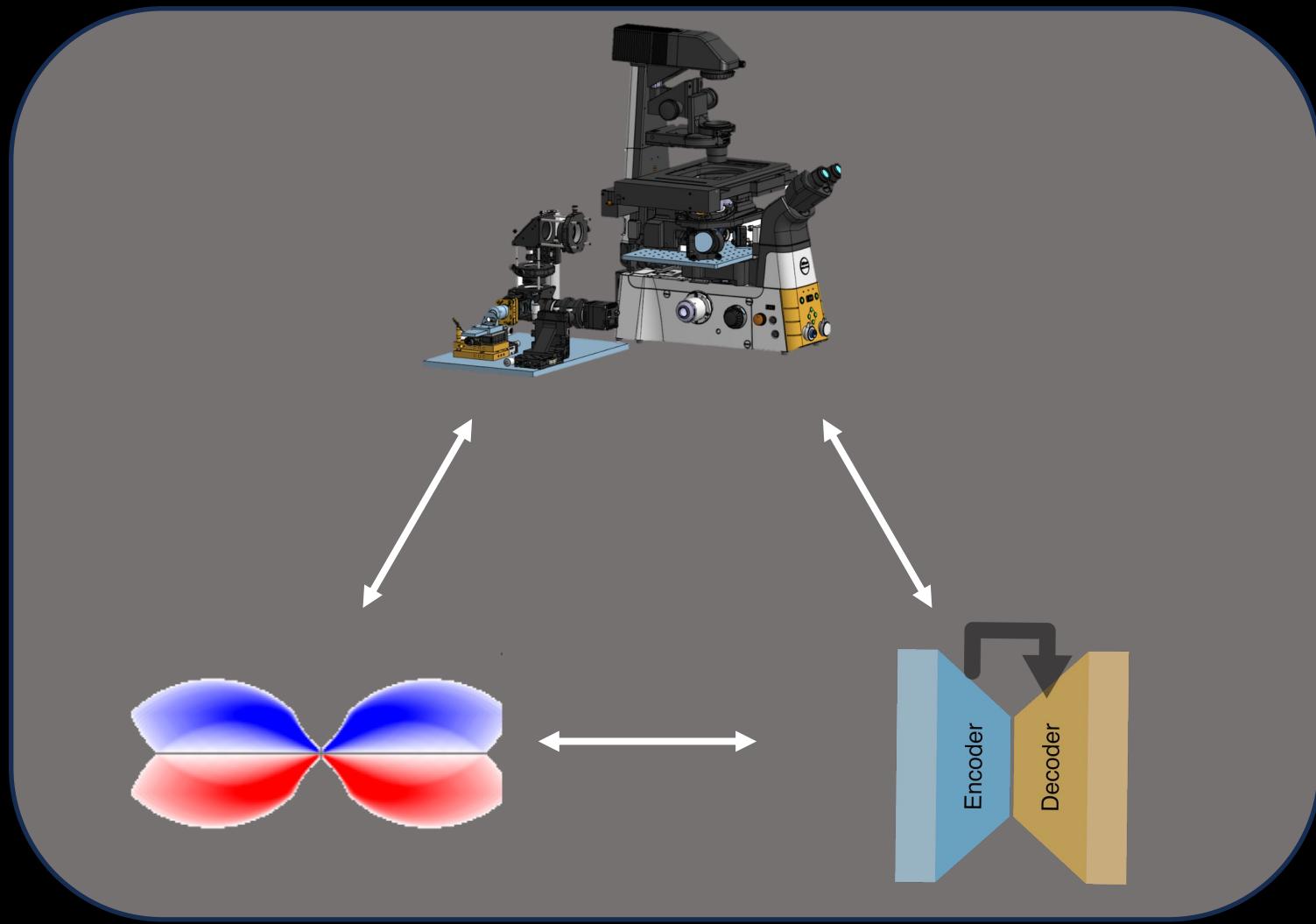


function

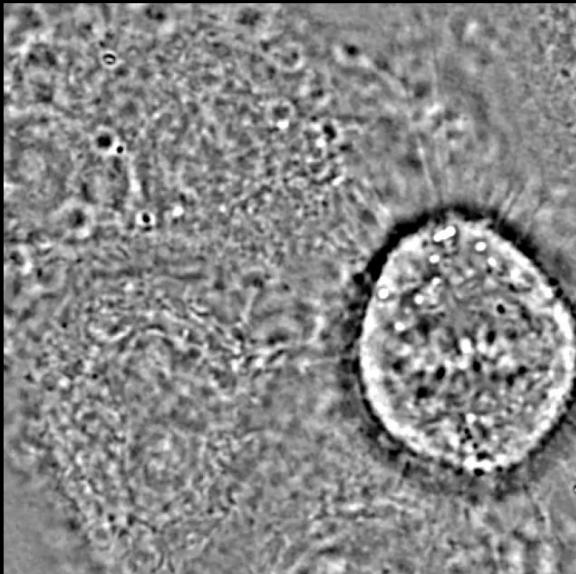
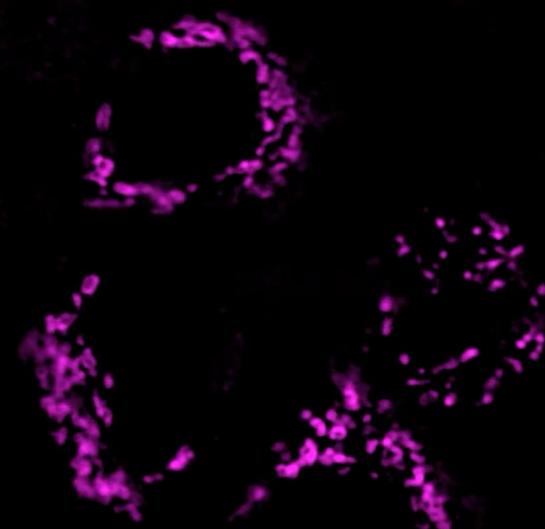
Multiplexing is hard just with optics!



Computational microscopy relaxes constraints

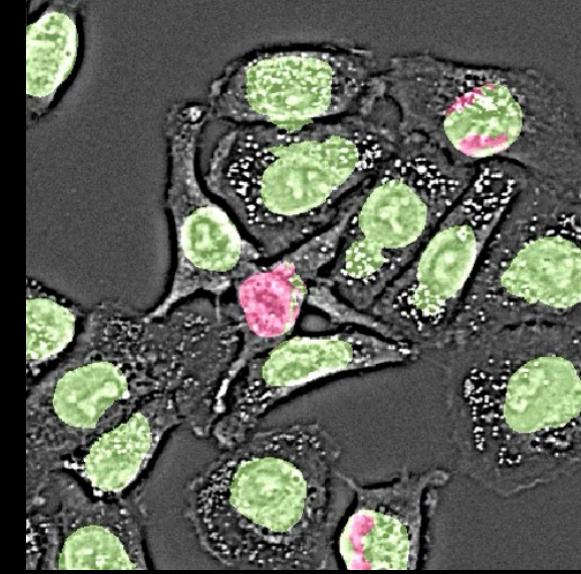


correlative live imaging



Guo et al., eLife 2020
Ivanov et al., Biomed. Opt. Exp., 2022
Yeh et al. Nature Methods, 2024
Ivanov et al., PNAS Nexus, 2024

learning cell dynamics



Guo et al., eLife 2020
Wu et al., Mol. Biol. Cell, 2022
Ivanov et al., PNAS Nexus, 2024
Liu et al., bioRxiv 2024



mehta-lab / `recOrder`



mehta-lab / `waveorder`



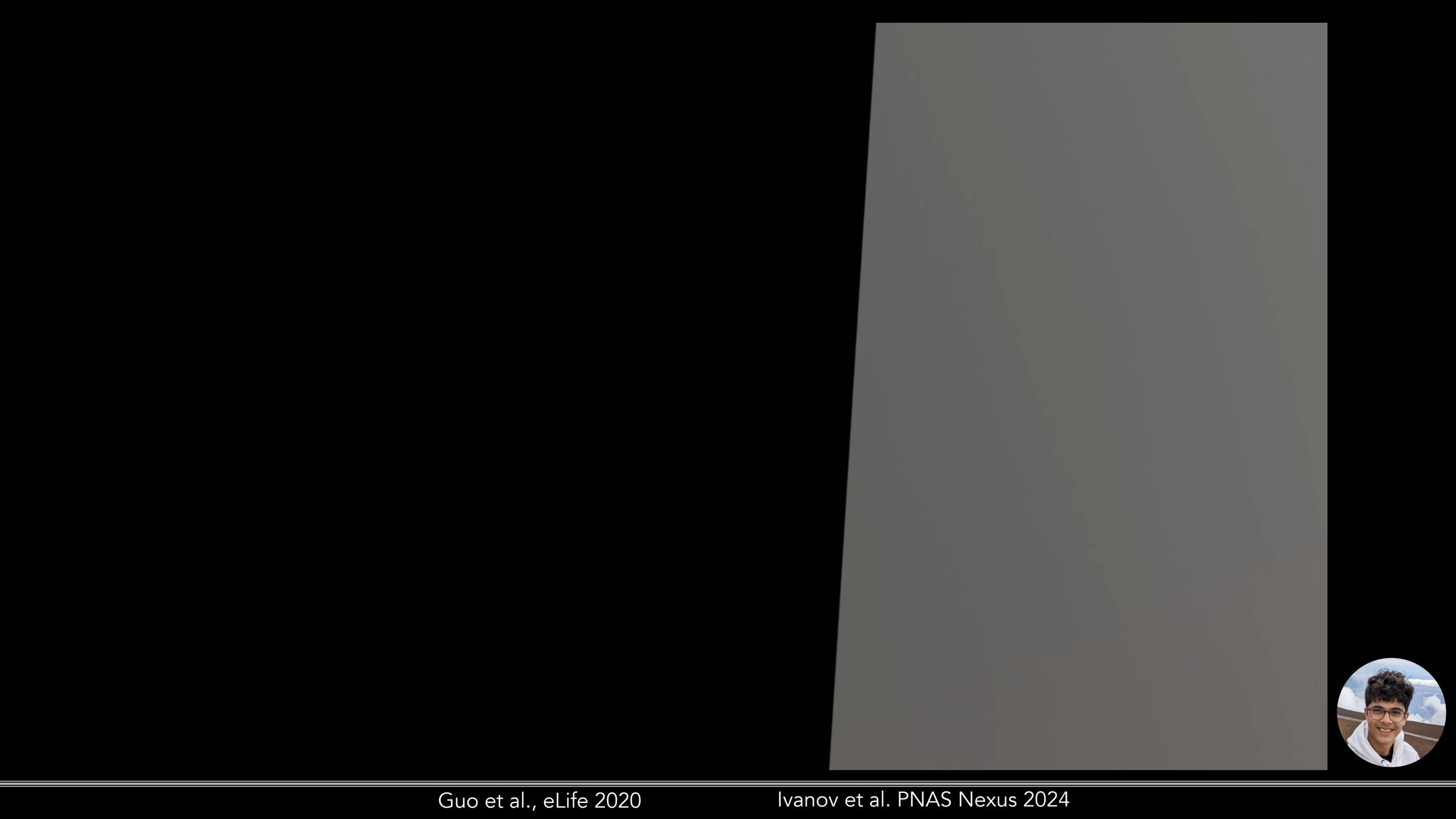
czbiohub-sf / `shrimPy`



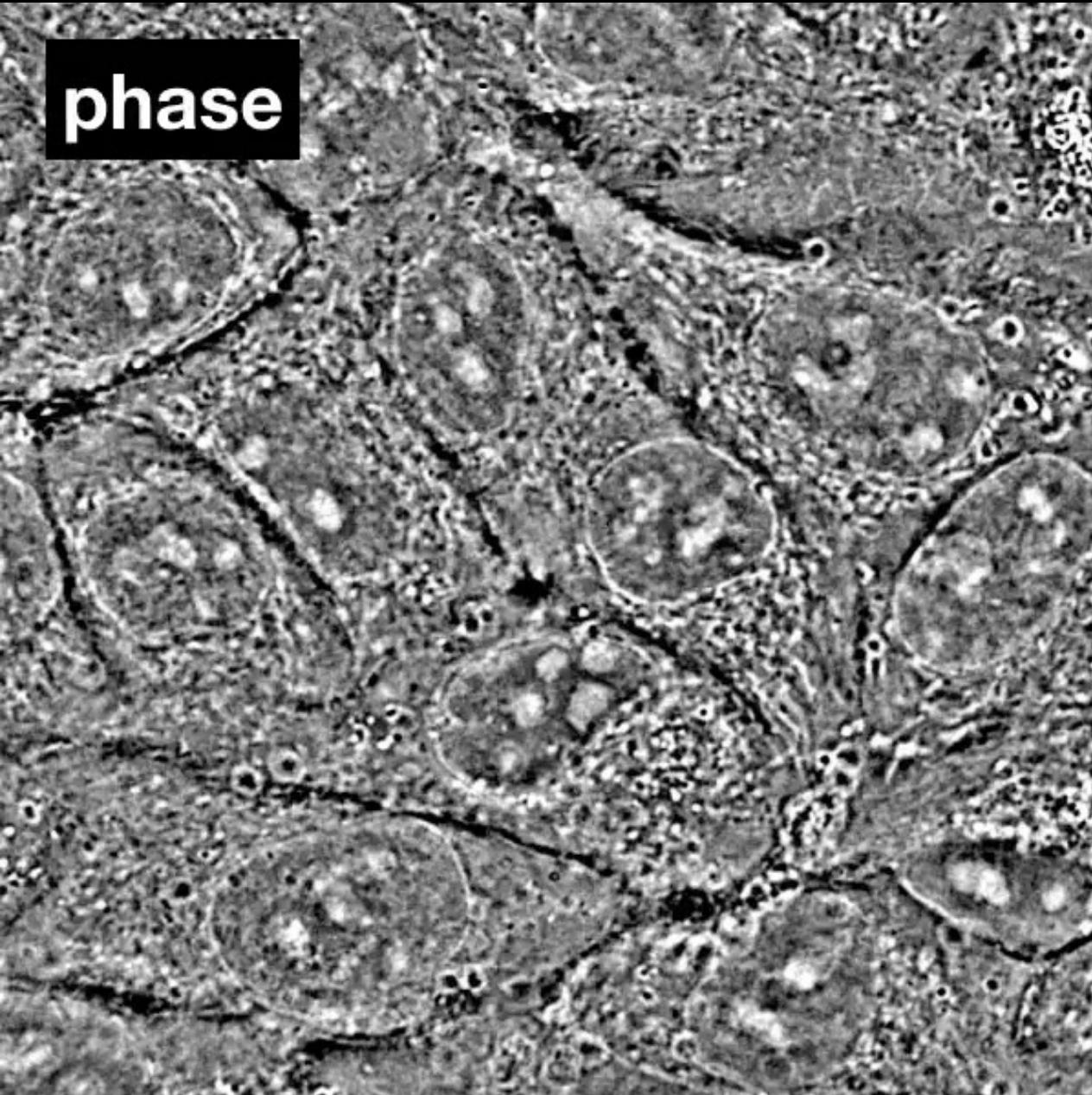
mehta-lab / `VisCy`



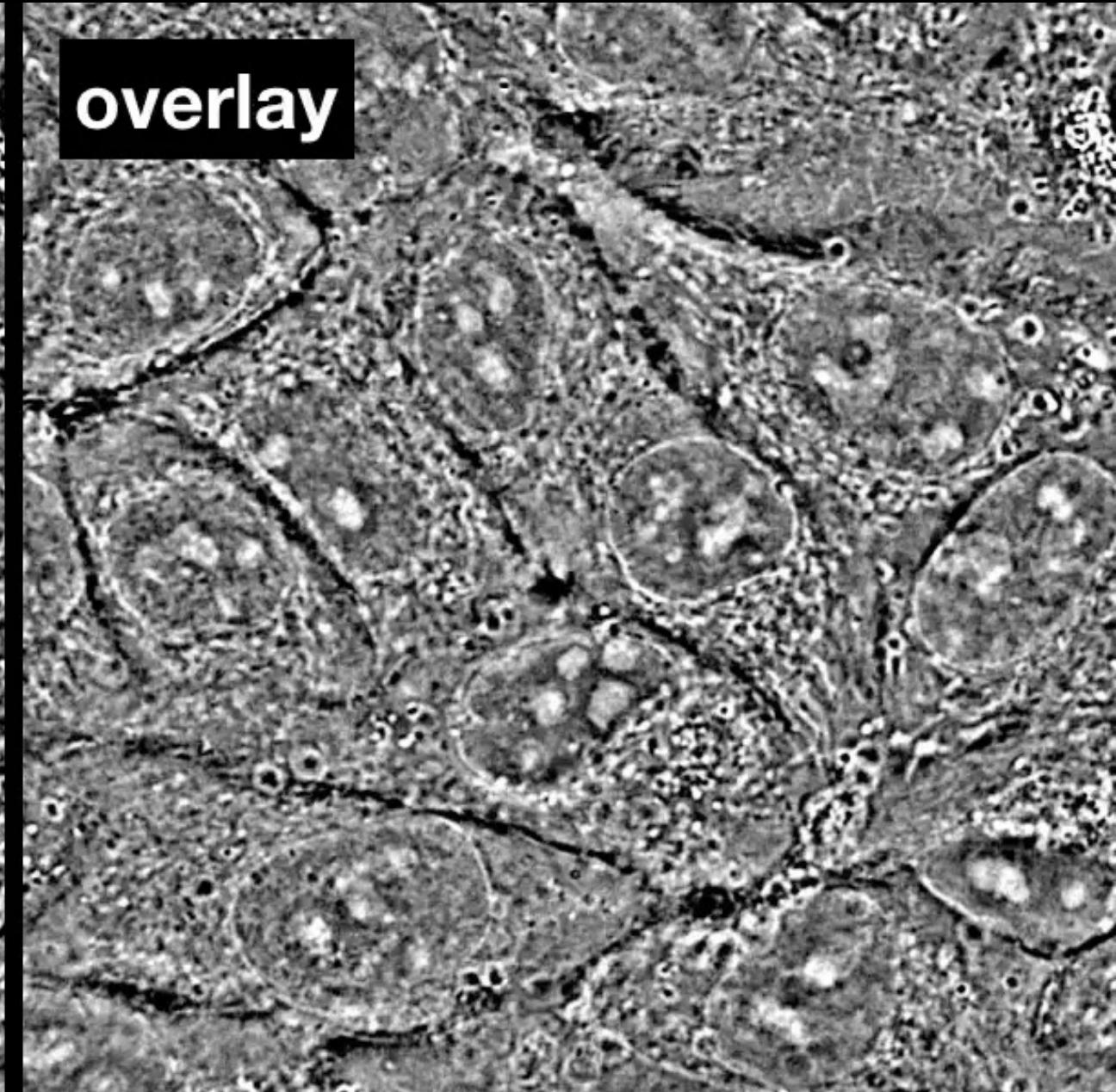
czbiohub-sf / `iohub`



phase

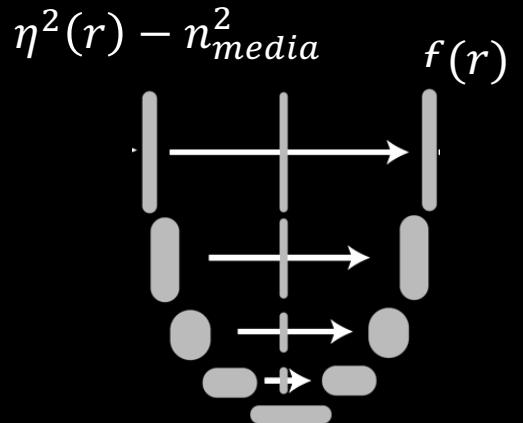
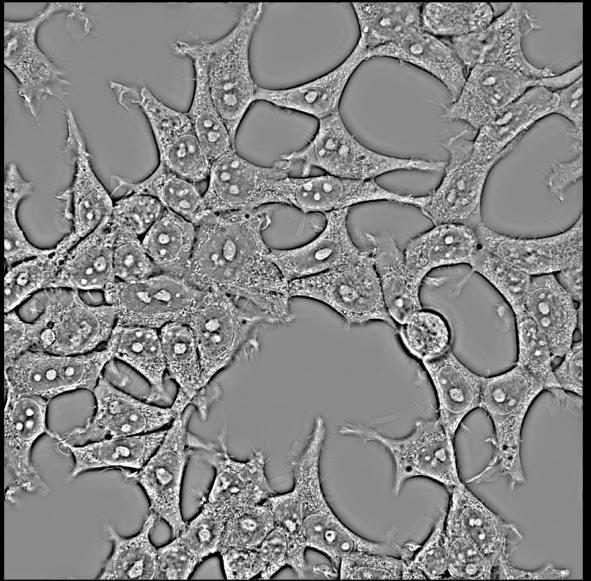


overlay

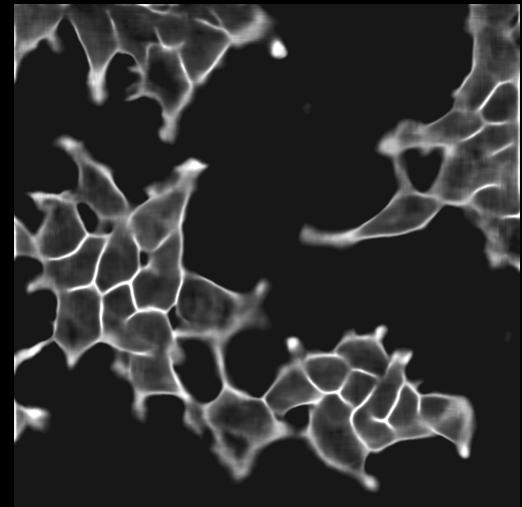
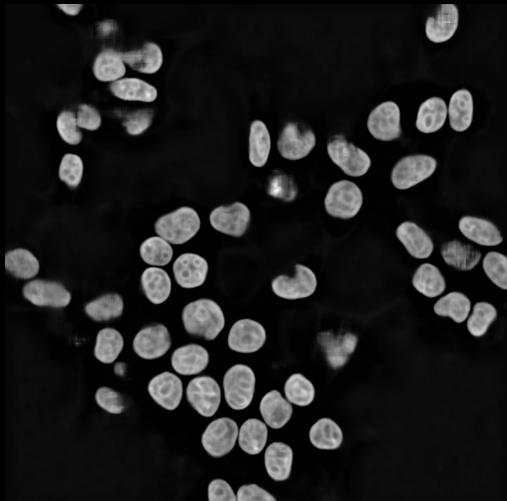


Robust virtual staining of landmarks

local permittivity



fluorophore density



Ziwen
Liu



Eduardo
Hirata-
Miyasaki



Soorya
Pradeep



Ivan
Ivanov



Talon
Chandler



Christian
Foley



Johanna
Rahm



Chad
Liu



Manu
Leonetti



Hunter
Woosley



Carolina
Arias

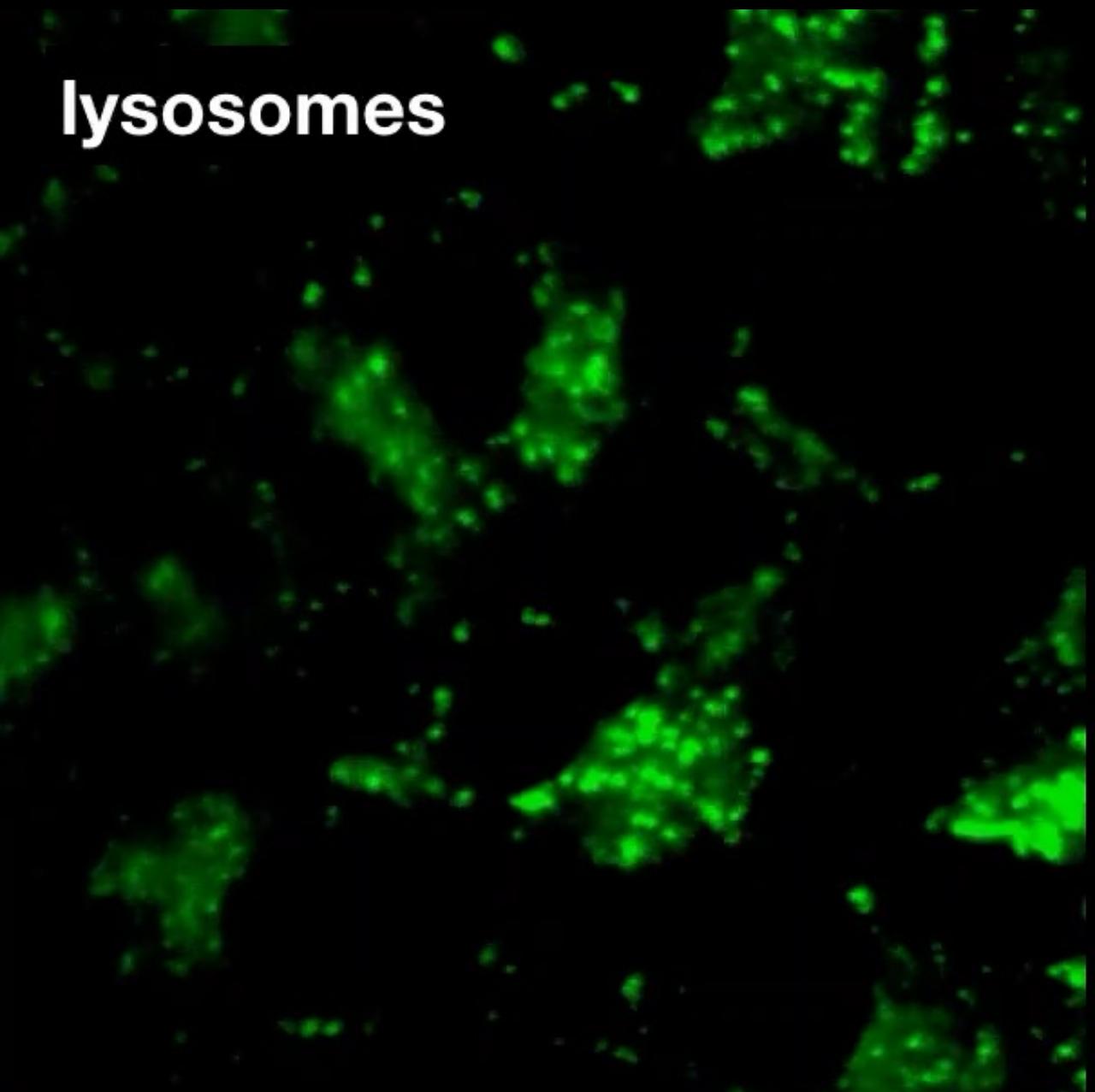


Ranen
Aviner

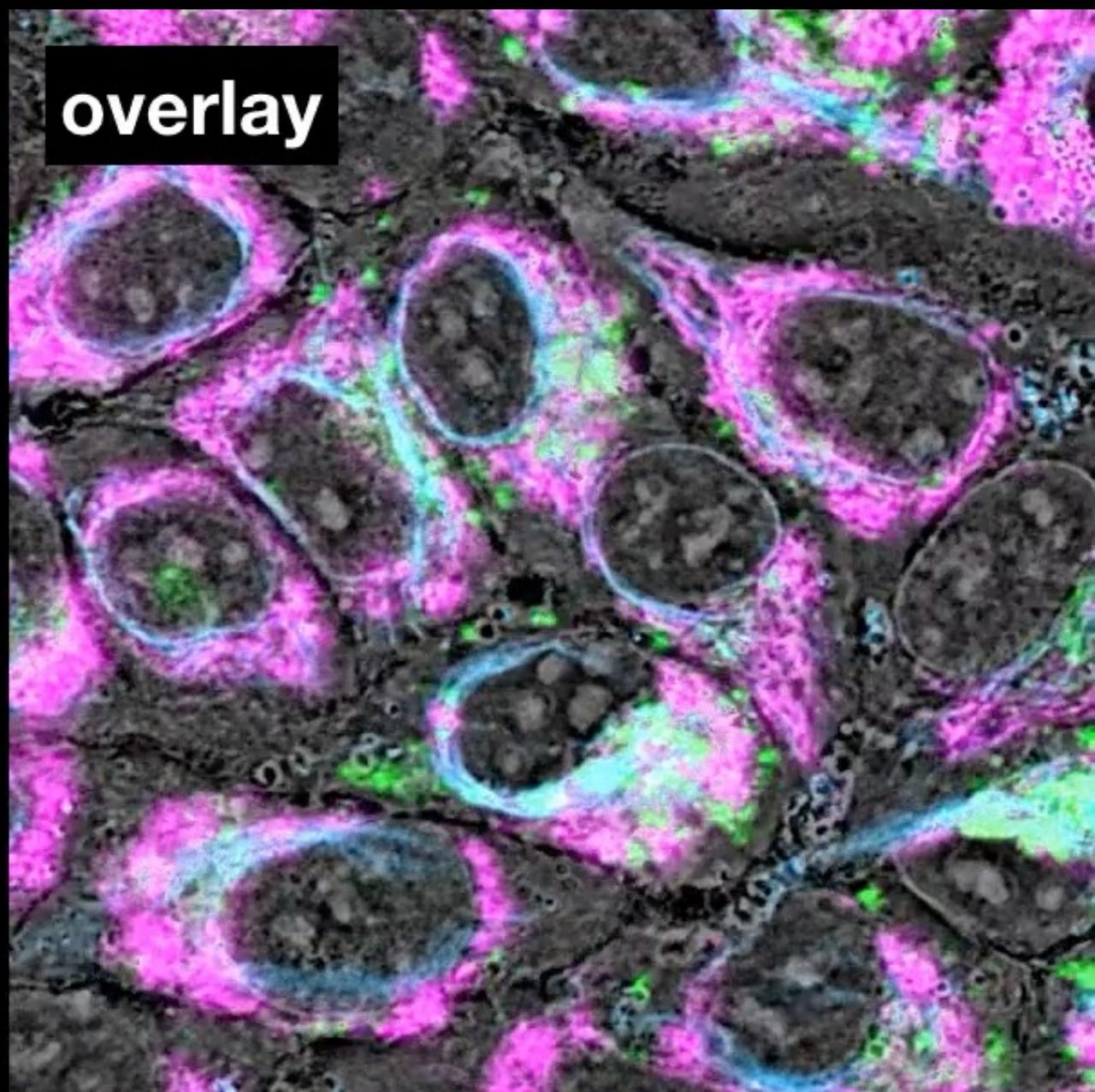


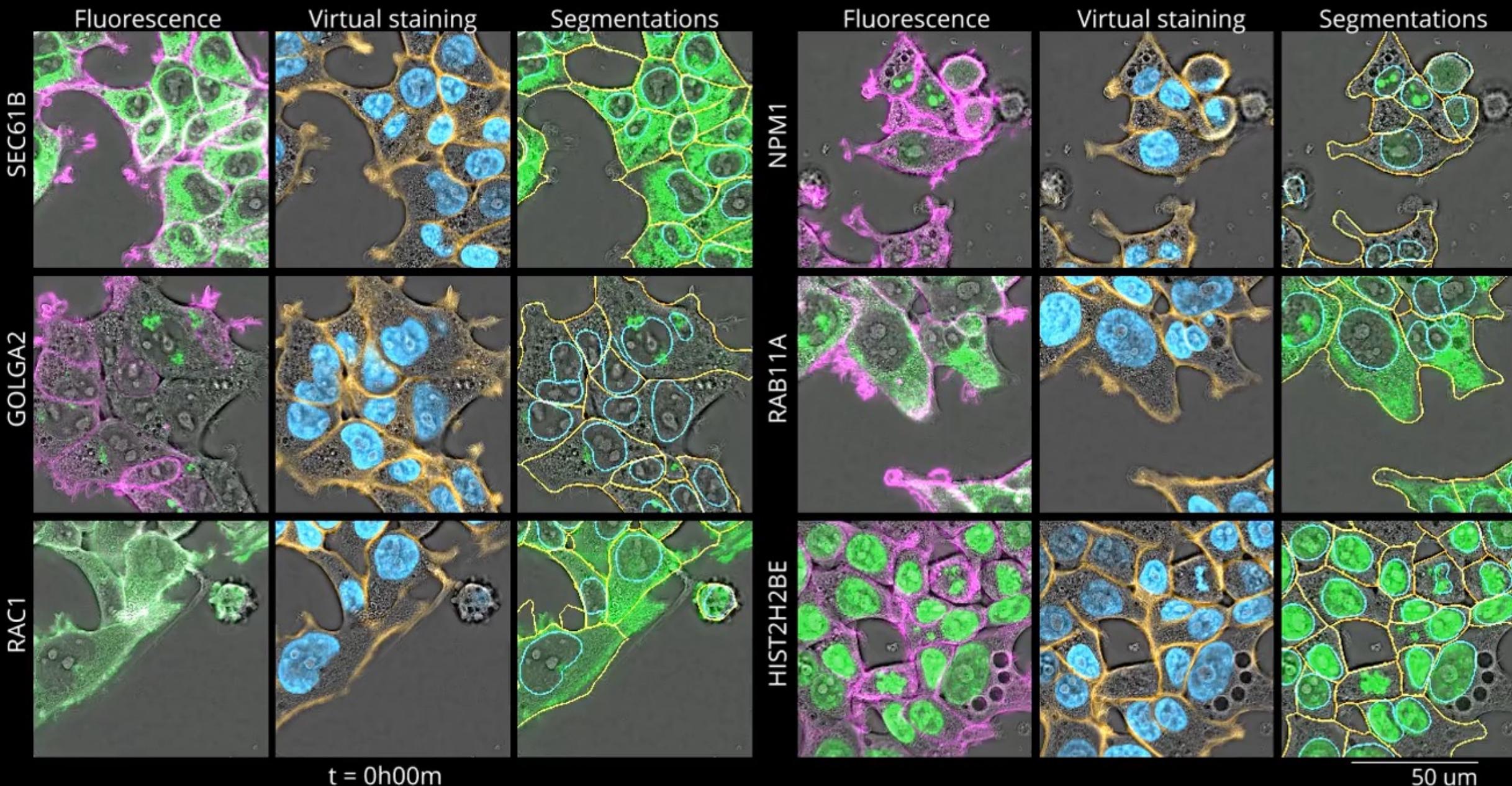
Adrian
Jacobo

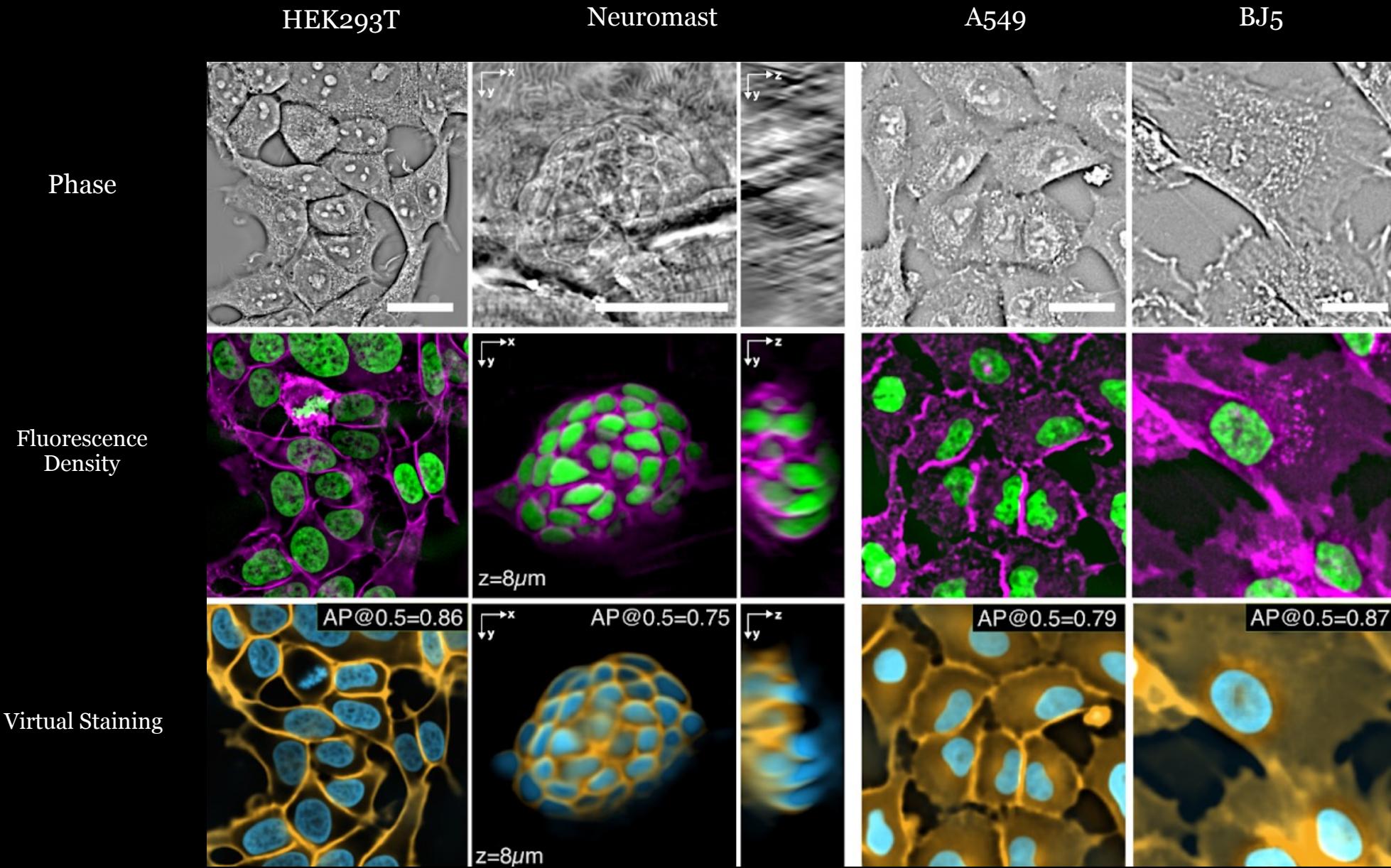
lysosomes



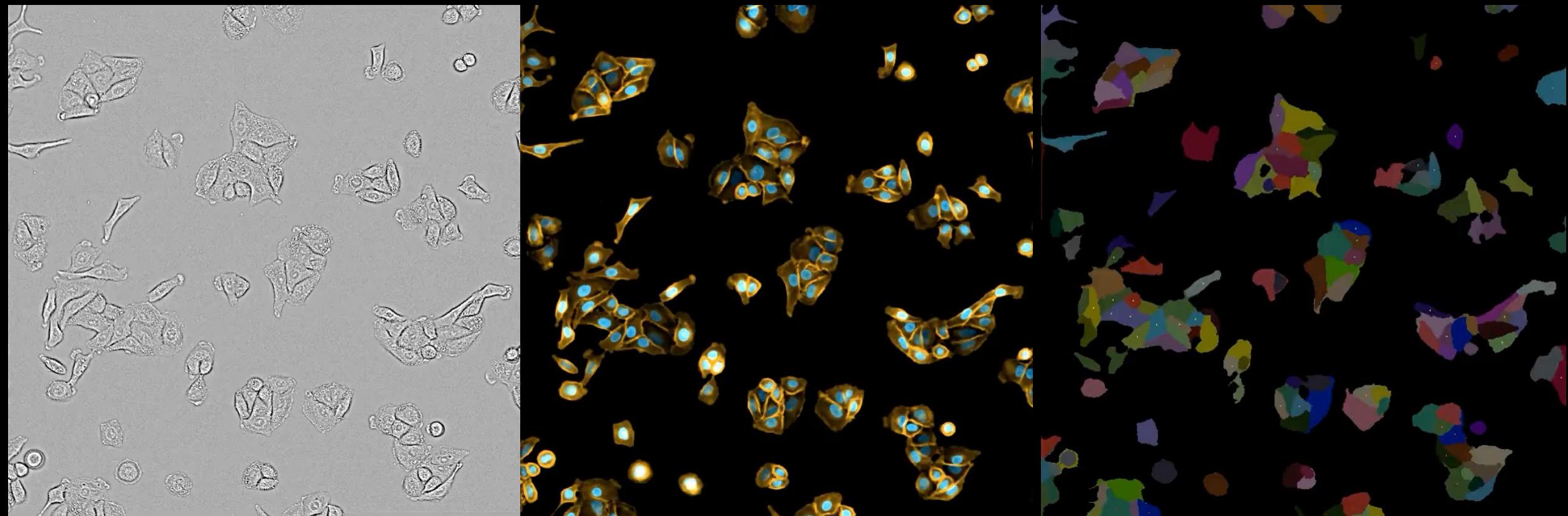
overlay





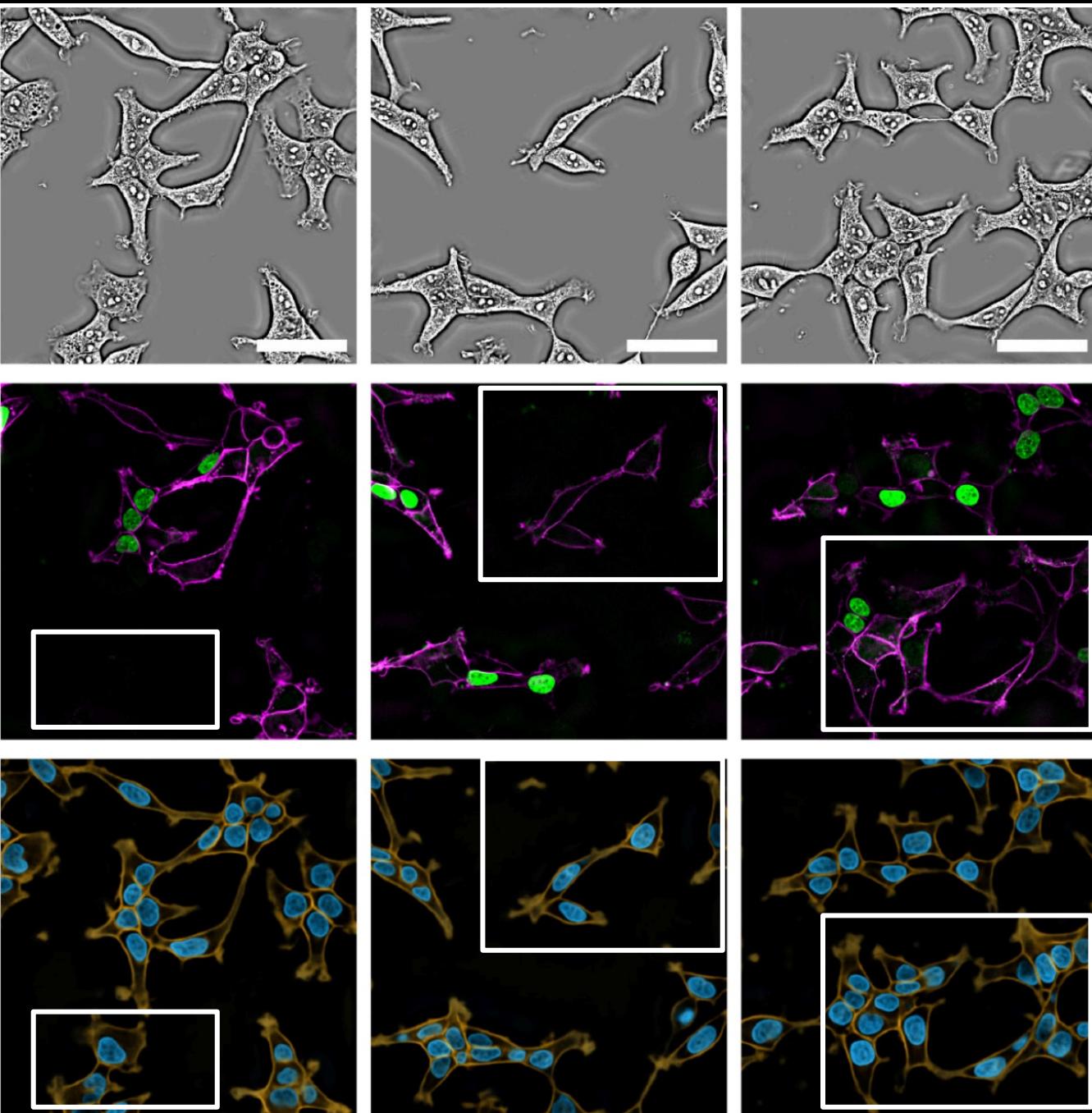


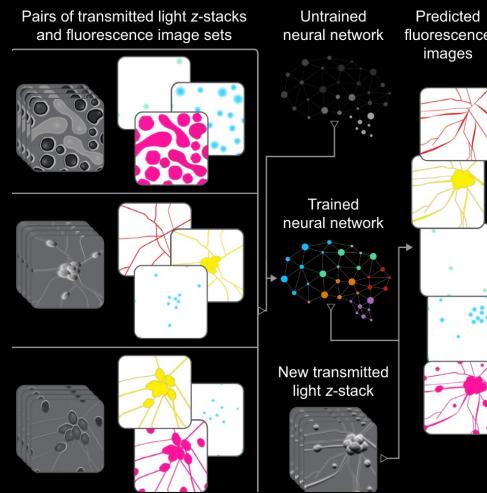
...across cell types



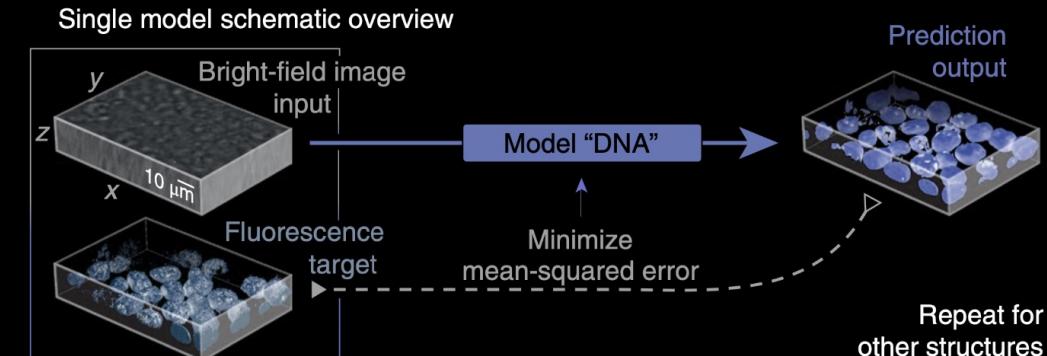
...across cell states

More robust than
experimental stain

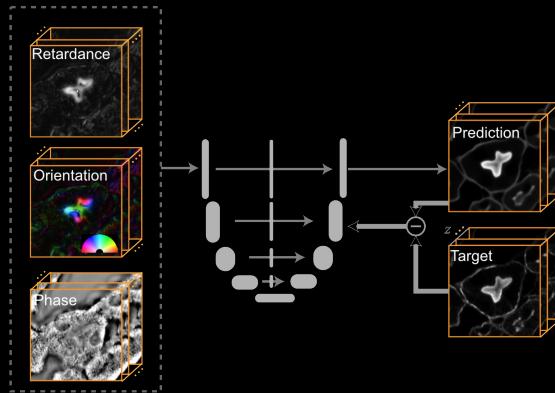




2D In silico labeling (Inception)
(Christiansen et al., Cell 2018)



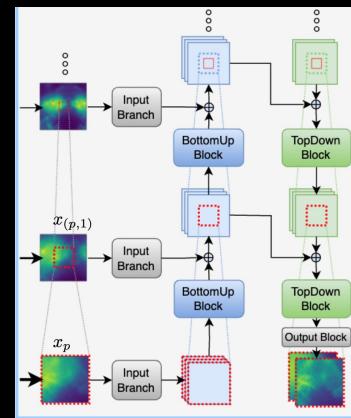
3D Label-free determination (3D U-Net)
(Ounkomol et al., Nature Methods, 2018)



3D image translation (2.5D U-Net)
(Guo et al., eLife, 2020;
Liu et al., bioRxiv, 2024)

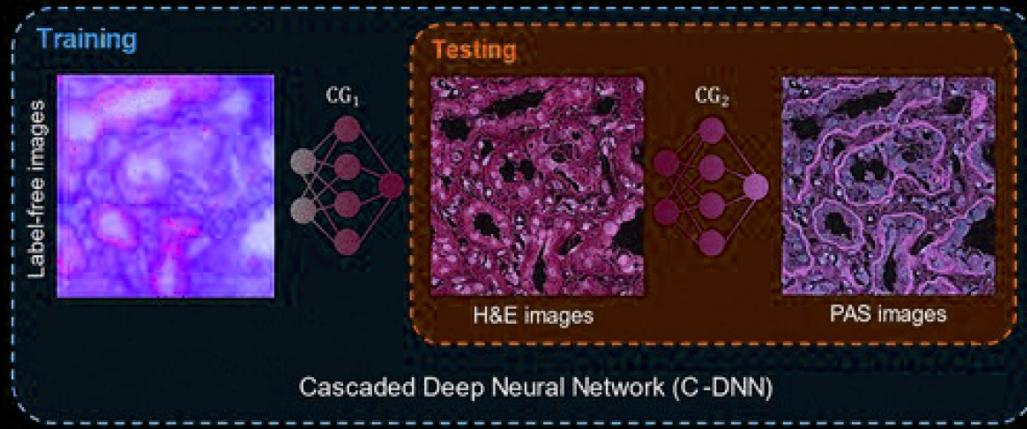


2D Pix2Pix (conditional GAN)
(Zhu et al., ICCV 2017; Isola et al., CVPR 2017; Tonks arXiv 2024)

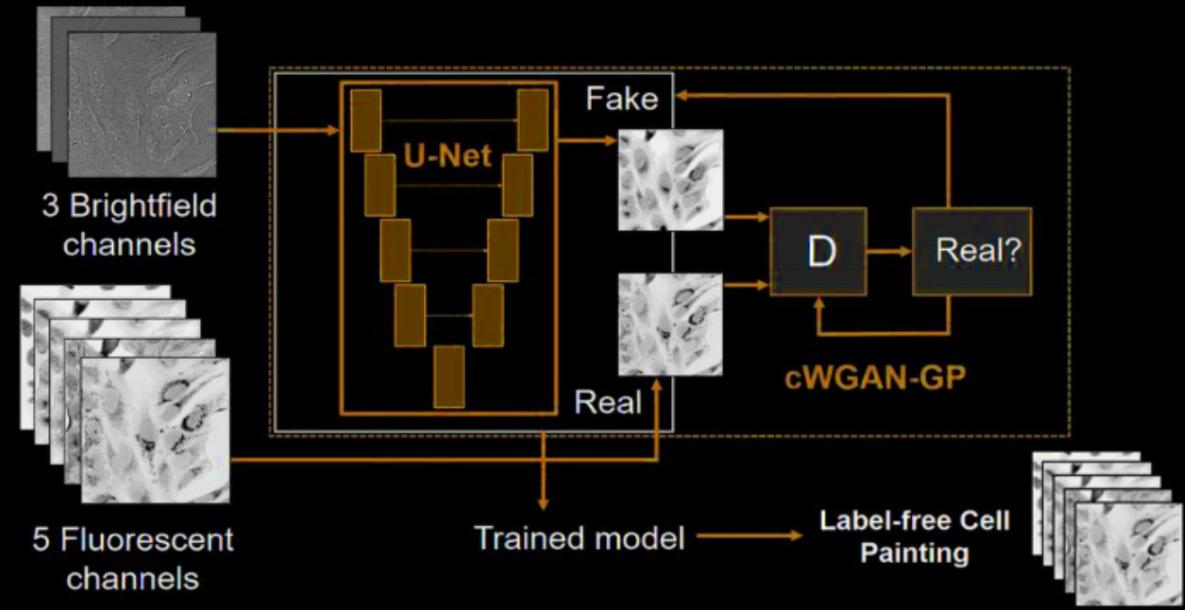


2D image splitting (HVAE)
Ashesh ICCV 2023

virtual histology & live cell painting

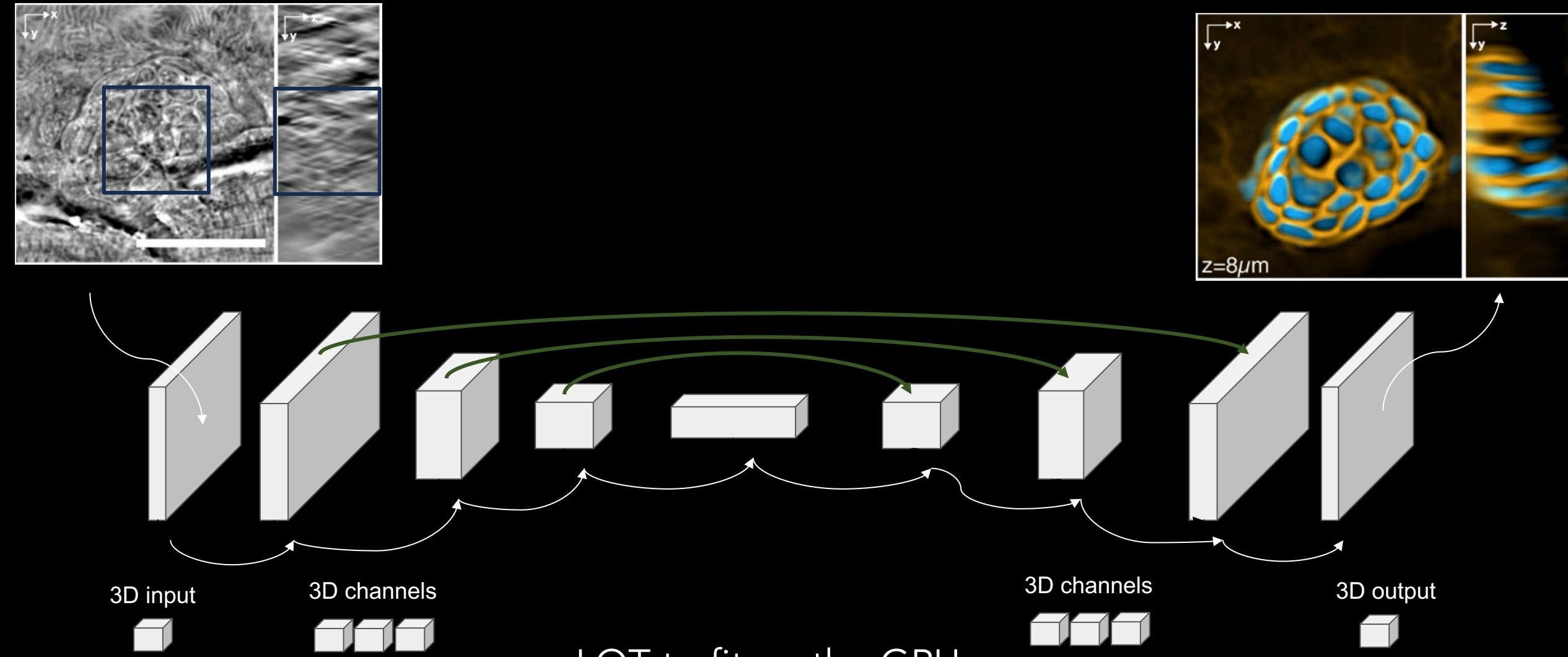


Yang...Ozcan ACS Photonics, 2022

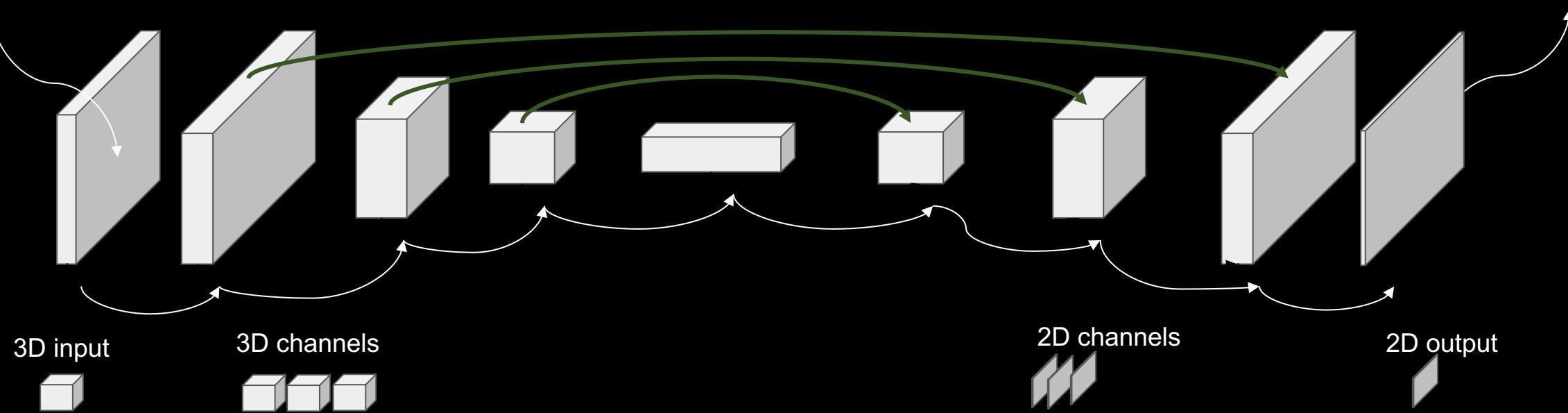
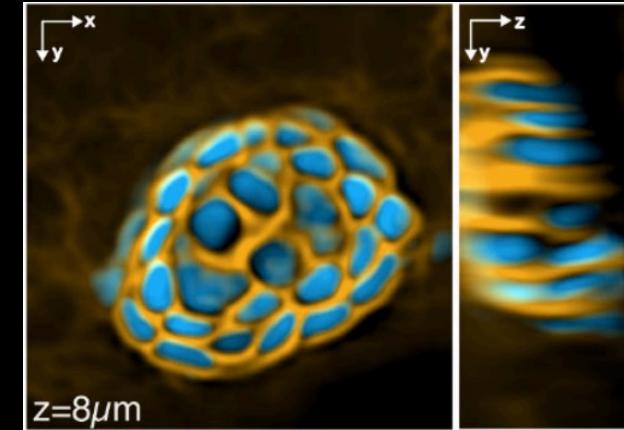
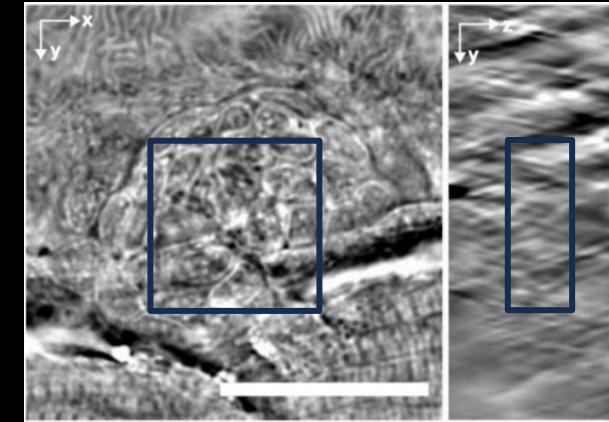


Oscar...Wang (AstraZeneca R&D), Scientific Reports, 2022

3D U-Net



2.5D U-Net



UNeXt

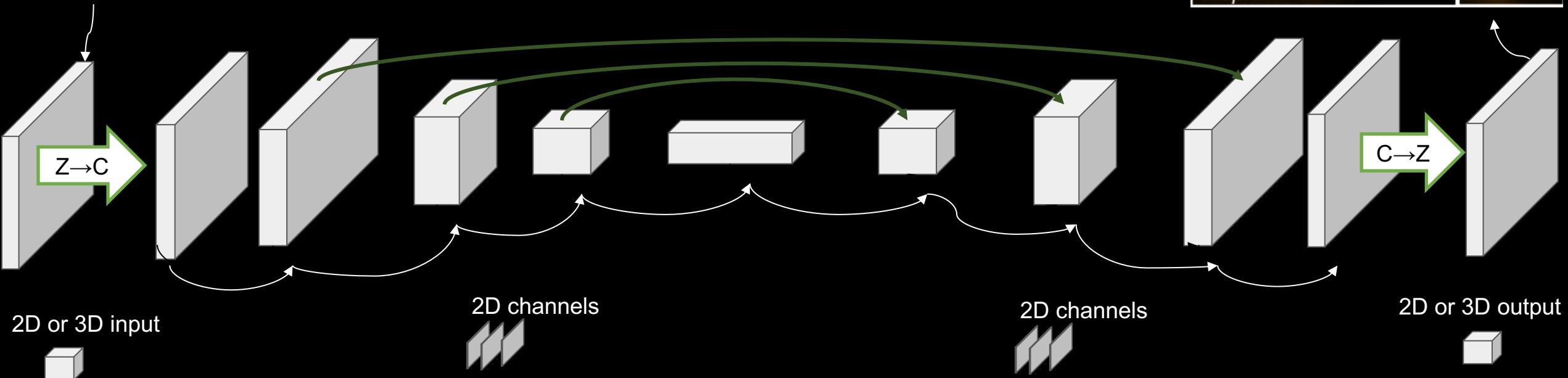
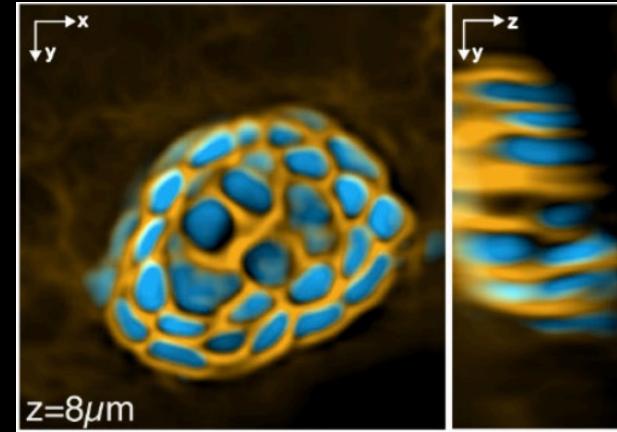
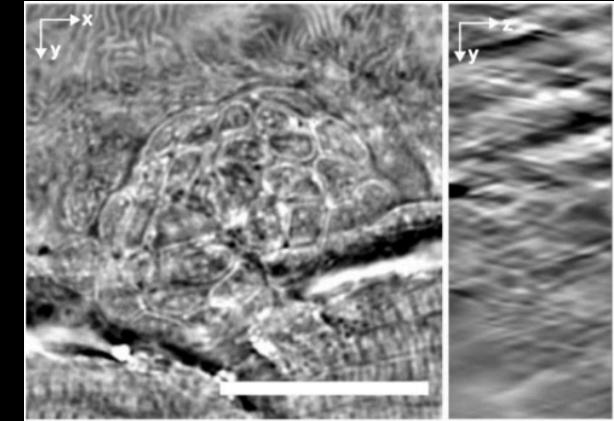
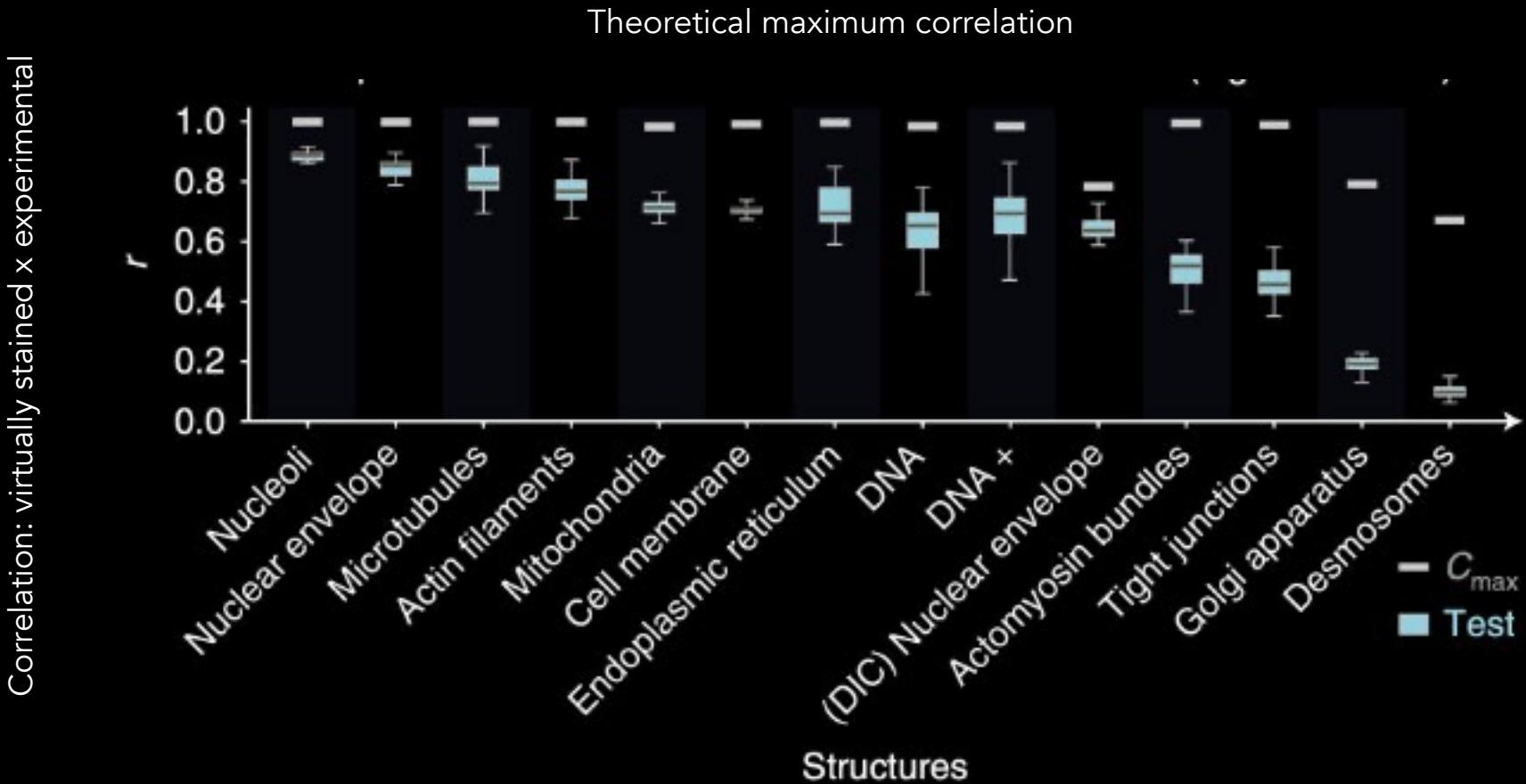


Table 1**In silico labeling in selected publications.**

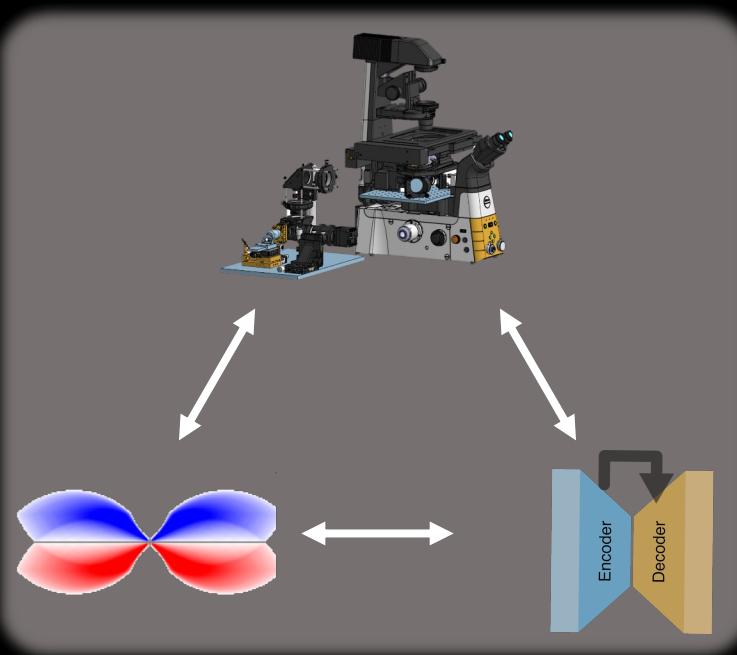
Reference	Paper title	Label-free modality	#Dimensions	Cell types	Subcellular structures	Term used	Deep learning network architecture	Evaluation metrics
Ounkomol et al., 2018 [11]	Label-free prediction of three-dimensional fluorescence images from transmitted light microscopy	bright-field, DIC	3	HEK-293, HT-1080, hiPSC	(+) microtubules, nuclear envelope, nucleoli (-) DNA, ER, cell membrane, actin filaments, mitochondria (-) desmosomes, actomyosin bundles, Golgi apparatus, tight junctions	label-free prediction	U-Net	PCC
Christiansen et al., 2018 [12]	In silico labeling: Predicting fluorescent labels in unlabeled images	bright-field, phase contrast, DIC	3	human motor neurons from iPSCs, primary rat cortical cultures, human breast cancer line	nuclei + neurons/dendrites/dead cells/membrane/axons	In silico labeling	multiscale sub-networks (towers) and one or more pixel-distribution-valued predictors (heads)	PCC, ▲ (see legend)
LaChance and Cohen 2020 [13]	Practical fluorescence reconstruction microscopy for large samples and low-magnification imaging	phase contrast, DIC	2	MDCK, KC, HUVEC	(+) nuclei (-) actin, cell–cell junctions	FRM (Fluorescence reconstruction microscopy)	U-Net	PCC, P, ▲▲ (see legend)
Wang et al., 2021 [14]	Global voxel transformer networks for augmented microscopy	Same as Ounkomol et al., 2018				augmented microscopy	U-Net with attention	PCC
Cheng et al., 2021 [15]	Single-cell cytometry via multiplexed fluorescence prediction by label-free reflectance microscopy	multimodal reflectance microscope	2	HeLa	(+) DNA, actin, endosome (-) Golgi apparatus	digital labeling	U-Net	PCC
Jo et al., 2021 [16]	Label-free multiplexed microtomography of	QPI	3	NIH3T3, CRL-1658, COS-7, HEK293, HeLa, astrocytes	(+) actin, mitochondria, lipid droplets, plasma	in silico labeling	NAS, U-Net	PSNR, PCC, SSIM

consistency of label-free encoding



Ounkomol et al., Nature Methods, 2018

Neural networks often don't generalize across imaging systems or cell types ...



Physics-based inverse algorithms:

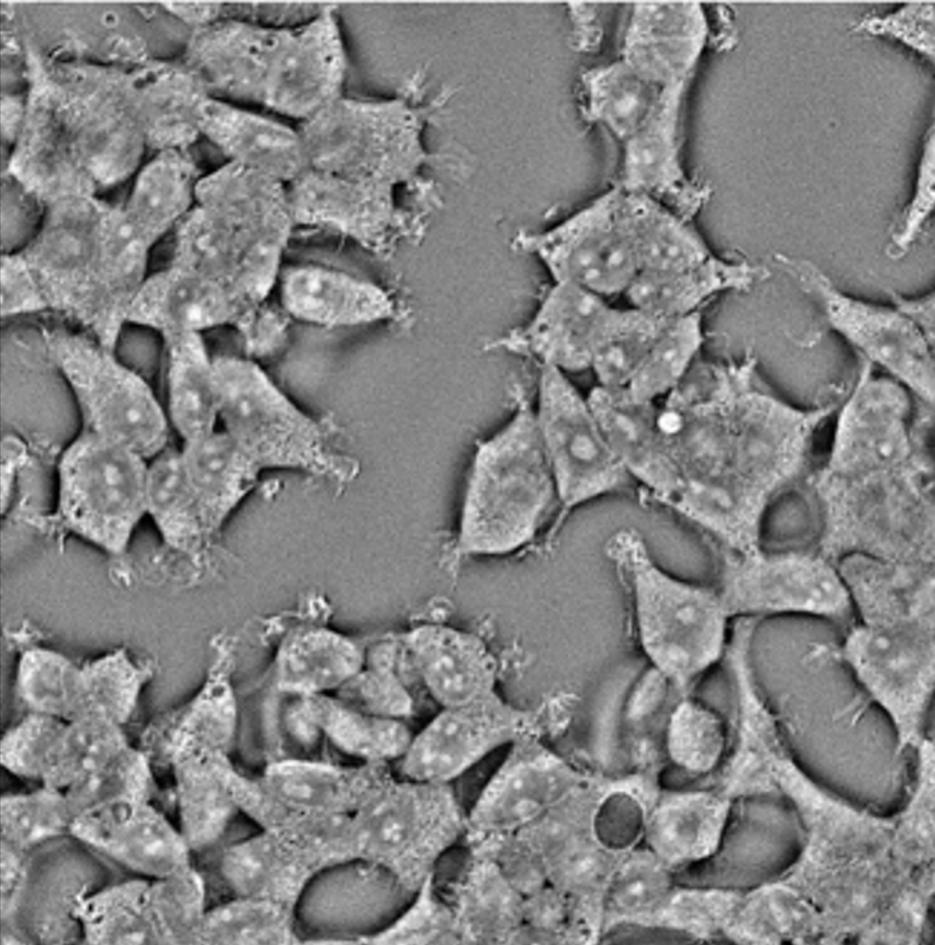
- Cells and tissues that scatter once
- Diverse imaging configuration

Learned models:

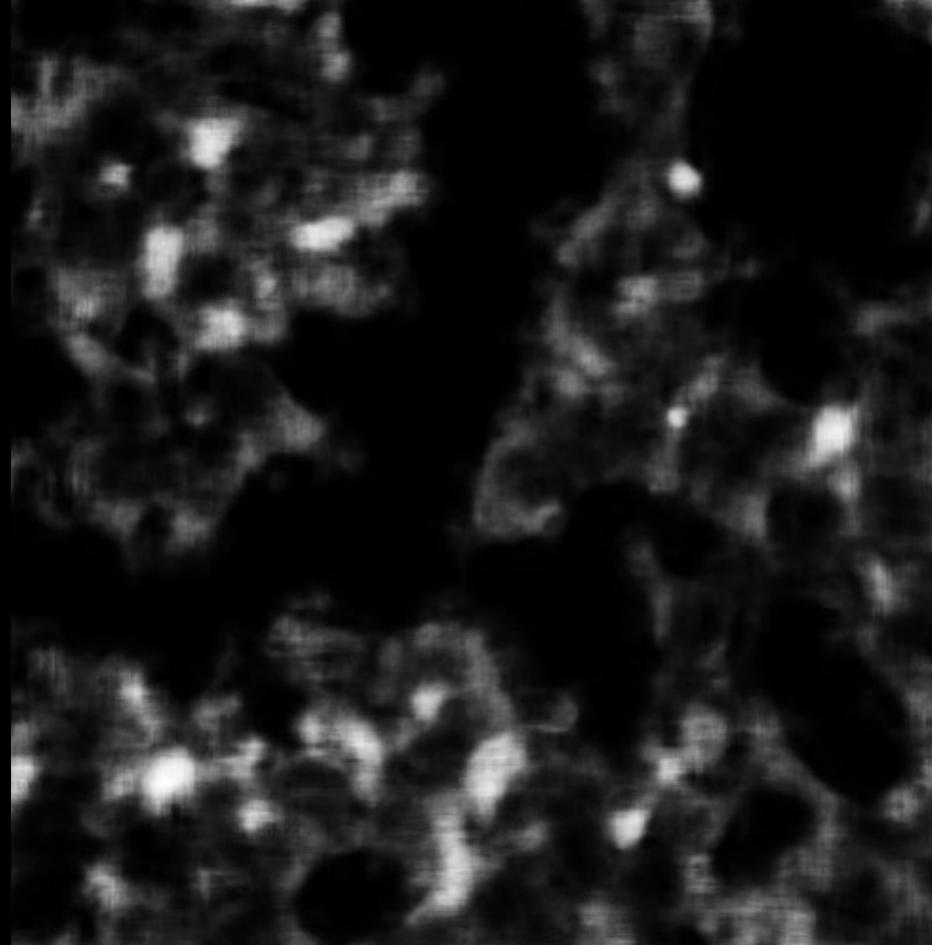
- Cells and tissues *in distro*
- Specific imaging configurations

Generalization across imaging conditions

Different noise than training data

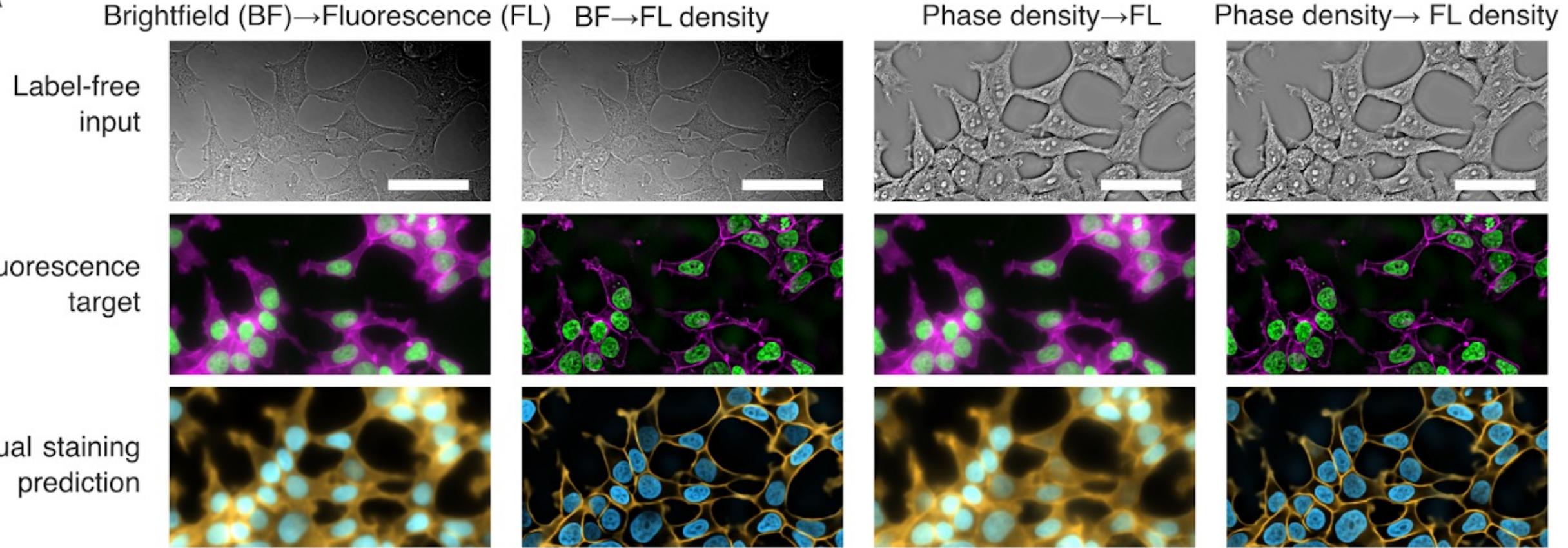


Prediction of nuclei

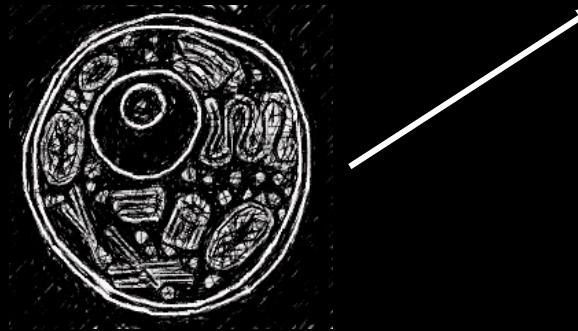
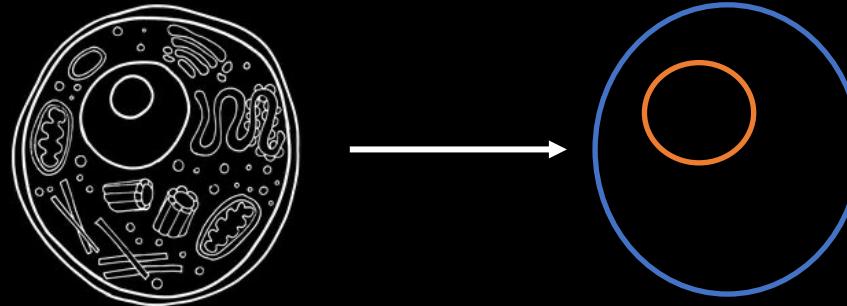


Towards robust virtual staining ...

deconvolution improves accuracy

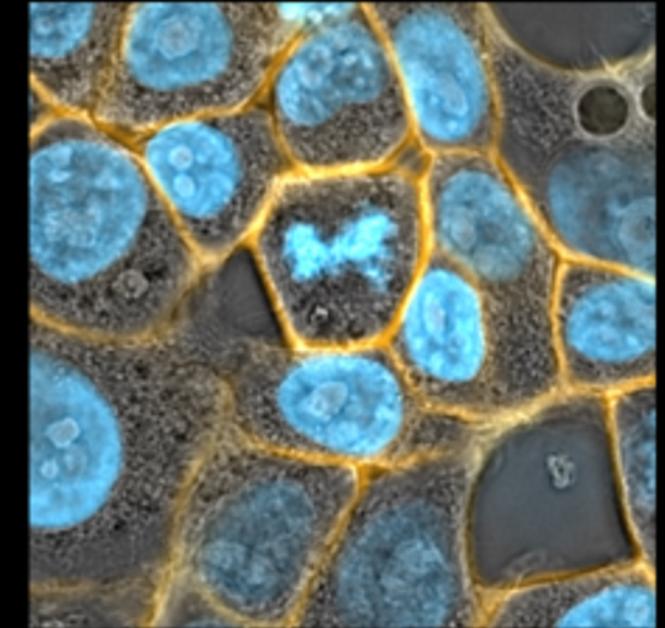
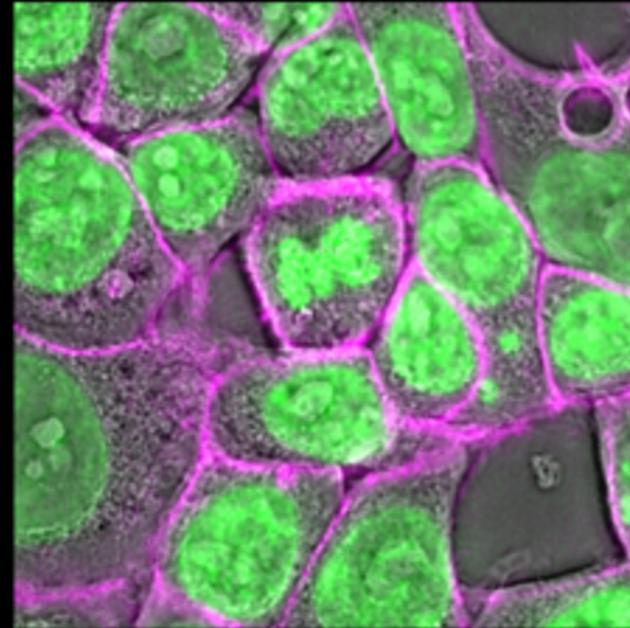


augmentations to the rescue!

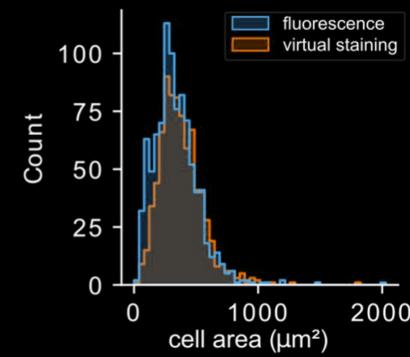
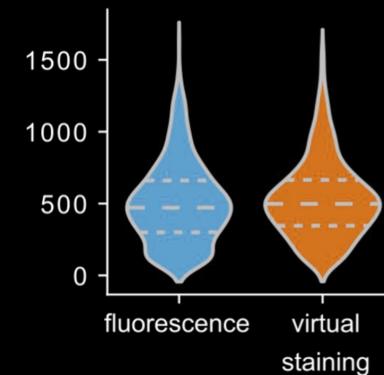


generalization across microscopes

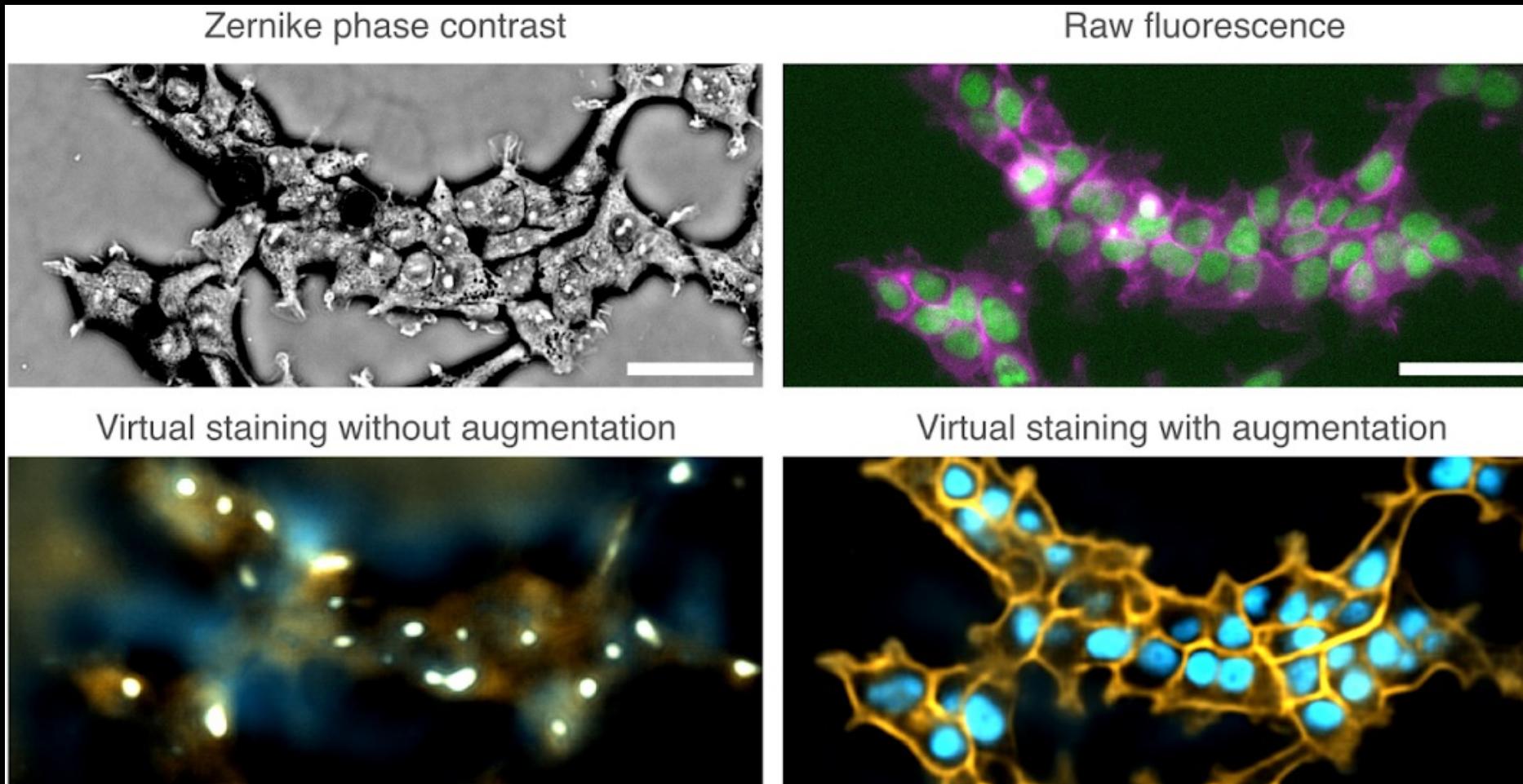
Training data from scope 1



Test data from scope 2

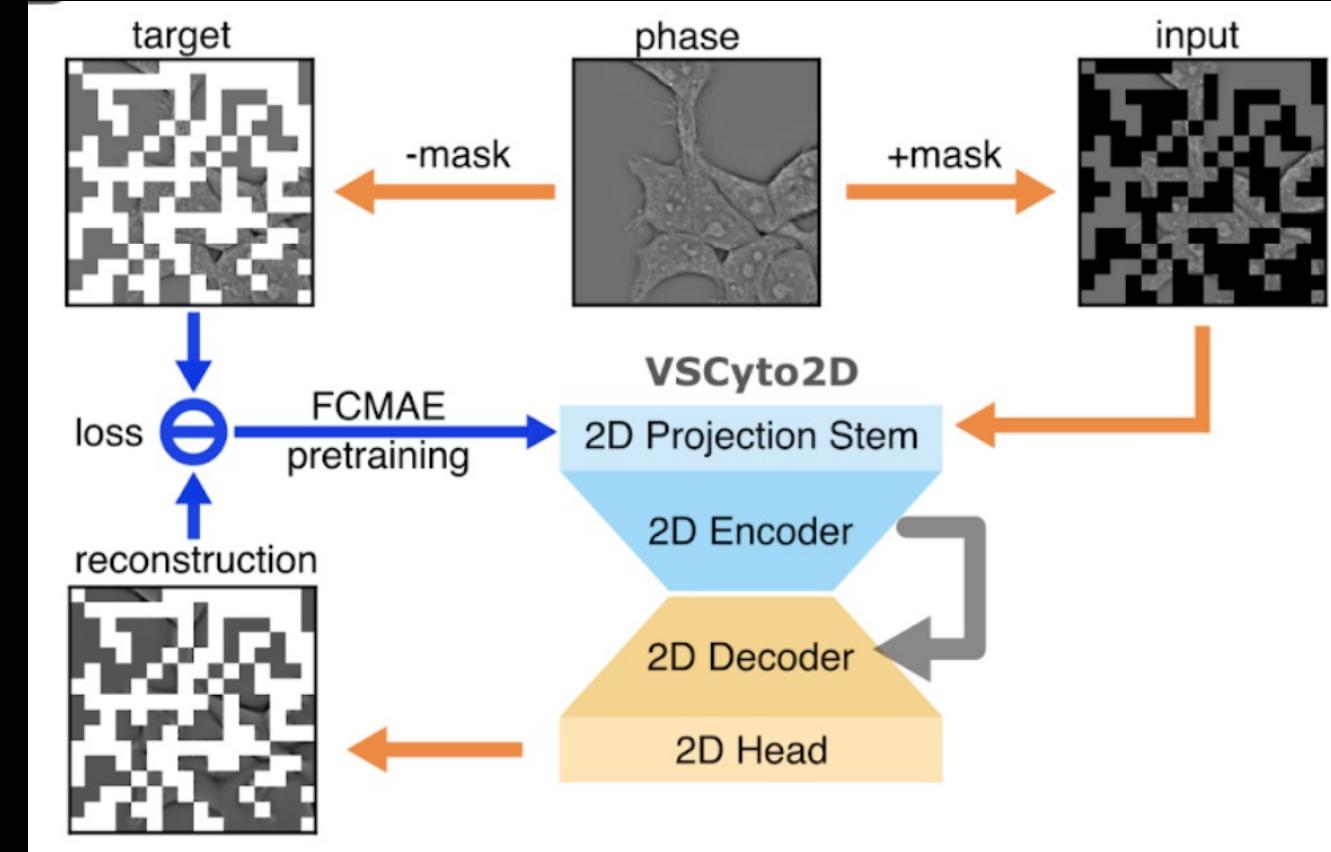
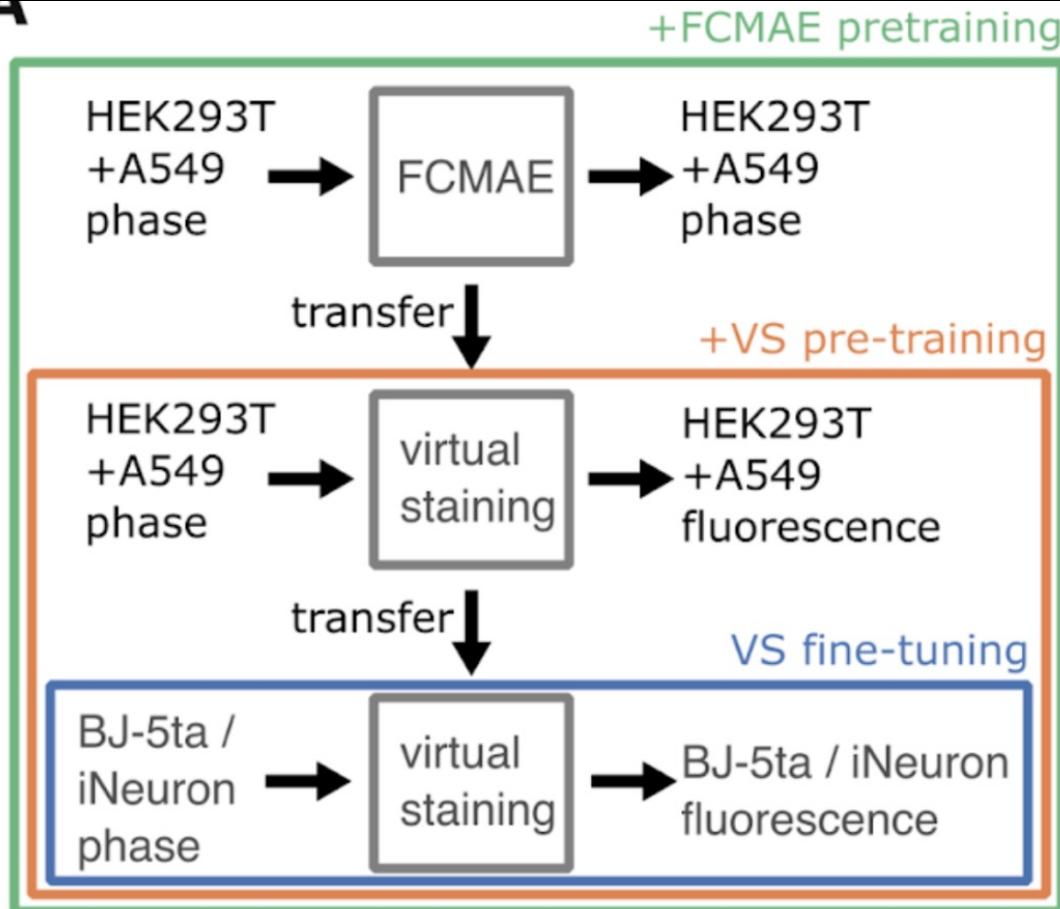


Zero-shot generalization to new contrast



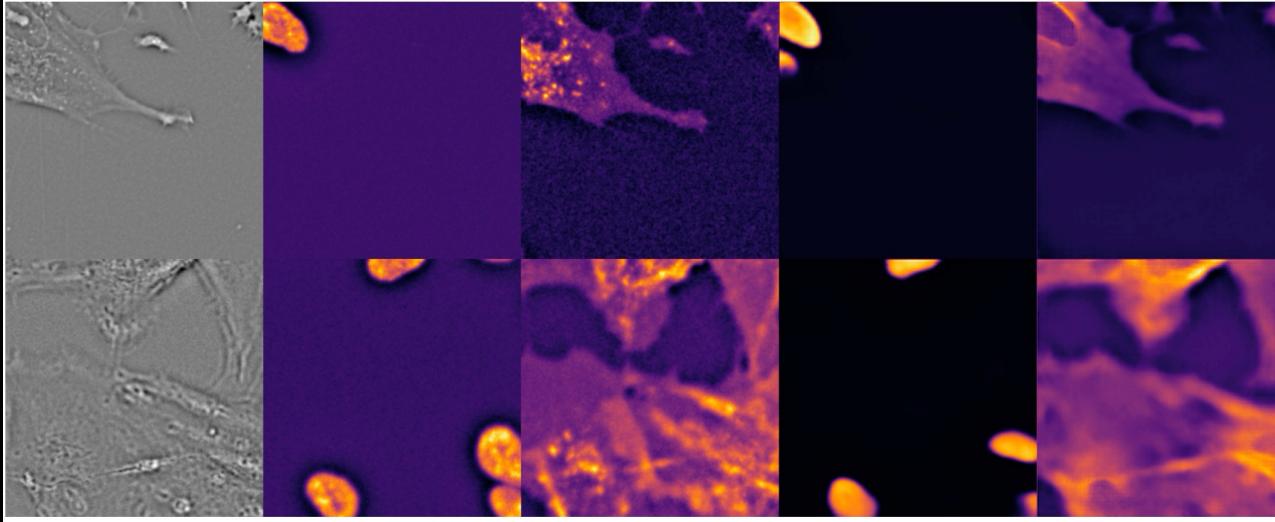
generalization across cell types

A

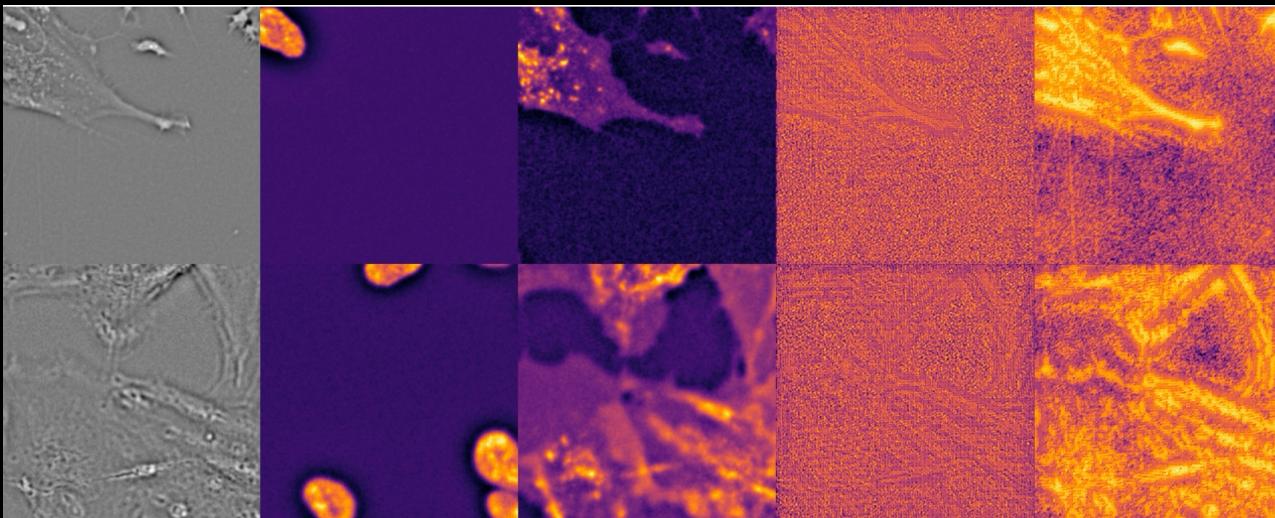


Pretraining for few-shot generalization

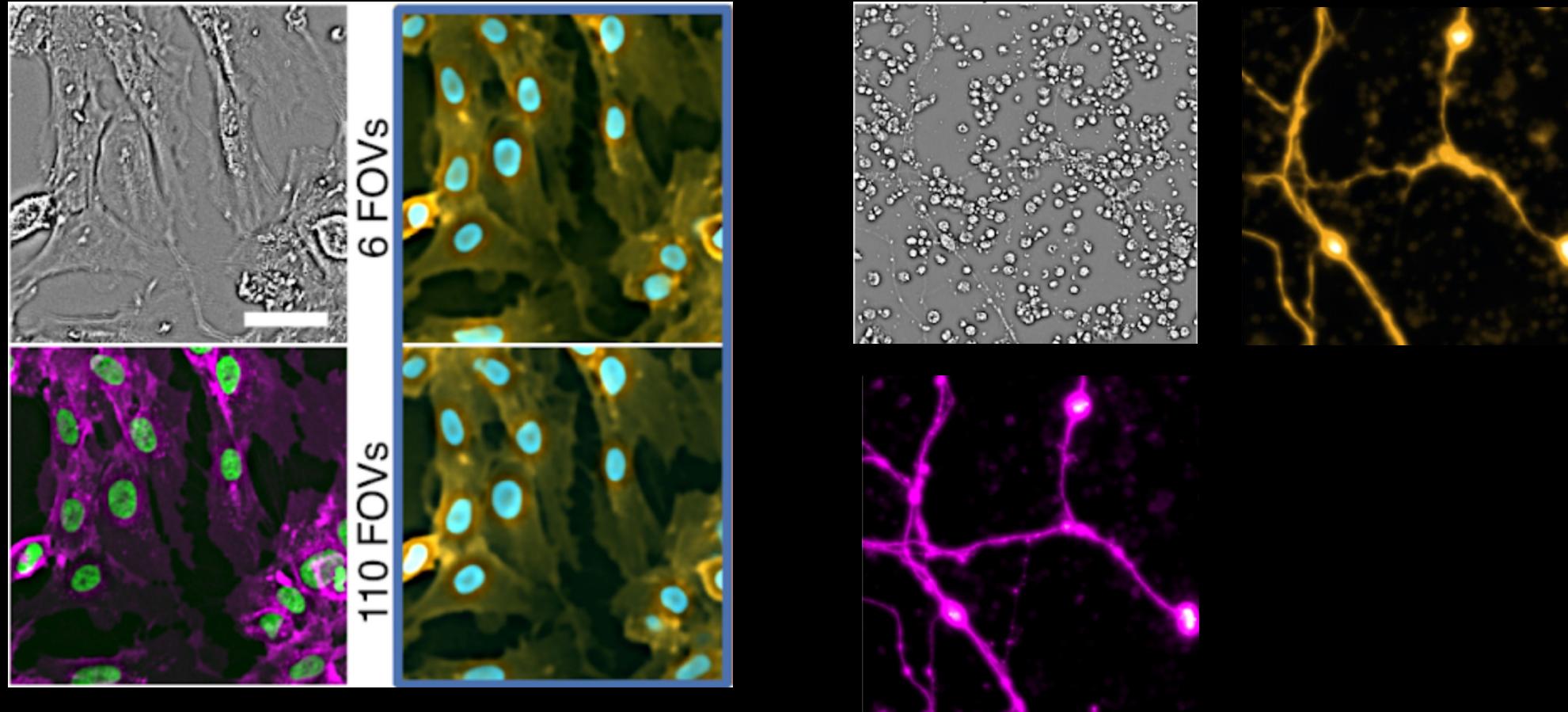
Pretrained +
fine-tuned

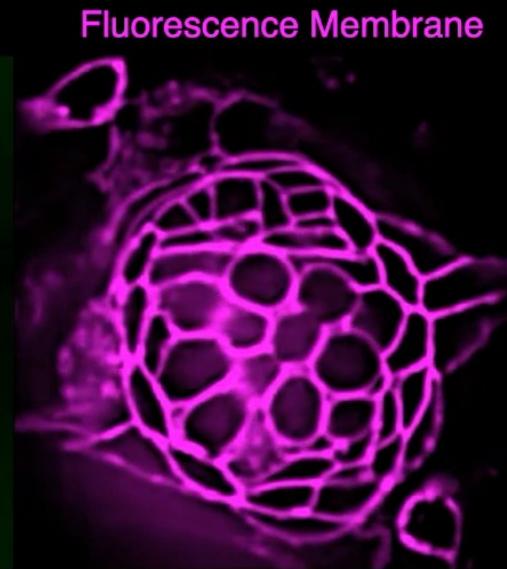
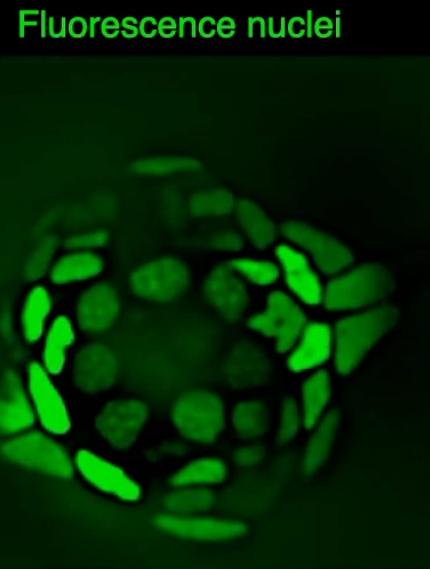
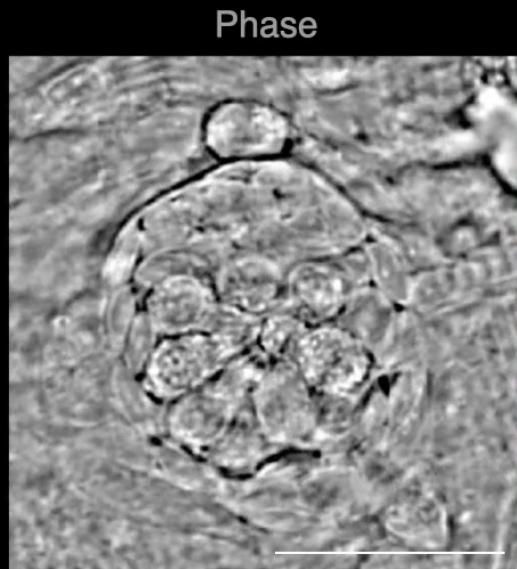


Trained from
scratch

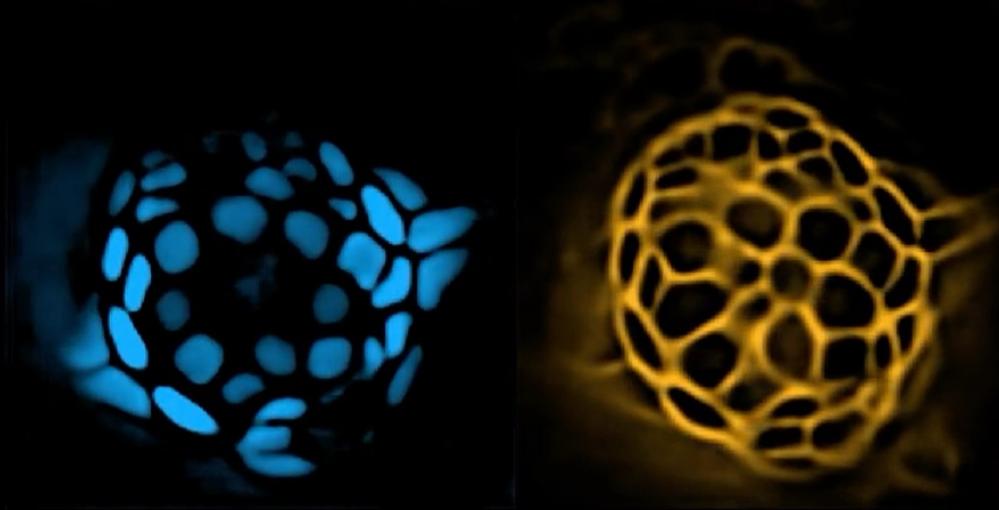


generalization across cell types





$t = 0\text{h}00\text{m}$, $z=5.94\mu\text{m}$

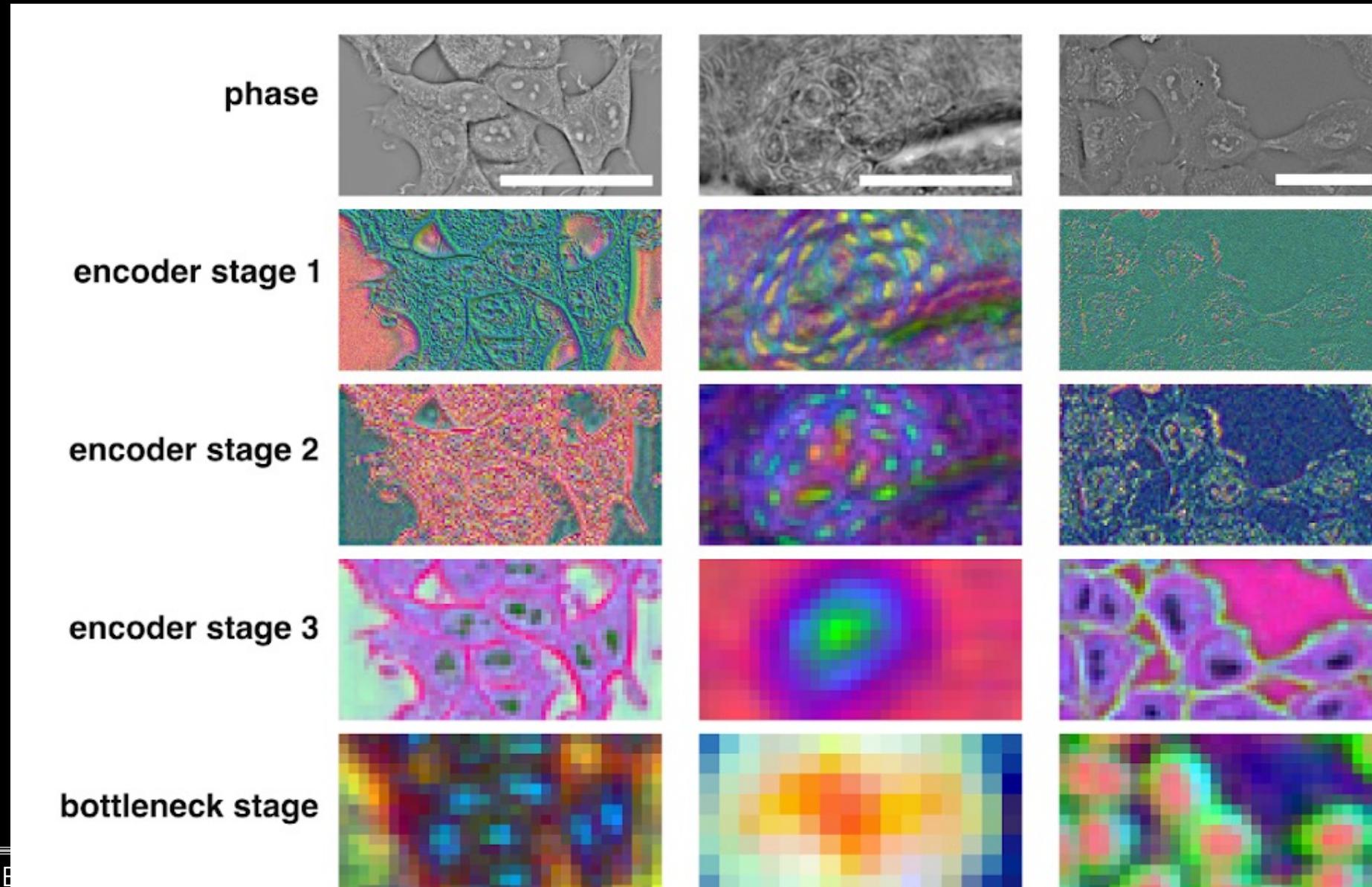


Virtual Staining nuclei

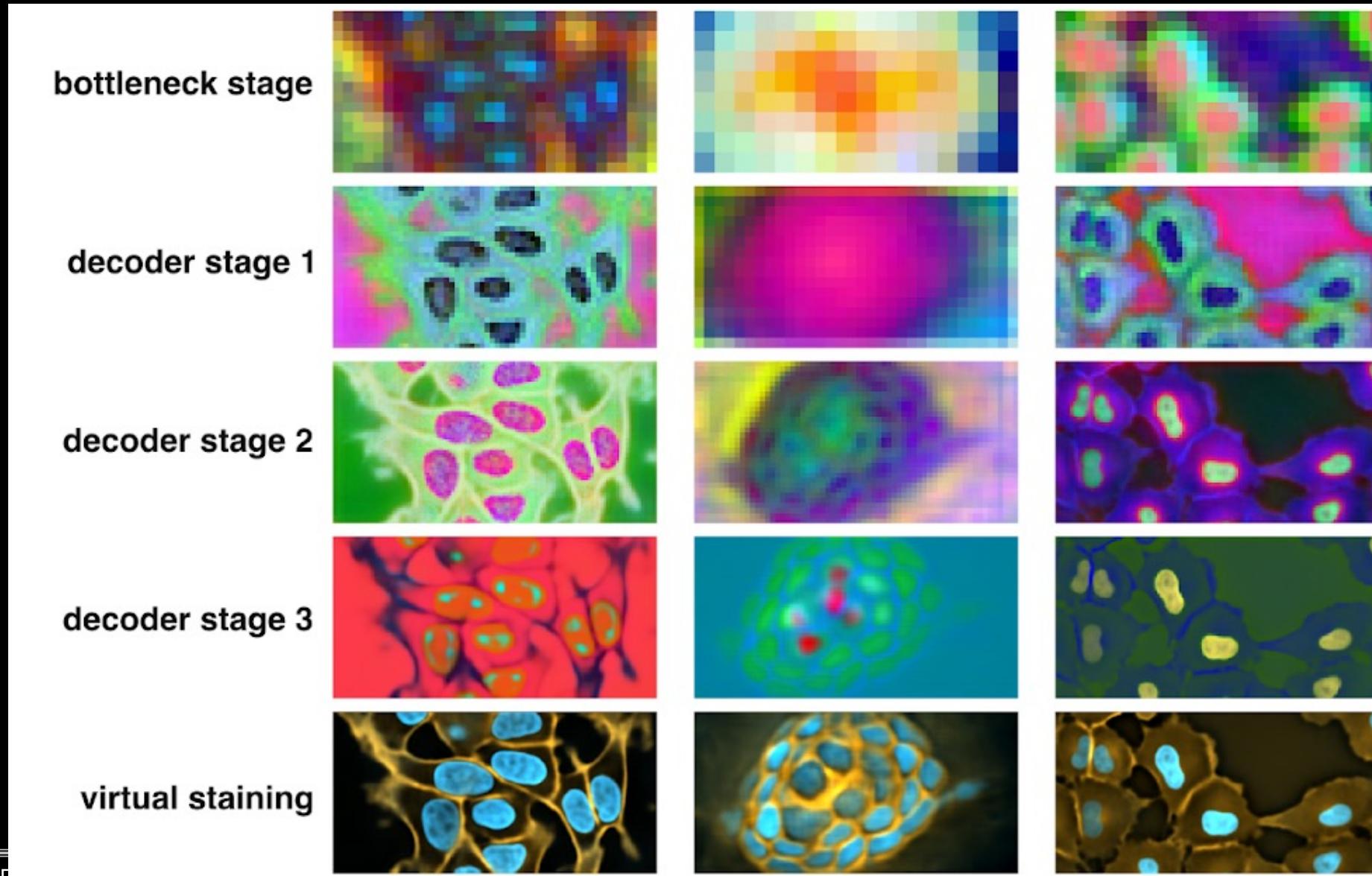
Virtual Staining membrane

...fine tune with sparse supervision

Multi-scale representation



Multi-scale representation



Thank you!

Computational Microscopy



CZ Biohub



Collaborators

