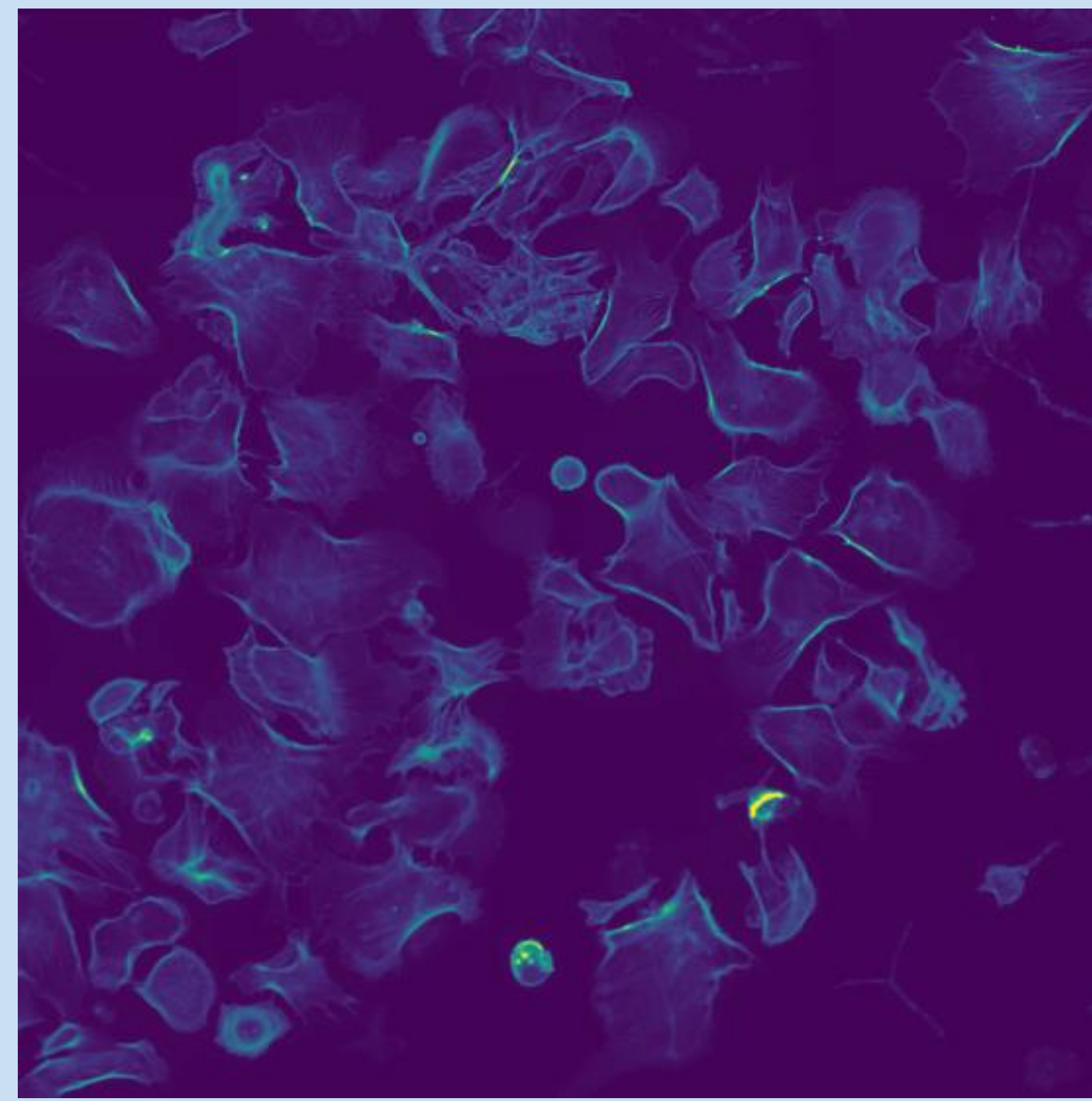


Deep Learning-based Segmentation and Deblurring of Microscopic Cell Images

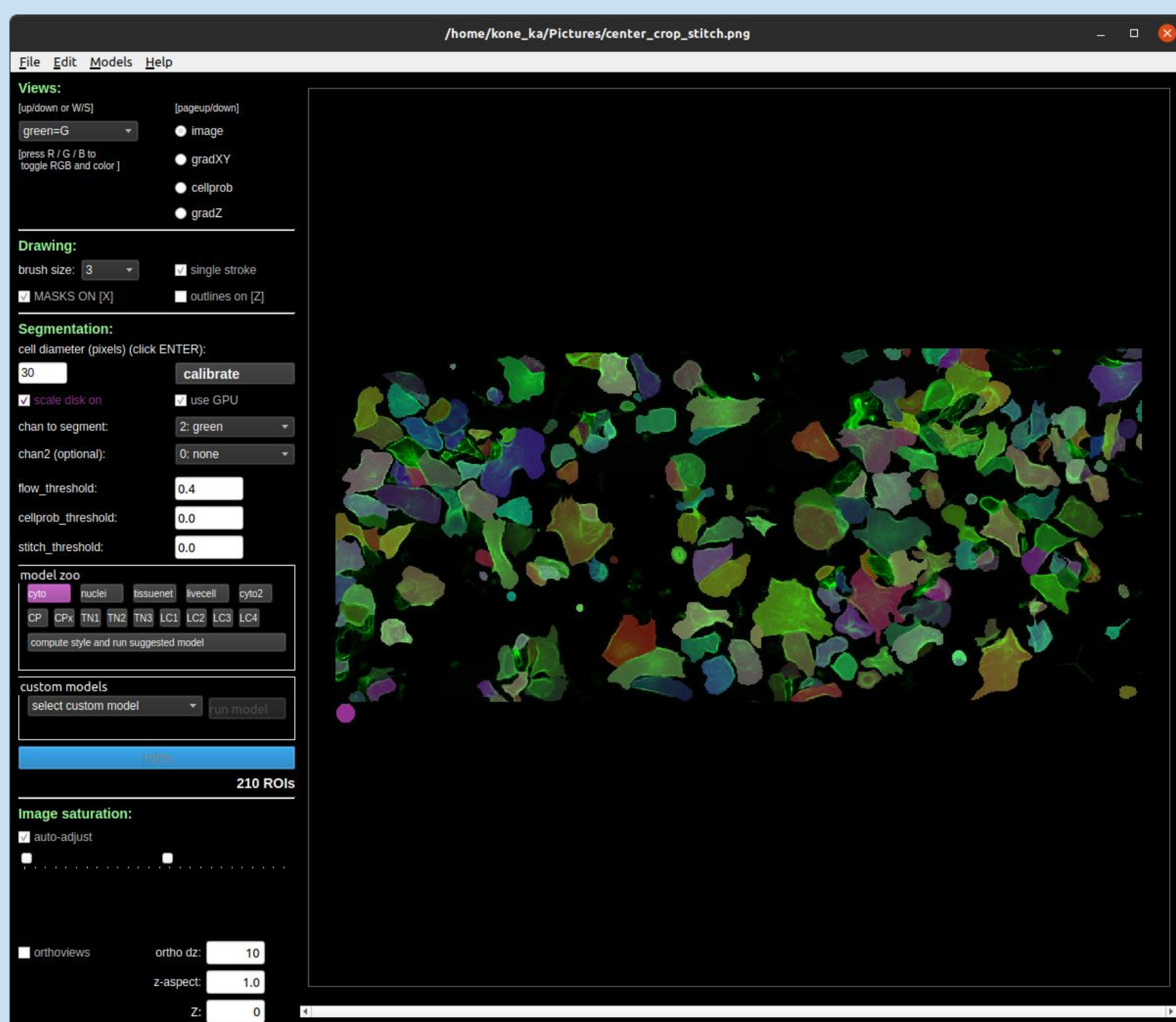
STED Microscope Image



From ME-BIO

Cellpose [1]

- × Generalist algorithm for **cellular segmentation** (www.cellpose.org)
- × **U-Net** architecture
- × **Human-in-the-loop** training possible via GUI



[1] Stringer, C., Wang, T., Michaelos, M. et al. Cellpose: a generalist algorithm for cellular segmentation. Nat Methods 18, 100–106 (2021). <https://doi.org/10.1038/s41592-020-01018-x>
 [2] Unpaired Image-to-Image Translation using Cycle-Consistent Adversarial Networks. Jun-Yan Zhu*, Taesung Park*, Phillip Isola, Alexei A. Efros. In ICCV 2017. (* equal contributions)

Deblurring

Results

Motivation

- × **Microscopic images** from fluorescence or STED microscopy can contain **blurry parts** due to diffraction barrier, astigmatism, defects and human error
- × Tracing of **intracellular filamentous structures** not possible/difficult
- × Use a **CycleGAN** to translate blurry images into sharp images

Method

- × Using <https://github.com/junyanz/pytorch-CycleGAN-and-pix2pix> [2] implementation
- × Results based on **103 paired images** with blurry (domain X) and sharp (domain Y) version
- × Based on domain X/Y, G/F generates fake images in domain Y/X
- × D_X/D_Y discriminates if image is from domain X/Y
- × Adversarial loss: $\mathbb{E}_{y \sim p_{\text{data}}(y)} [\log D_Y(y)] + \mathbb{E}_{x \sim p_{\text{data}}(x)} [\log(1 - D_Y(G(x)))]$
- × Cycle consistency loss: $\mathbb{E}_{x \sim p_{\text{data}}(x)} [\|F(G(x)) - x\|_1] + \mathbb{E}_{y \sim p_{\text{data}}(y)} [\|G(F(y)) - y\|_1]$

Cell Filament Analysis

Results

Motivation

- × **Manual analysis** of the differences between actin fibers from cells grown under microgravity compared to cells grown under standard conditions entails an **extremely high workload**
- × An **automated analysis** would benefit research areas that work with such filamentous cell data

Method

- × **Ridge Detection** via first Eigen value of the **Hessian Matrix** of the cell image
- × **Filtering** out **small filament** candidates
- × Calculating the **orientation** of individual filaments via **PCA**

