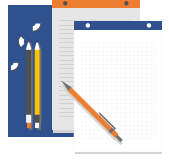


Optimizing Deep Learning Models for Cell Recognition in Fluorescence Microscopy: the Impact of Loss Functions on Performance and Generalization

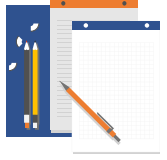
Luca Clissa, Antonio Macaluso, Antonio Zoccoli

luca.clissa2@unibo.it



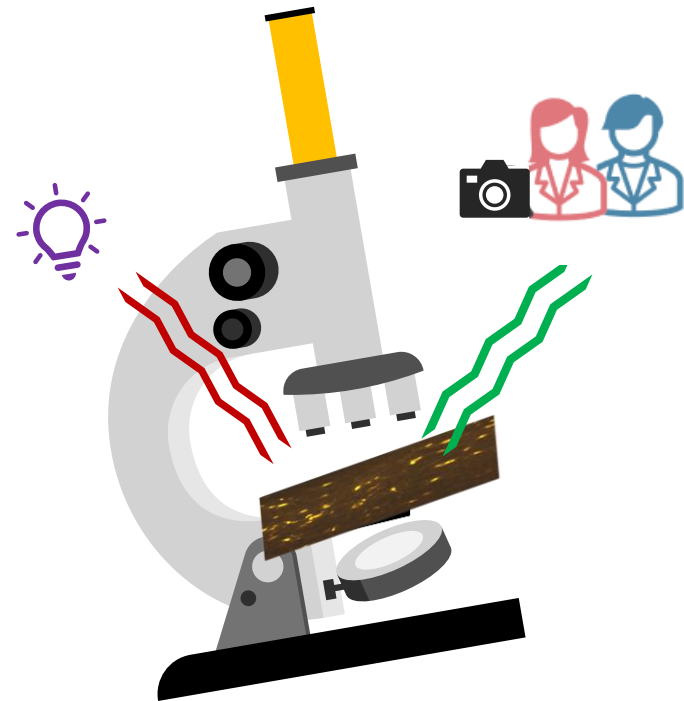
Outline

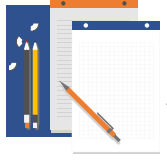
- Fluorescence Microscopy
- Challenges
- Experimental setup
- Results
- Conclusions



Fluorescence microscopy

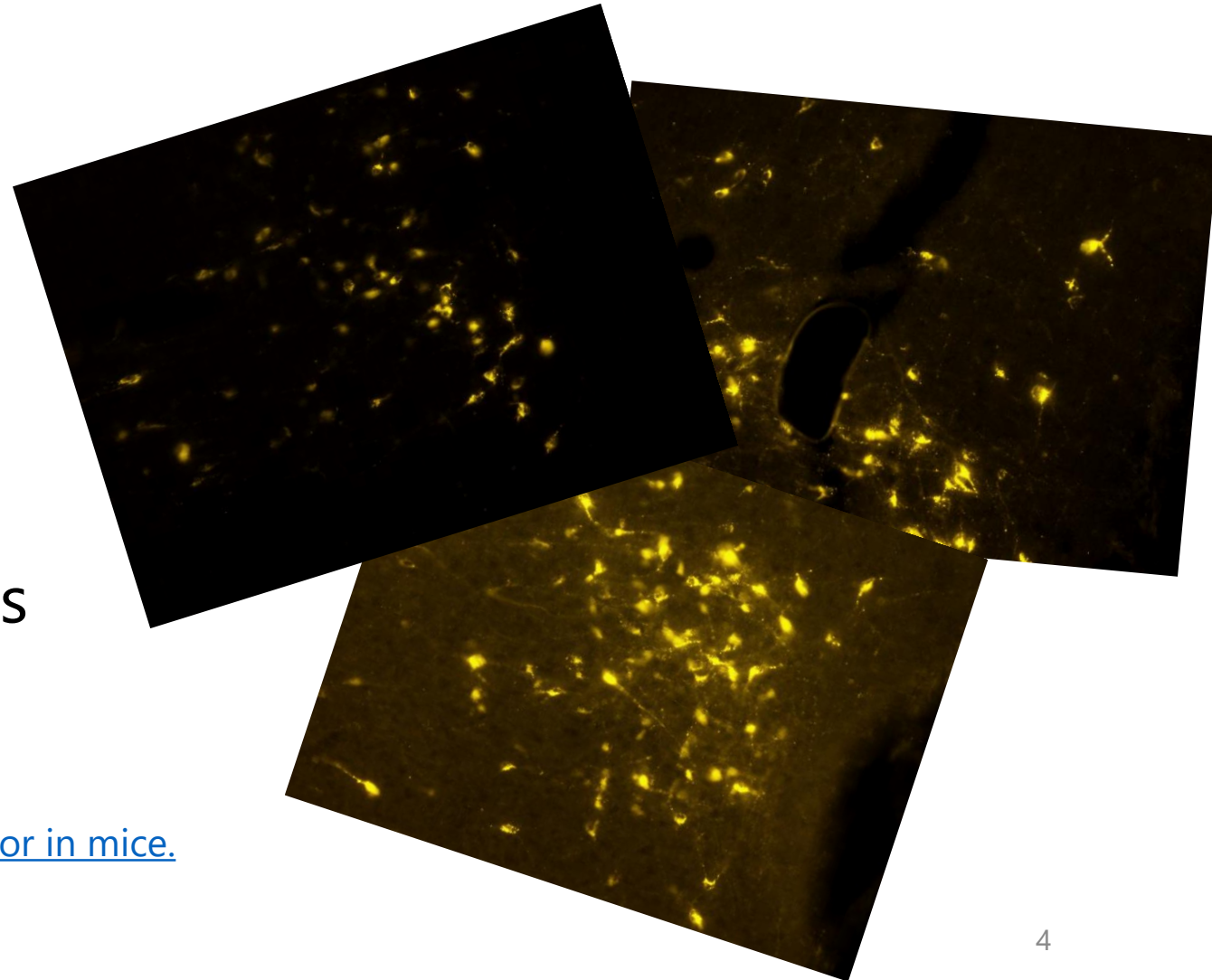
- Physics-based imaging technique
- Exploits light absorption/emission properties
- Used to mark/tag/stain biological compounds



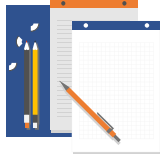


Applications

- Very popular in life science
 - Torpor onset [\[1\]](#)
- Cytoplasmatic neuronal structures
- Variability in shape, size and color hue
- **Goal:** count stained structures

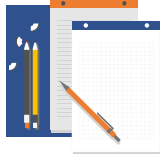


[\[1\] Hitrec, T., et al.: Neural control of fasting-induced torpor in mice. Scientific Reports 9\(1\) \(oct 2019\).](#)



Problem

- Manual processing
 - Time-consuming
 - Error-prone
 - Subjectivity of borderline cases
- Hard to adapt Deep Learning solutions
 - Domain shift
 - Few in-domain annotated datasets
 - How to train? How to evaluate?

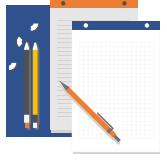


Contributions

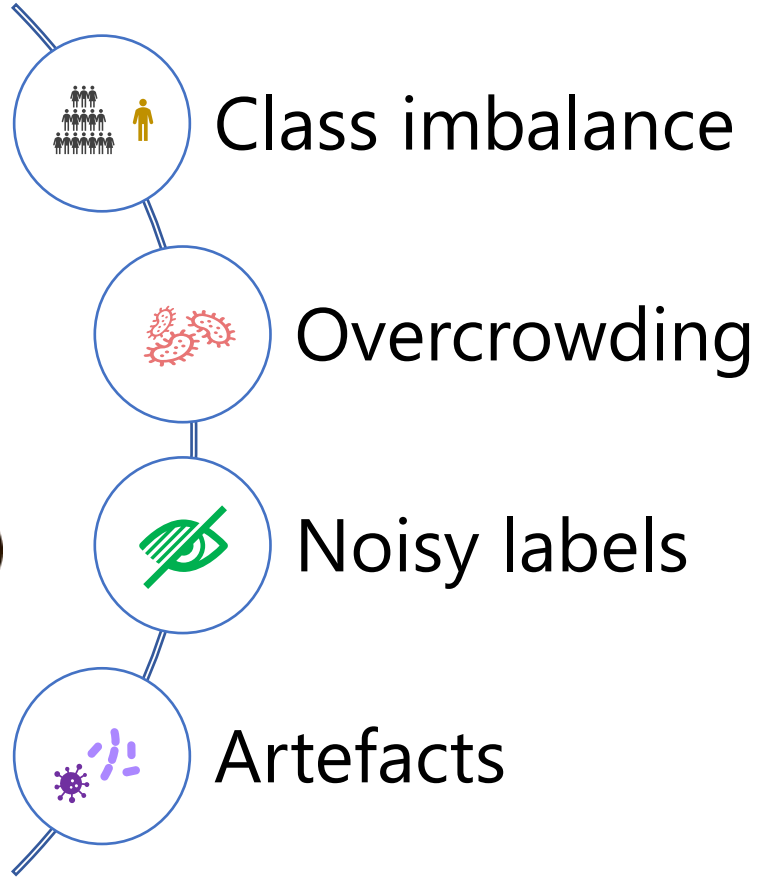
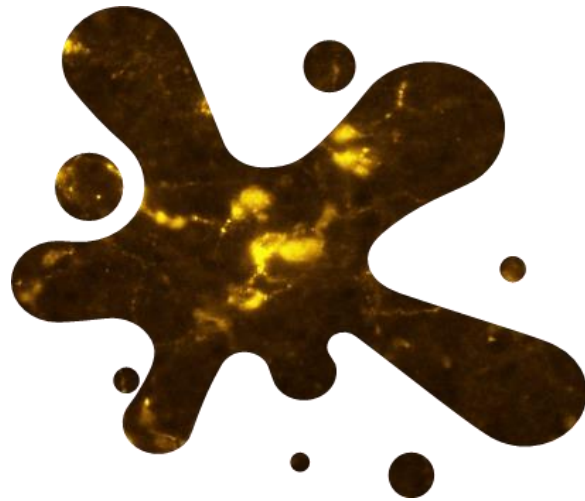
- Semantic segmentation using c-ResUnet [\[2\]](#) and Fluorescent Neuronal Cells v2 dataset (FNC v2) [\[3\]](#)
- Show the impact of loss functions on model performance
 - 60 ablation studies
 - 6 loss functions
- Inspect pros and cons of several evaluation metrics
- Discuss characteristics affecting out-of-sample generalization

[\[2\] Morelli, R., et al.: Automating cell counting in fluorescent microscopy through deep learning with c-ResUnet. Scientific Reports 11\(1\), 22920 \(2021\).](#)

[\[3\] Clissa, L., et al.: Fluorescent neuronal cells v2: Multi-task, multi-format annotations for deep learning in microscopy. arXiv preprint \(submitted to Scientific Data\) \(2023\)](#)



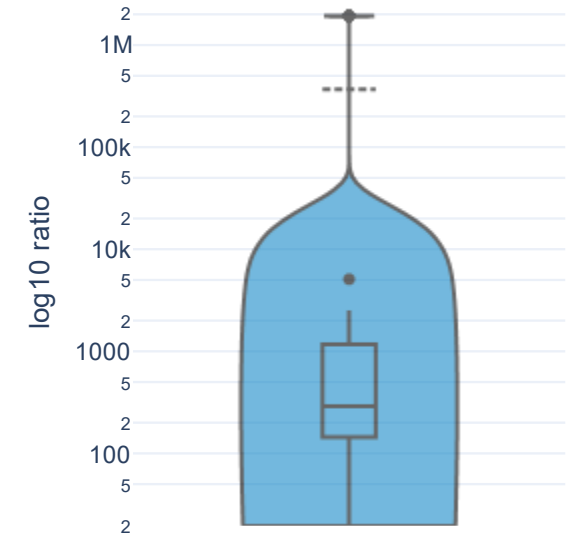
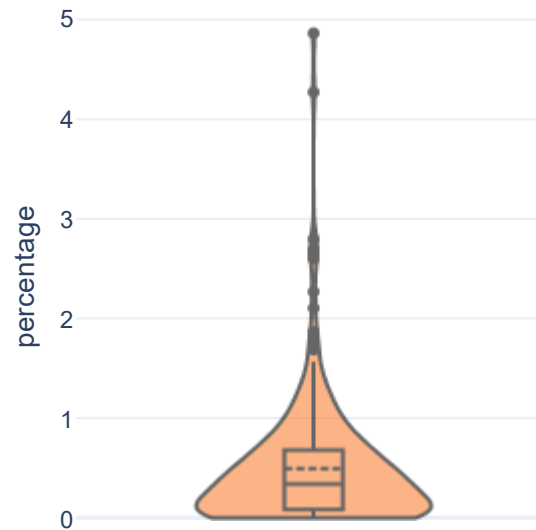
Challenges





Class imbalance

quantile	signal (%)	signal ratio
mean	0.50	367k
s.d.	0.61	756k
min	0	19.57
10%	0	92.39
25%	0.09	145.35
50%	0.34	291.10
75%	0.68	1k
90%	1.07	1.9M
max	4.86	1.9M



10 μ m

cells agglomerate

cells agglomerate

non-marked cell
type: shaded

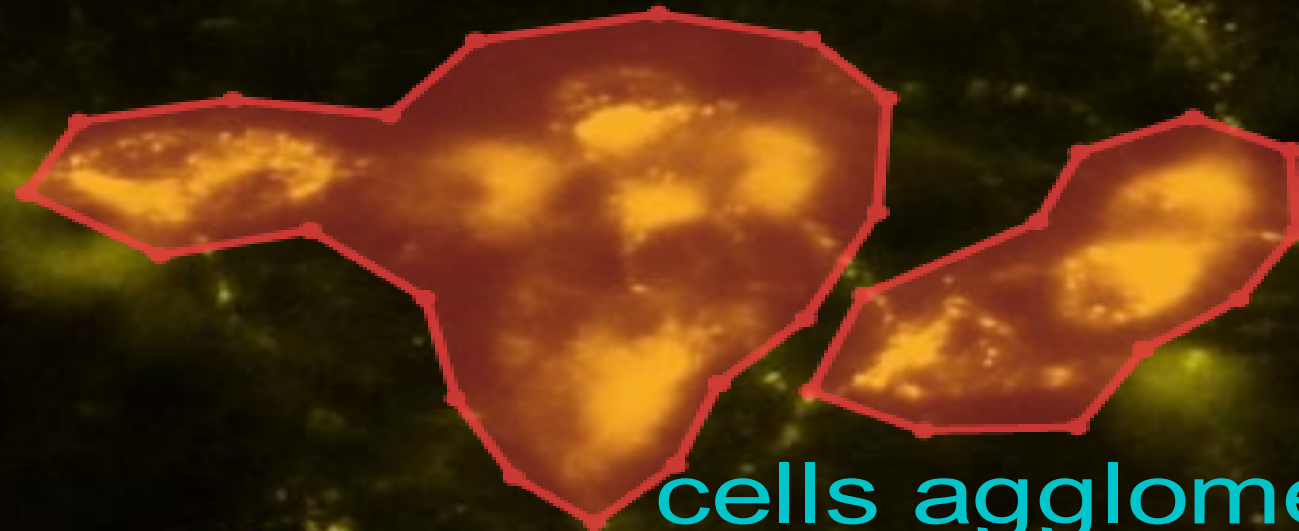
marked cell
type: dotted

cells agglomerate
cells agglomerate

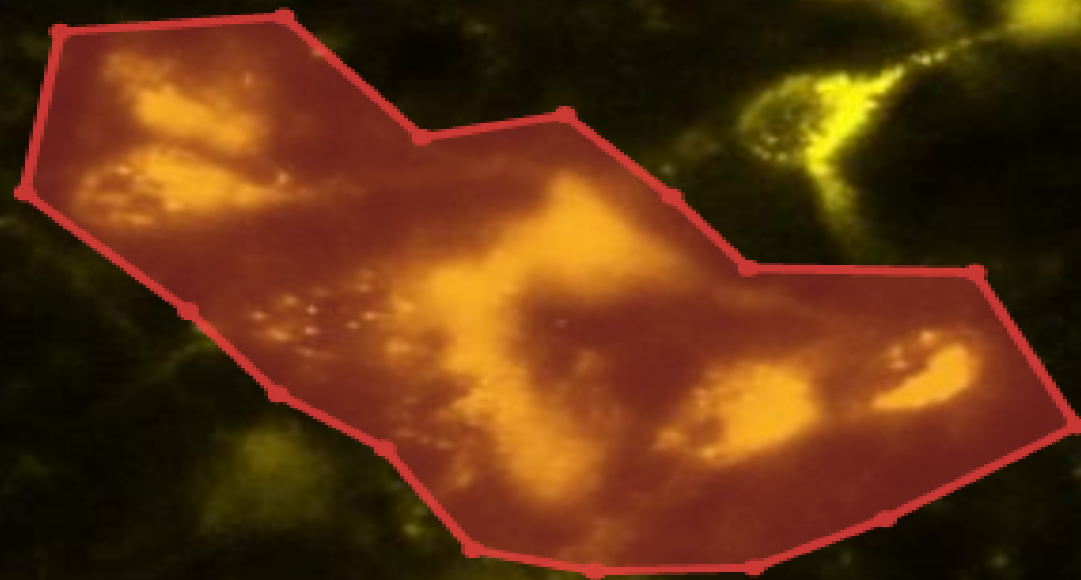
cells agglomerate

non-marked cell
type: dotted

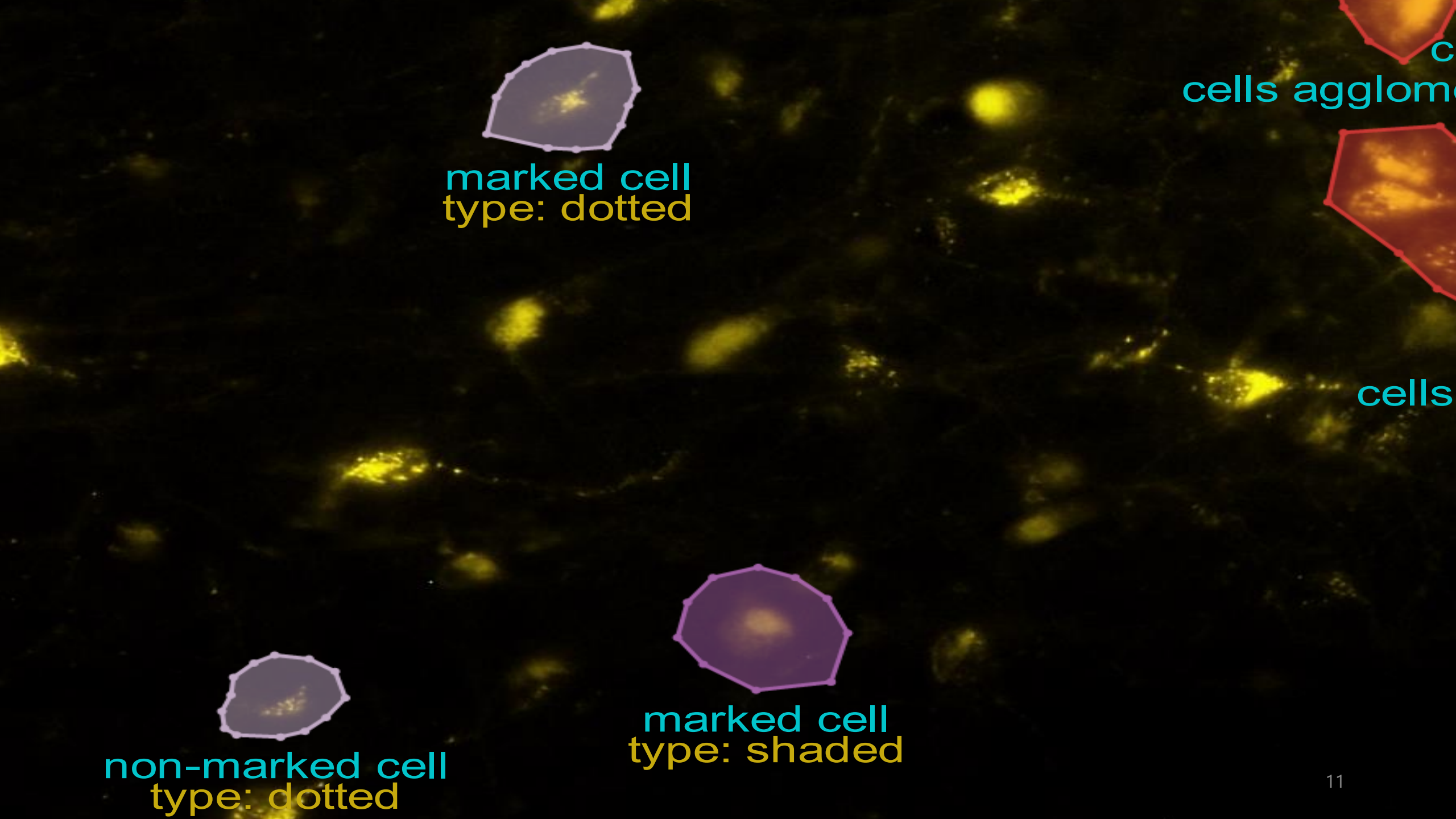
marked cell
type: shaded



cells agglomerate
cells agglomerate



cells agglomerate



marked cell
type: dotted

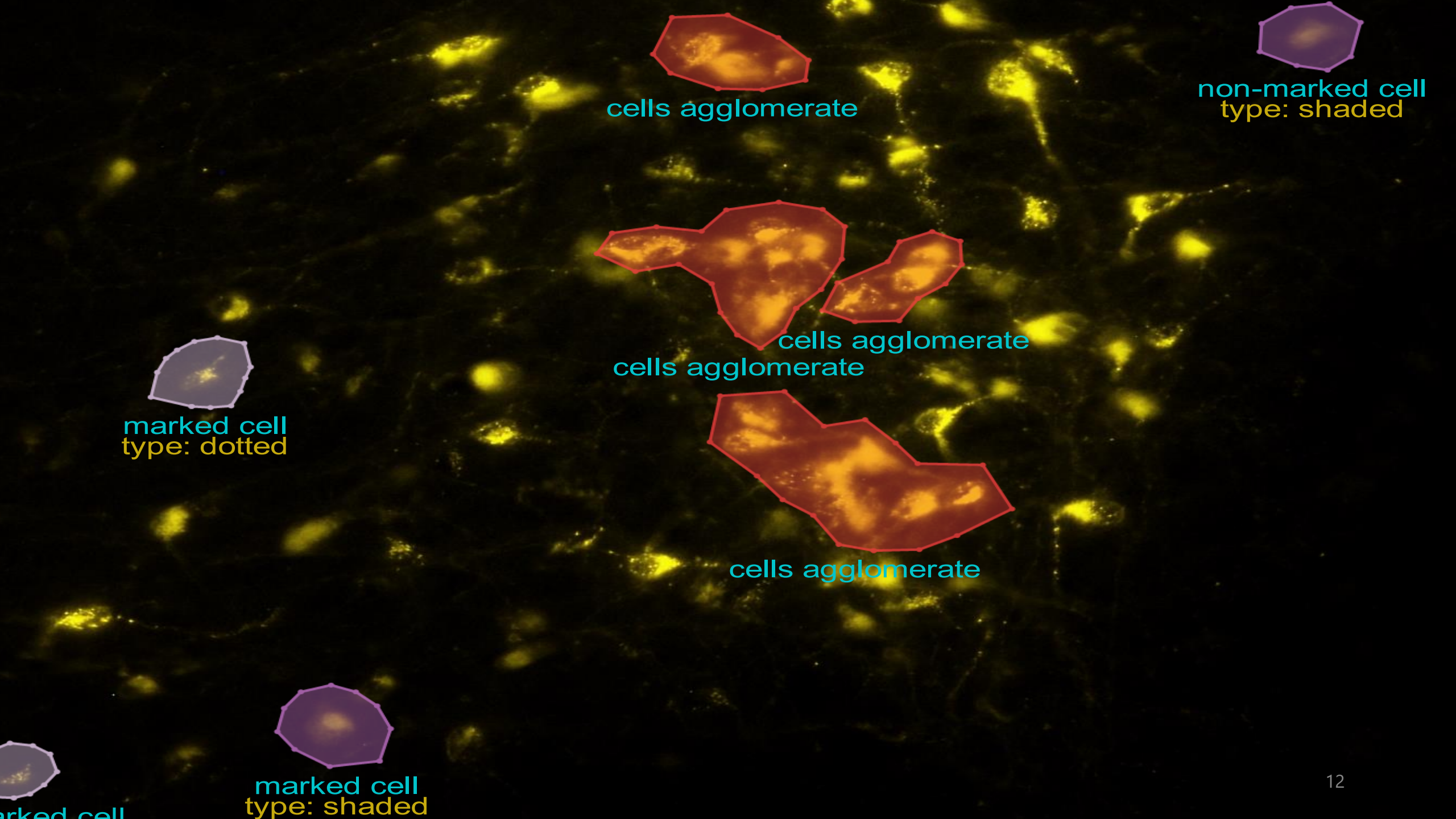
cells agglom



cells

non-marked cell
type: dotted

marked cell
type: shaded



cells agglomerate

non-marked cell
type: shaded

cells agglomerate
cells agglomerate

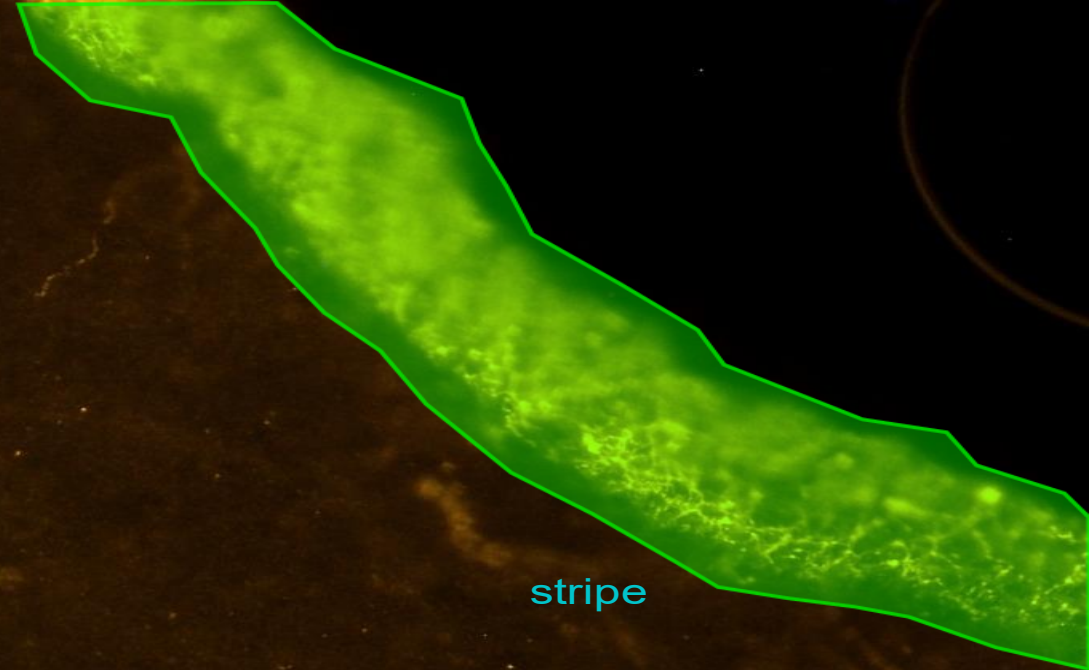
marked cell
type: dotted

cells agglomerate

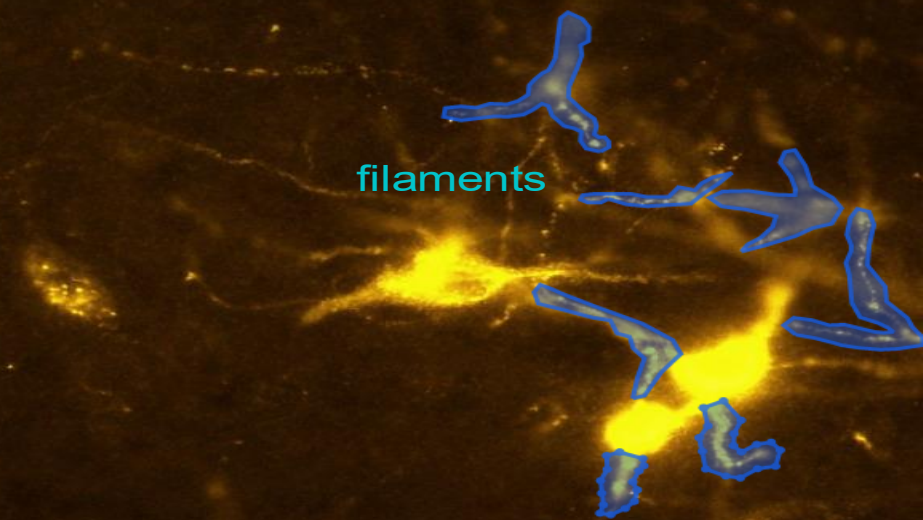
marked cell
type: shaded

marked cell

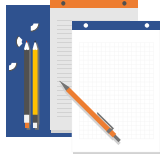
10 μm



stripe

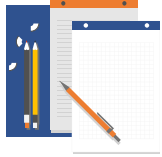


filaments



Ablation studies

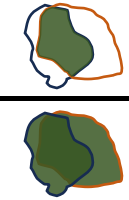
- 4 alternative losses
 - Weighted Binary Cross Entropy (BCE): $w_{cell} = 50, 100, 200$; $w_{bkgd} = 1$
 - Dice Loss
 - Focal Loss
 - Focal Tversky Loss
- 2 combined losses
 - CombinedLoss = $\lambda_1 BCE + \lambda_2 \text{Dice} + \lambda_3 \text{Focal}$
 - CombinedFTLoss = $\lambda_1 BCE + \lambda_2 \text{Dice} + \lambda_3 \text{Focal Tversky}$
 - Balanced: $\lambda_1 = 0.3, \lambda_2 = 0.3, \lambda_3 = 0.4$
 - Overcrowd: $\lambda_1 = 0.2, \lambda_2 = 0.5, \lambda_3 = 0.3$
 - CellViT: $\lambda_1 = 0.5, \lambda_2 = 0.3, \lambda_3 = 0.5$



Evaluation Metrics

- Segmentation

- Mean Intersection over Union (mIoU) = $\frac{\text{Intersection}}{\text{Union}}$
- threshold: 0.4



- Detection

- Centers distance
- threshold: 40 pixels (mean cell diameter)

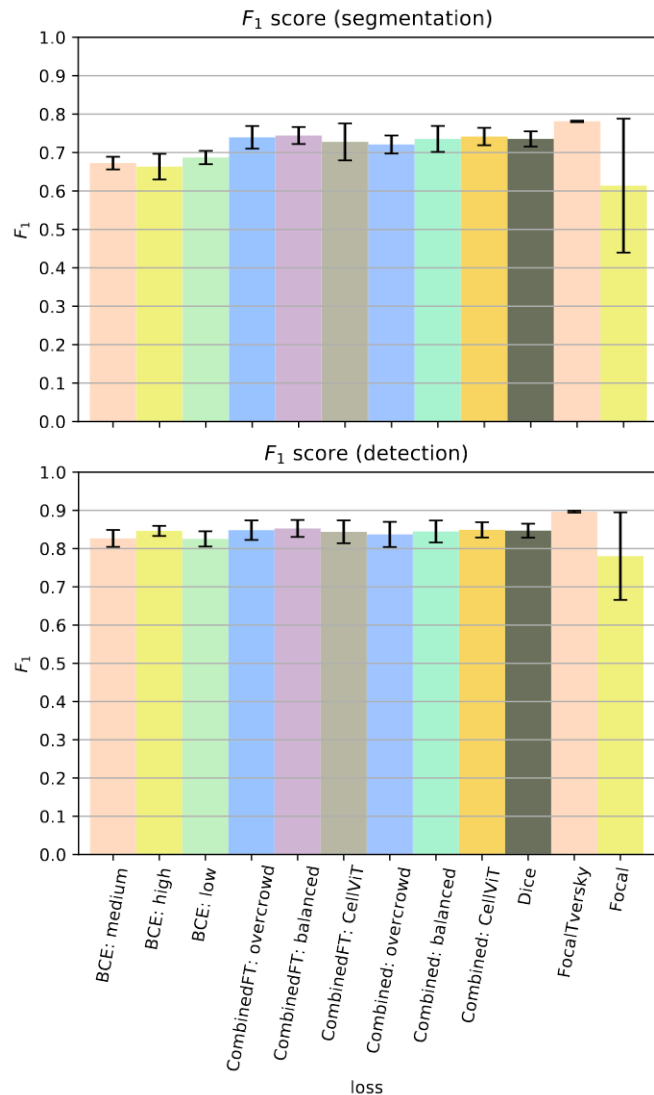


- Counting

- Mean Absolute Error
- Median Absolute Error
- Mean Percentage Error: $\frac{(n_t - n_p)}{\max(n_t, 1)} * 100$



Segmentation & Detection

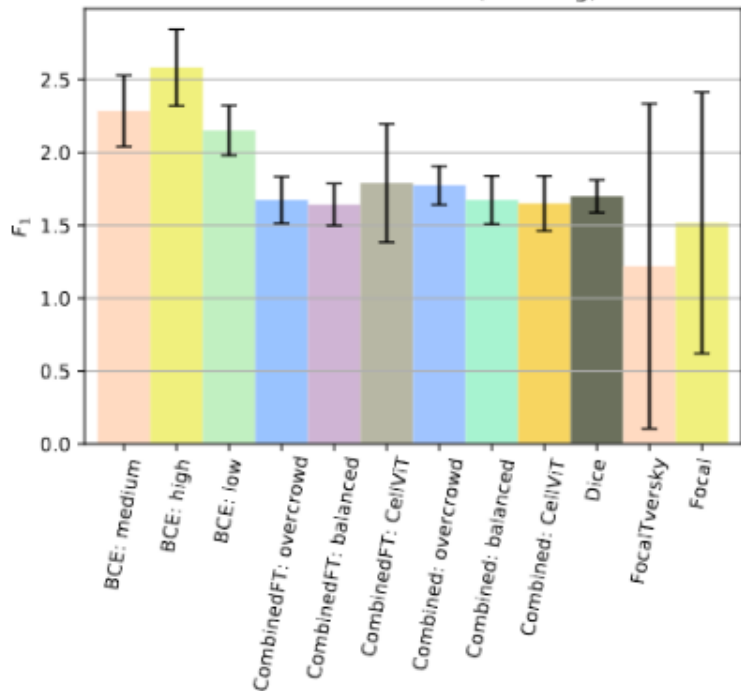


Loss	F1 score (IoU)	F1 score (distance)
BCE: medium	0.673±0.017	0.827±0.022
BCE: high	0.663±0.033	0.846±0.013
BCE: low	0.687±0.017	0.825±0.020
CombinedFT: overcrowd	0.740±0.029	0.848±0.026
CombinedFT: balanced	0.744±0.022	0.853±0.022
CombinedFT: CellViT	0.728±0.048	0.844±0.030
Combined: overcrowd	0.721±0.023	0.837±0.033
Combined: balanced	0.735±0.034	0.845±0.029
Combined: CellViT	0.742±0.023	0.849±0.020
Dice	0.735±0.020	0.847±0.018
Focal Tversky	0.781±0.002	0.897±0.003
Focal	0.614±0.027	0.780±0.034

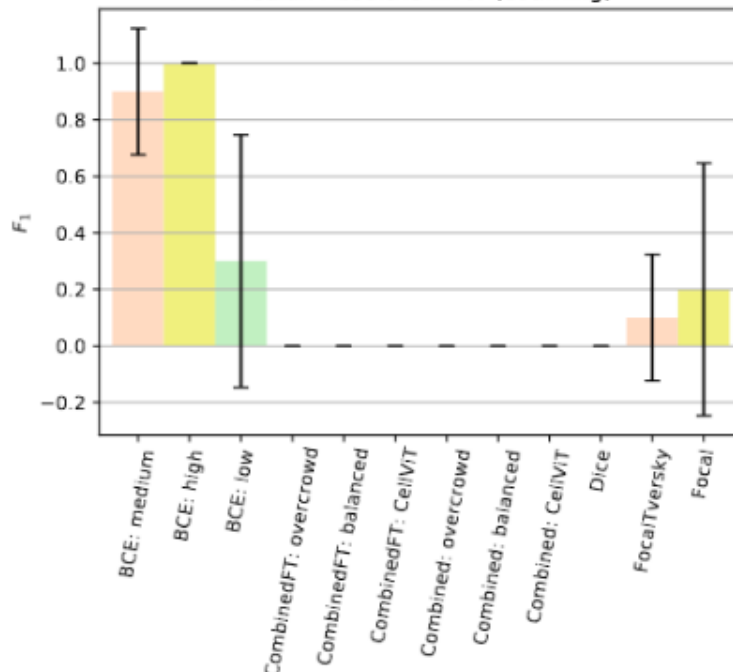


Counting

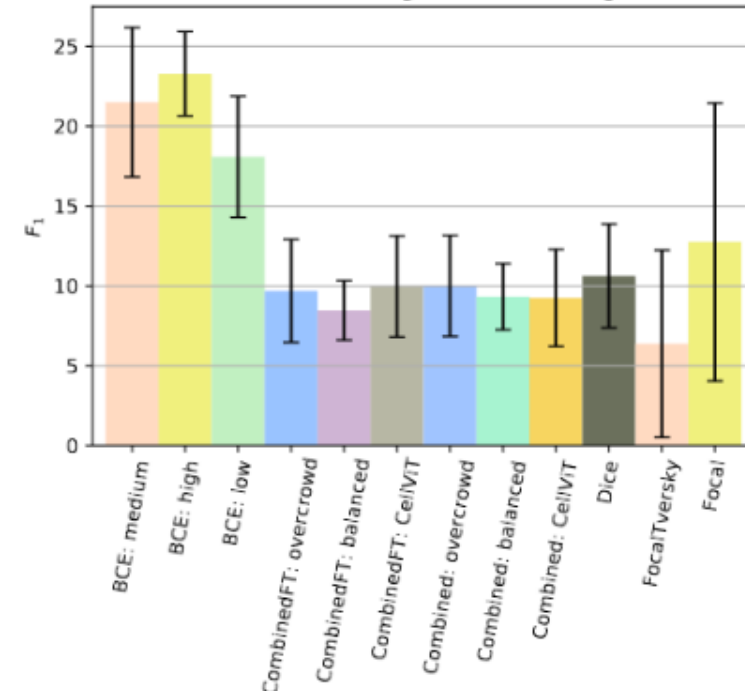
Mean Absolute Error (counting)



Median Absolute Error (counting)



Mean Percentage Error (counting)



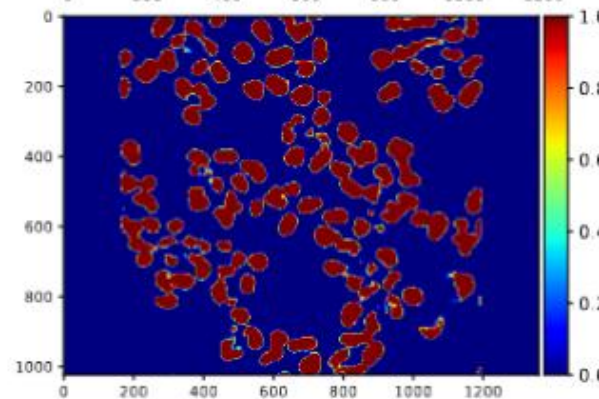
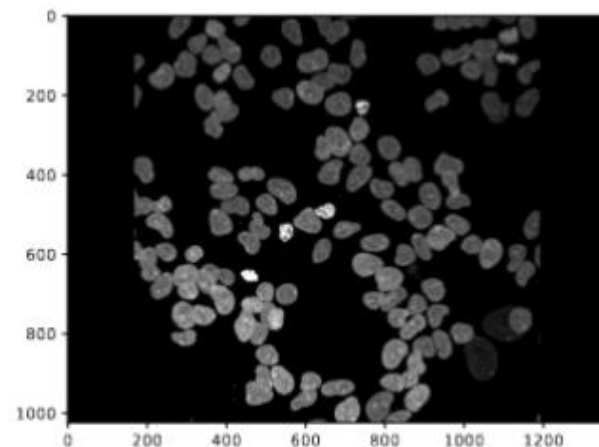
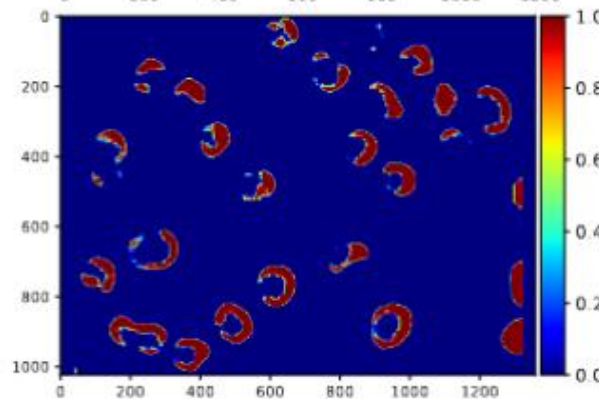
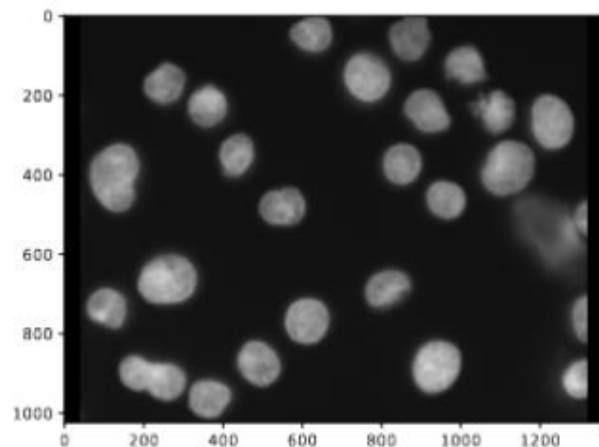
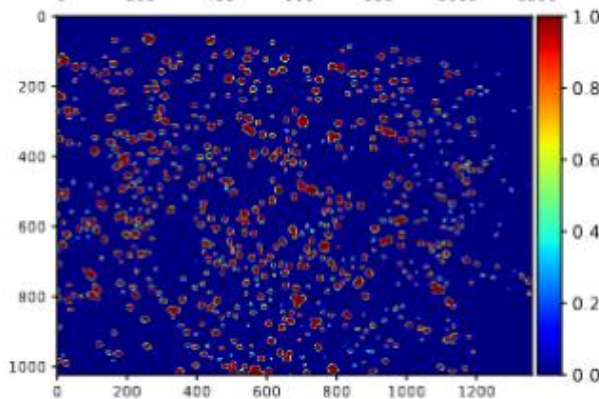
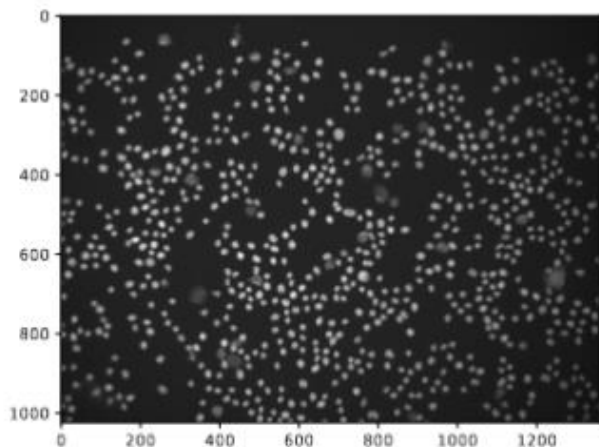
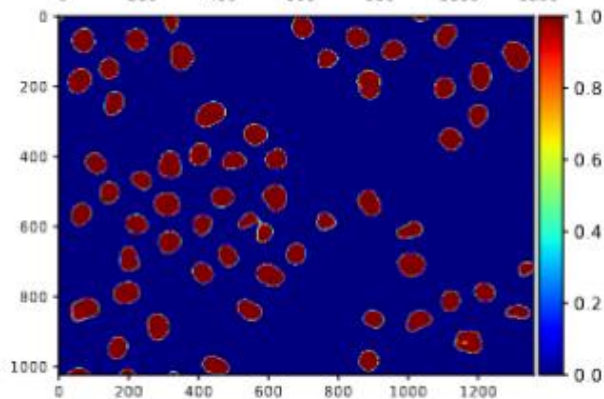
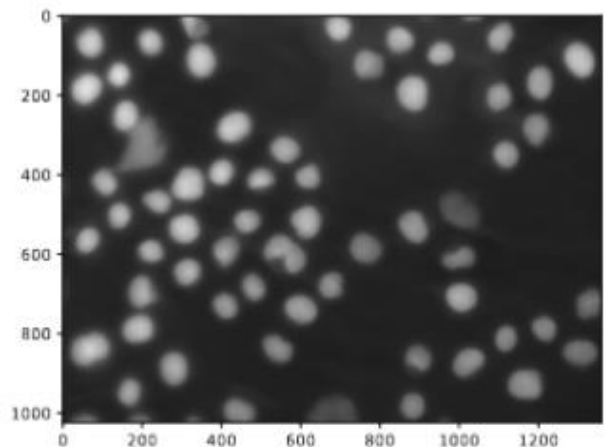
BCE: medium *BCE: high* *BCE: low* *CombinedFT: overcrowd* *CombinedFT: balanced* *CombinedFT: CellViT* *Combined: overcrowd* *Combined: balanced* *Combined: CellViT* *Dice* *Focal Tversky* *Focal*

	<i>BCE: medium</i>	<i>BCE: high</i>	<i>BCE: low</i>	<i>CombinedFT: overcrowd</i>	<i>CombinedFT: balanced</i>	<i>CombinedFT: CellViT</i>	<i>Combined: overcrowd</i>	<i>Combined: balanced</i>	<i>Combined: CellViT</i>	<i>Dice</i>	<i>Focal Tversky</i>	<i>Focal</i>
MAE	2.286±0.245	2.583±0.263	2.151±0.171	1.674±0.161	1.643±0.144	1.791±0.406	1.774±0.132	1.674±0.166	1.651±0.188	1.700±0.113	1.220±1.115	1.517±0.365
MedAE	0.9±0.224	1±0	0.3±0.447	0±0	0±0	0±0	0±0	0±0	0±0	0±0	0.1±0.224	0.2±0.4
MPE	21.498±4.679	23.281±2.637	18.079±3.791	9.682±3.221	8.465±1.859	9.968±3.163	10±3.160	9.322±2.074	9.249±3.039	10.621±3.247	6.373±1.607	12.747±3.525



Out-of-sample generalization

[4] S-BSST265 dataset

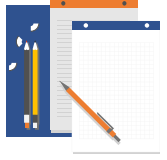


(a) Medium-sized, sharp objects

(b) Small objects

(c) Uneven texture and filling

(d) Overcrowding

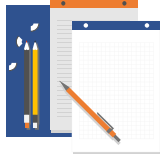


Conclusions

- Focal Tversky loss overperforms other losses
- Still some troubles separating crowded objects
→ careful post-processing needed (hole filling, small objects, watershed)
- Combined losses are competitive
 - Better tuning of lambda weights
- Generalization
 - High variability
 - Dedicated augmentation may help
 - Panoptic loss
- Integrating more metrics enables more comprehensive assessment

Thanks for your attention!

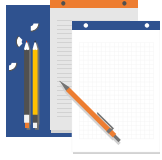
Questions?



References

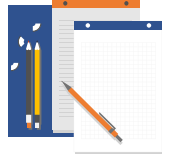
- [\[1\] Hitrec, T., et al.: Neural control of fasting-induced torpor in mice. Scientific Reports 9\(1\) \(oct 2019\).](#)
- [\[2\] Morelli, R., et al.: Automating cell counting in fluorescent microscopy through deep learning with c-ResUnet. Scientific Reports 11\(1\), 22920 \(2021\).](#)
- [\[3\] Clissa, L., et al.: Fluorescent neuronal cells v2: Multi-task, multi-format annotations for deep learning in microscopy. arXiv preprint \(submitted to Scientific Data\) \(2023\)](#)
- [\[4\] Kromp, F., et al.: An annotated fluorescence image dataset for training nuclear segmentation methods. Scientific Data 7\(1\), 262 \(2020\)](#)
- [\[5\] Jadon, S.: A survey of loss functions for semantic segmentation. In: 2020 IEEE conference on computational intelligence in bioinformatics and computational biology \(CIBCB\). pp. 1–7 \(2020\)](#)

Backup



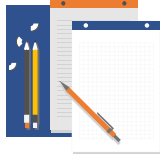
Ablation studies configurations

	BCE	Dice	Focal	Focal Tversky	Combined	Combined FT
Hyperparameters w_cell		smooth	gamma	gamma	$\lambda_1, \lambda_2, \lambda_3$	$\lambda_1, \lambda_2, \lambda_3$
Values	[50, 100, 200]	1×10^{-6}	2	2	balanced: [0.3, 0.3, 0.4] overcrowd: [0.2, 0.5, 0.3] CellViT: [0.5, 0.3, 0.5]	balanced: [0.3, 0.3, 0.4] overcrowd: [0.2, 0.5, 0.3] CellViT: [0.5, 0.3, 0.5]



Training setup

- Adam optimizer
- Learning rate test for initial LR
- 200 epochs
- Cyclical learning rates
- Best model on validation dice coefficient



Loss functions

- **Weighted Binary Cross Entropy:** higher weight to underrepresented class

$$L_{W-BCE}(y, \hat{y}) = -(\beta * y \log(\hat{y}) + (1 - y) \log(1 - \hat{y}))$$

- **Dice Loss:** targets segmentation performance directly, low impact of small objects

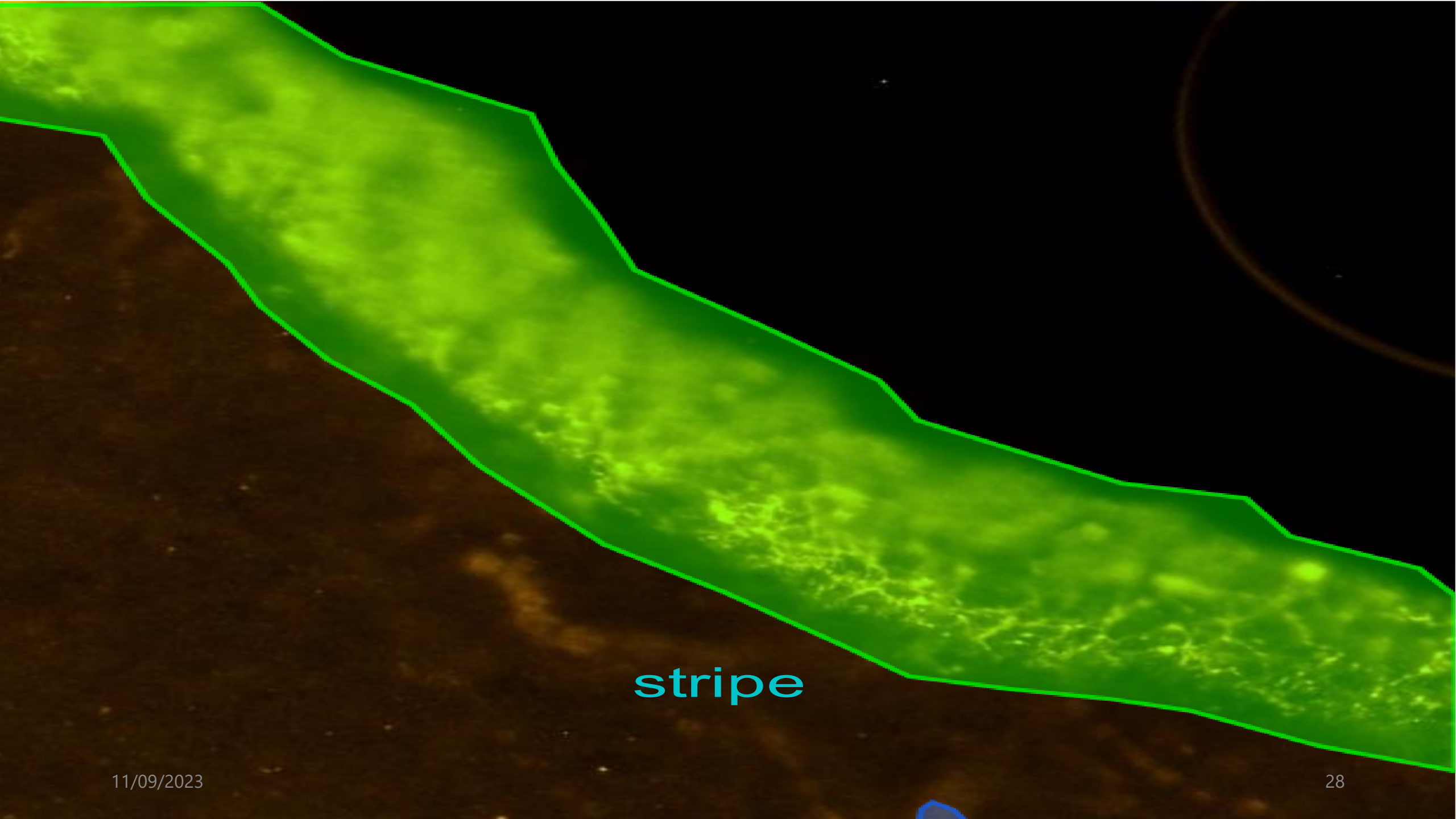
$$DL(y, \hat{p}) = 1 - \frac{2y\hat{p} + 1}{y + \hat{p} + 1}$$

- **Focal Loss:** oversample hard examples

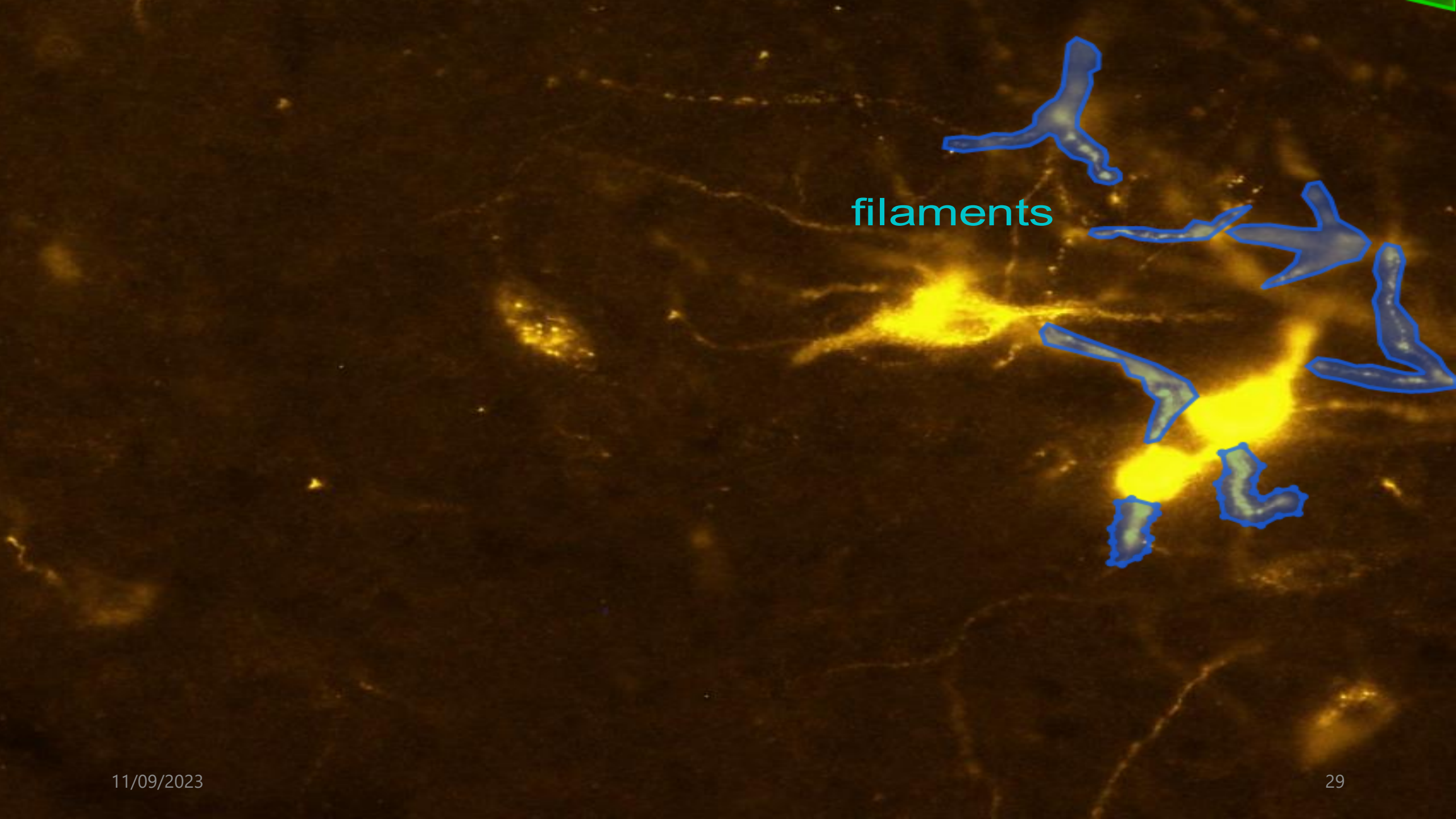
$$FL(p_t) = -\alpha_t (1 - p_t)^\gamma \log(p_t), \quad p_t = \begin{cases} p, & \text{if } y = 1 \\ 1 - p, & \text{otherwise} \end{cases}$$

- **Focal Tversky Loss:** bring together advantages of Dice and Focal losses

$$FTL = \sum_c (1 - TI_c)^\gamma, \quad TI(p, \hat{p}) = \frac{p\hat{p}}{p\hat{p} + \beta(1 - p)\hat{p} + (1 - \beta)p(1 - \hat{p})}$$



stripe



filaments

10 μm

artifact
size: small

artifact
size: small

artifact
size: big