

Midterm presentation

Automated recognition of NETs



Institute of Biochemistry - TiHo

Stakeholder

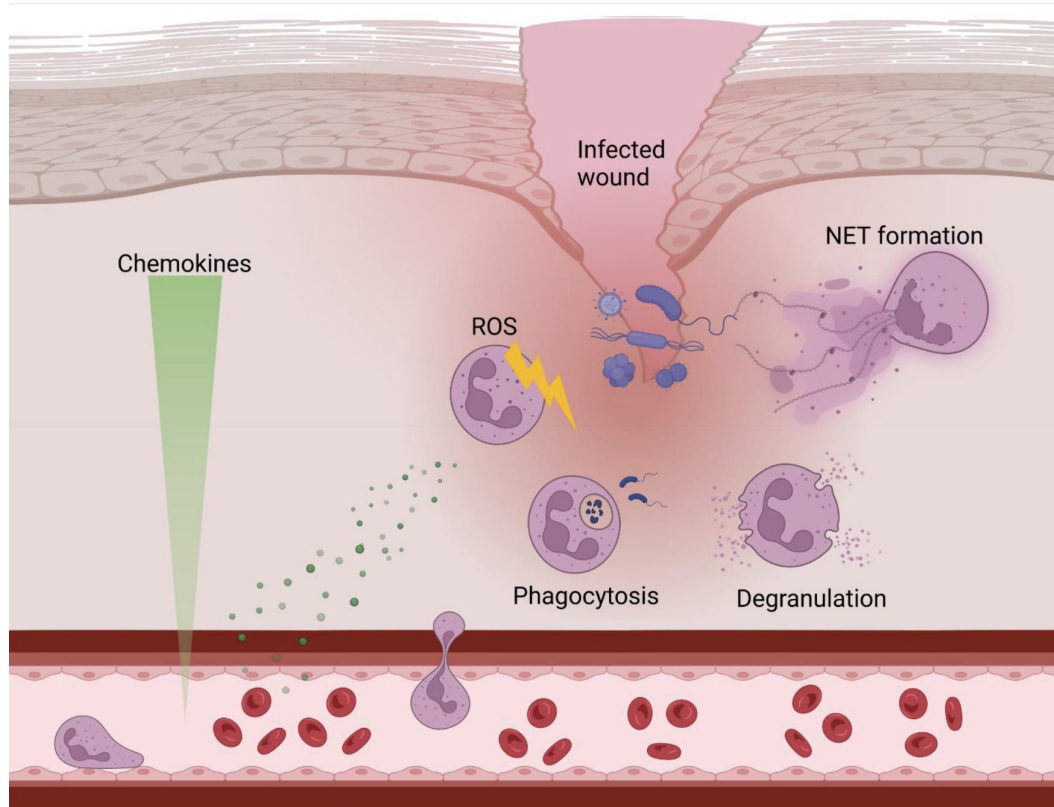


Interesting for anyone working in the NET field

Neutrophil Extracellular Traps

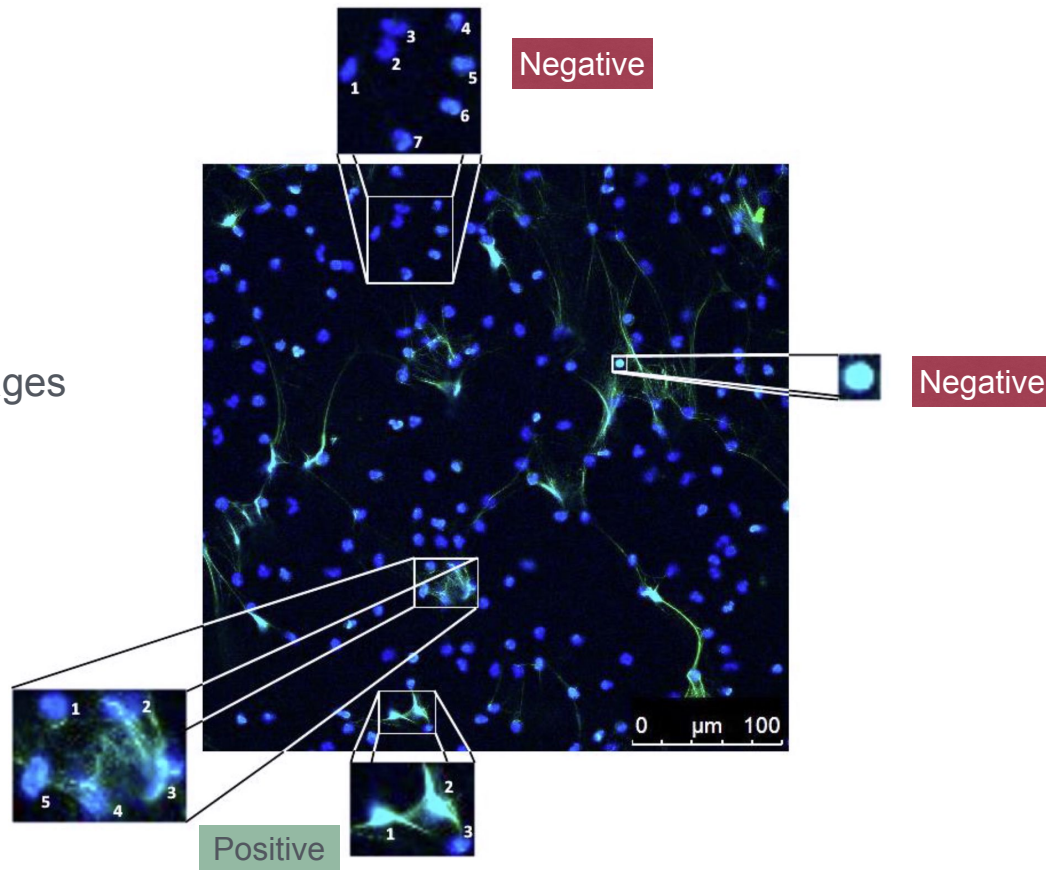
NETs

- Part of innate immune response mechanism
- Neutrophils perform programmed cell death (NETosis)
- NET: DNA backbone + (antimicrobial) proteins



How does our data look like?

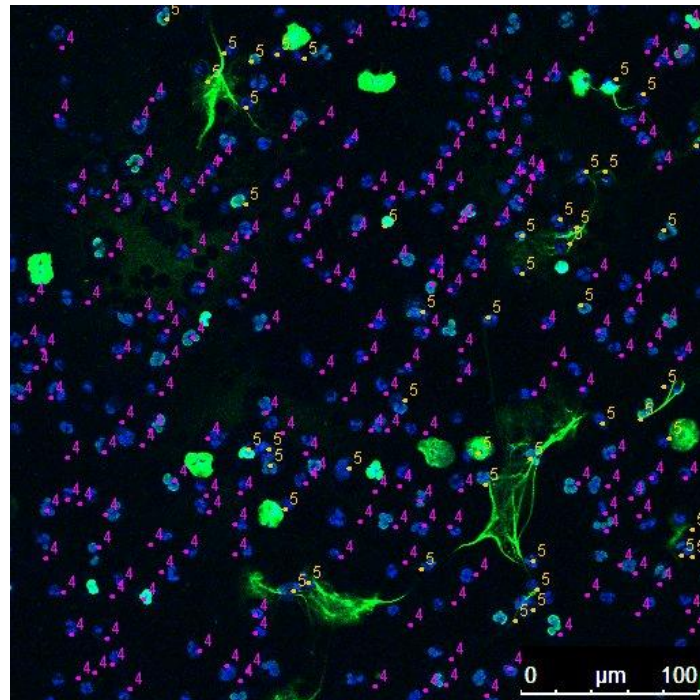
38 manually annotated images



Baseline models

Earlier attempts to automatise the NET counting

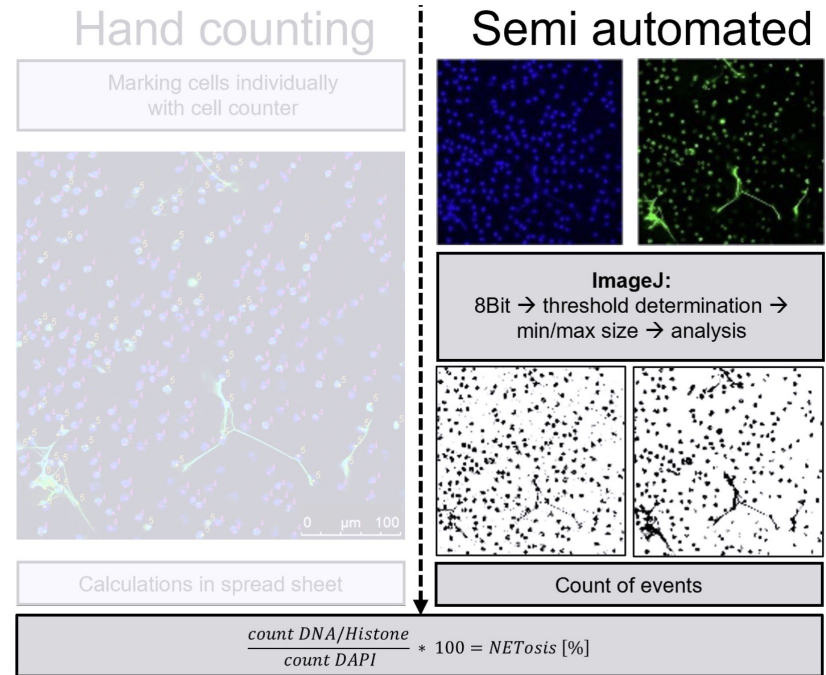
- Goal to reach: precision of hand counting



Baseline models

Earlier attempts to automatise the NET counting

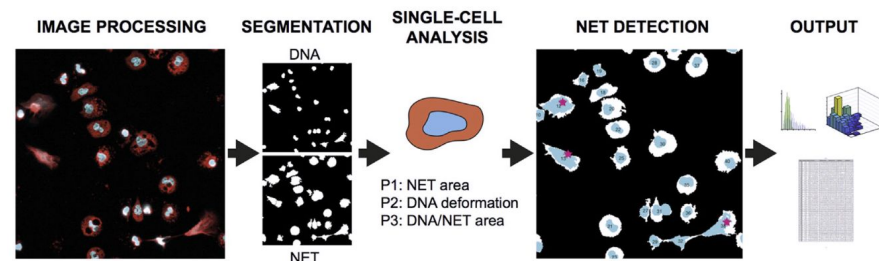
- Goal to reach: precision of hand counting
- Baseline models:
 - Semi automated with ImageJ



Baseline models

Earlier attempts to automatise the NET counting

- Goal to reach: precision of hand counting
- Baseline models:
 - Semi automated with ImageJ
 - Fully automated with NETQUANT (MATLAB)

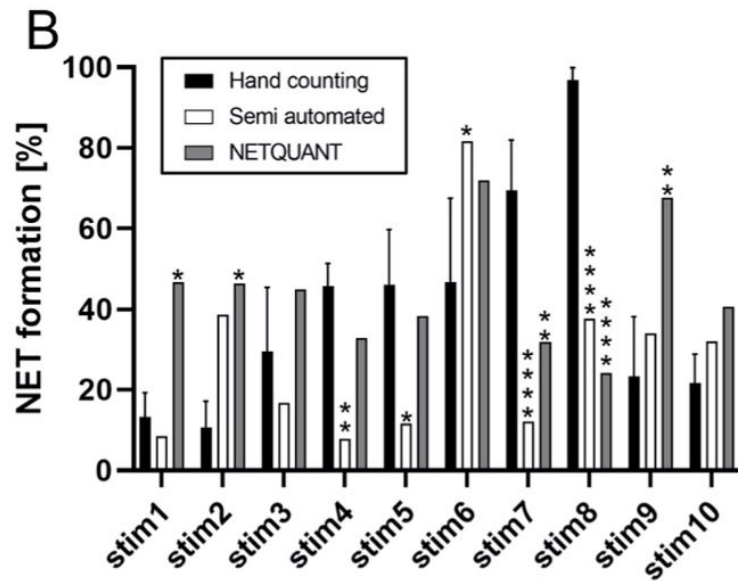


Mohanty et al., 2018

Baseline models

Earlier attempts to automatise the NET counting

- Goal to reach: precision of hand counting
- Baseline models:
 - Semi automated with ImageJ
 - Fully automated with NETQUANT (MATLAB)
- Overall poor performance of both models



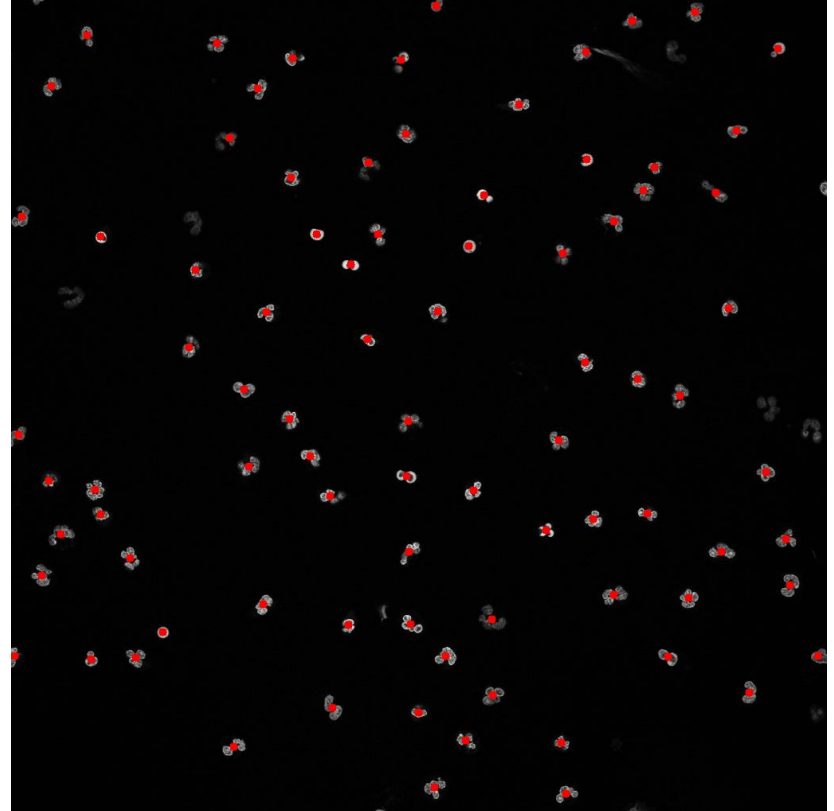
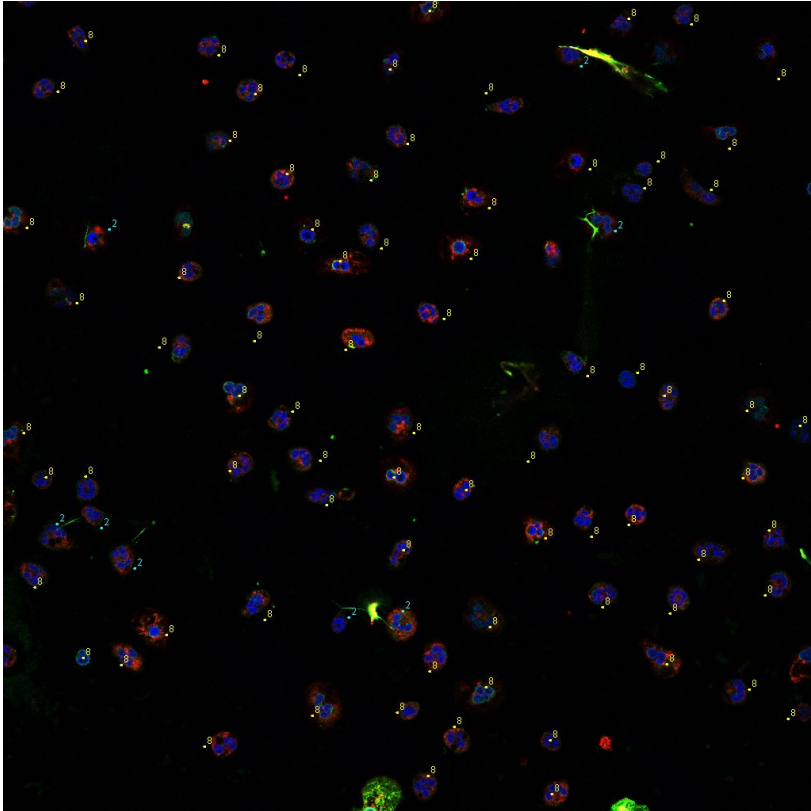
Henneck et al., 2023

Two modelling approaches

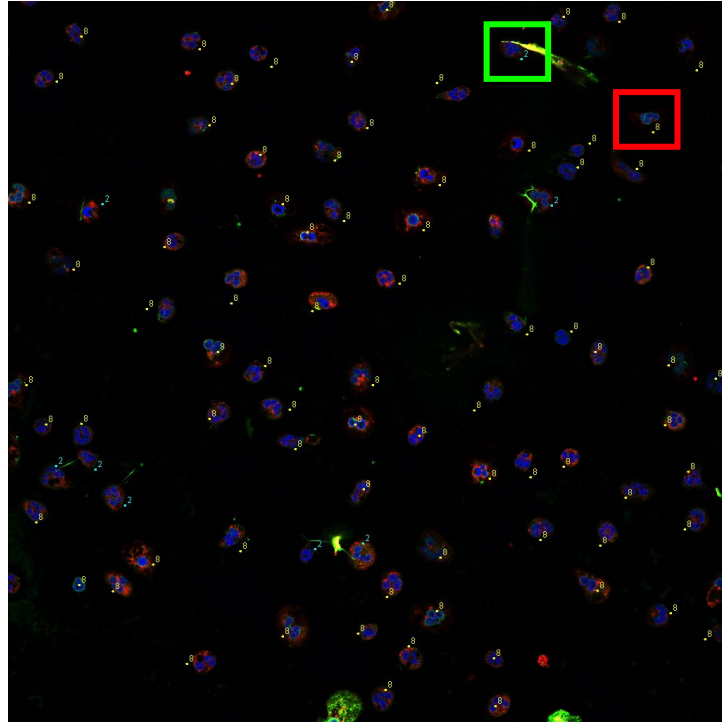
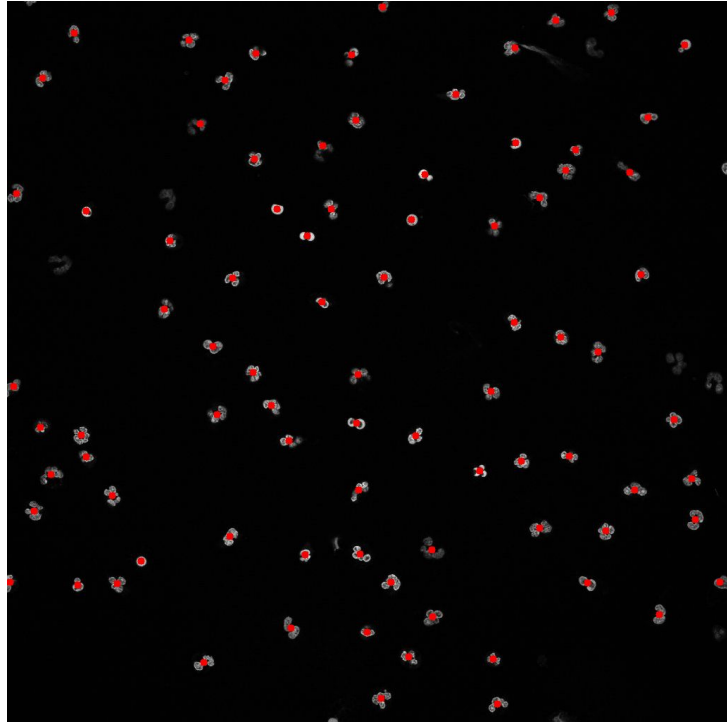
- Use recall or F1 score as main metrics to capture maximum positives (recall) and also taking false negatives and false positives into account (F1).
- Continue with two different approaches:
 - Subimaging
 - Segmentation

Model 1: Subimaging

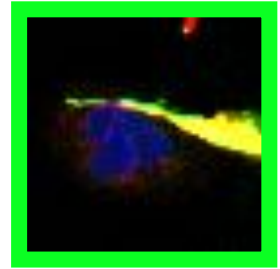
Get coordinates of nucleus centres in blue channel



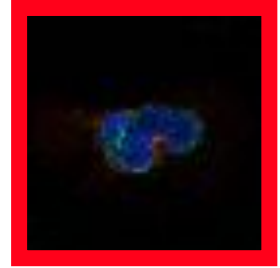
Use coordinates to make subimages of three color image



NET positive

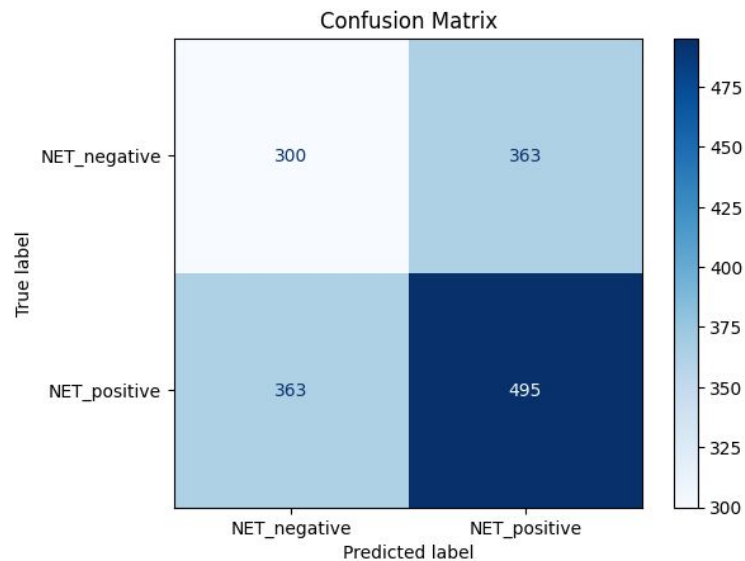


NET negative



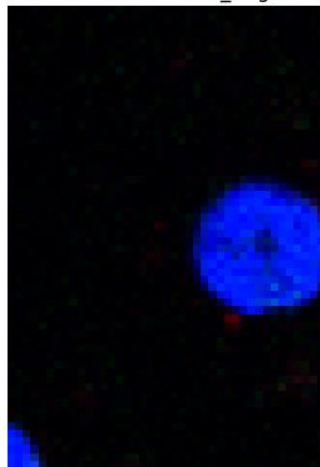
Best model: subimaging

Results: VGG16, F1 Metrics

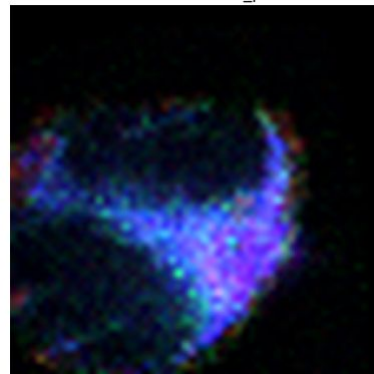


Classification Report:				
	precision	recall	f1-score	support
NET_negative	0.45	0.45	0.45	663
NET_positive	0.58	0.58	0.58	858
accuracy			0.52	1521
macro avg	0.51	0.51	0.51	1521
weighted avg	0.52	0.52	0.52	1521

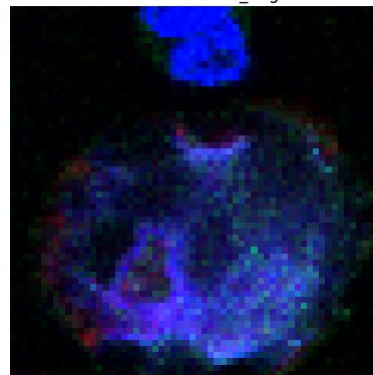
Predicted class: NET_negative



Predicted class: NET_positive



Predicted class: NET_negative



Model 2: Instance segmentation

Image segmentation using detectron2

Model pipeline

Original Image

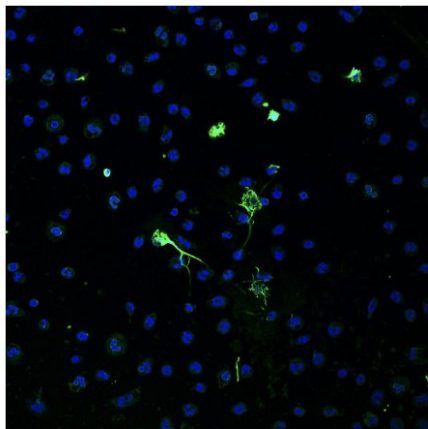
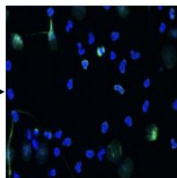
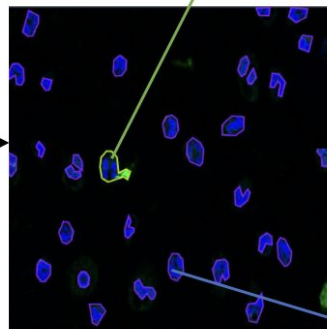


Image splitted



× 6

NETs



Instance
segmentation model



Detectron2

Non-Nets

Model Performance

Accumulating evaluation results...

DONE (t=0.02s).

Average Precision	(AP) @[IoU=0.50:0.95	area= all	maxDets=100]	= 0.571
Average Precision	(AP) @[IoU=0.50	area= all	maxDets=100]	= 0.759
Average Precision	(AP) @[IoU=0.75	area= all	maxDets=100]	= 0.671
Average Precision	(AP) @[IoU=0.50:0.95	area= small	maxDets=100]	= 0.328
Average Precision	(AP) @[IoU=0.50:0.95	area=medium	maxDets=100]	= 0.656
Average Precision	(AP) @[IoU=0.50:0.95	area= large	maxDets=100]	= 0.171
Average Recall	(AR) @[IoU=0.50:0.95	area= all	maxDets= 1]	= 0.061
Average Recall	(AR) @[IoU=0.50:0.95	area= all	maxDets= 10]	= 0.352
Average Recall	(AR) @[IoU=0.50:0.95	area= all	maxDets=100]	= 0.634
Average Recall	(AR) @[IoU=0.50:0.95	area= small	maxDets=100]	= 0.362
Average Recall	(AR) @[IoU=0.50:0.95	area=medium	maxDets=100]	= 0.712
Average Recall	(AR) @[IoU=0.50:0.95	area= large	maxDets=100]	= 0.263

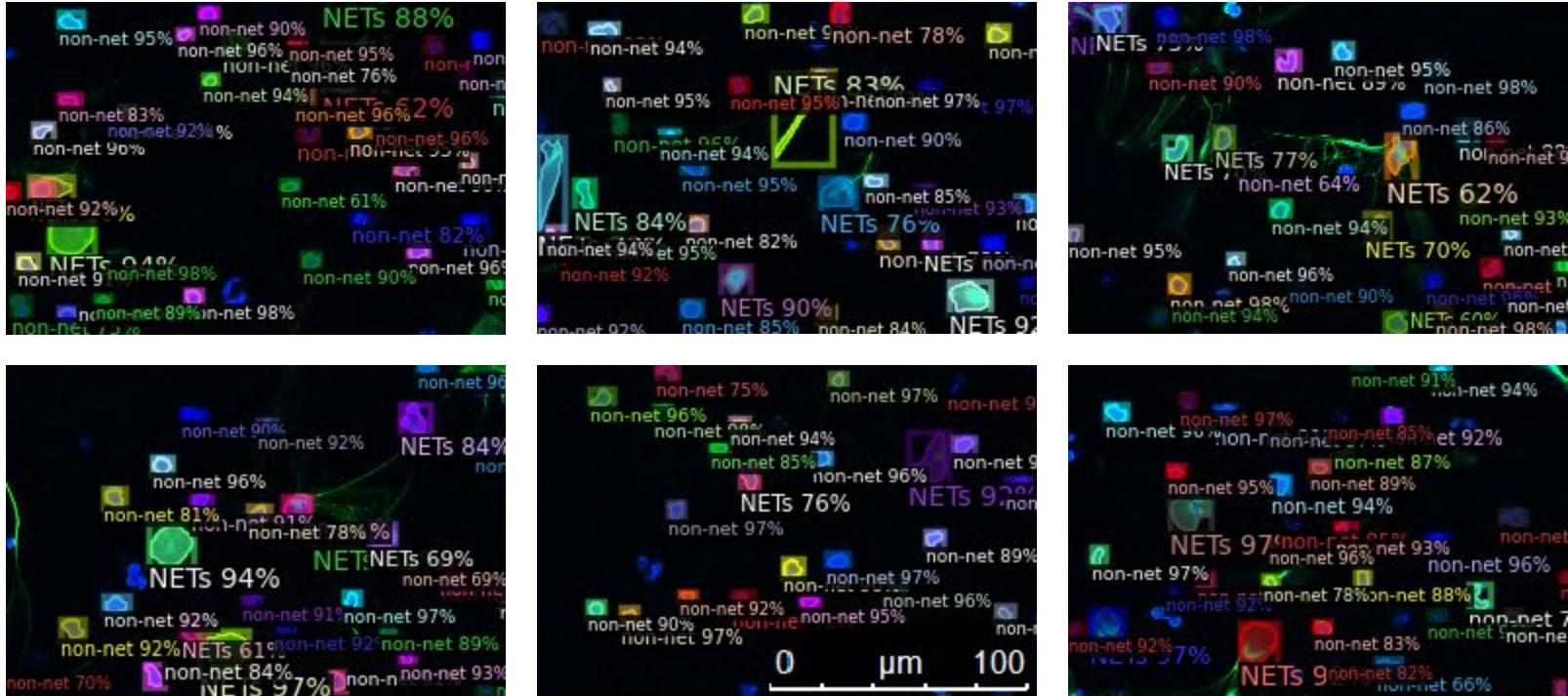
[06/06 14:38:16 d2.evaluation.coco_evaluation]: Evaluation results for bbox:

AP	AP50	AP75	APs	APm	APl
:-----:	:-----:	:-----:	:-----:	:-----:	:-----:
57.115	75.888	67.085	32.785	65.619	17.144

[06/06 14:38:16 d2.evaluation.coco_evaluation]: Per-category bbox AP:

category	AP	category	AP	category	AP
:-----:	:-----:	:-----:	:-----:	:-----:	:-----:
microscopic_images	nan	NETs	44.660	non-net	69.570

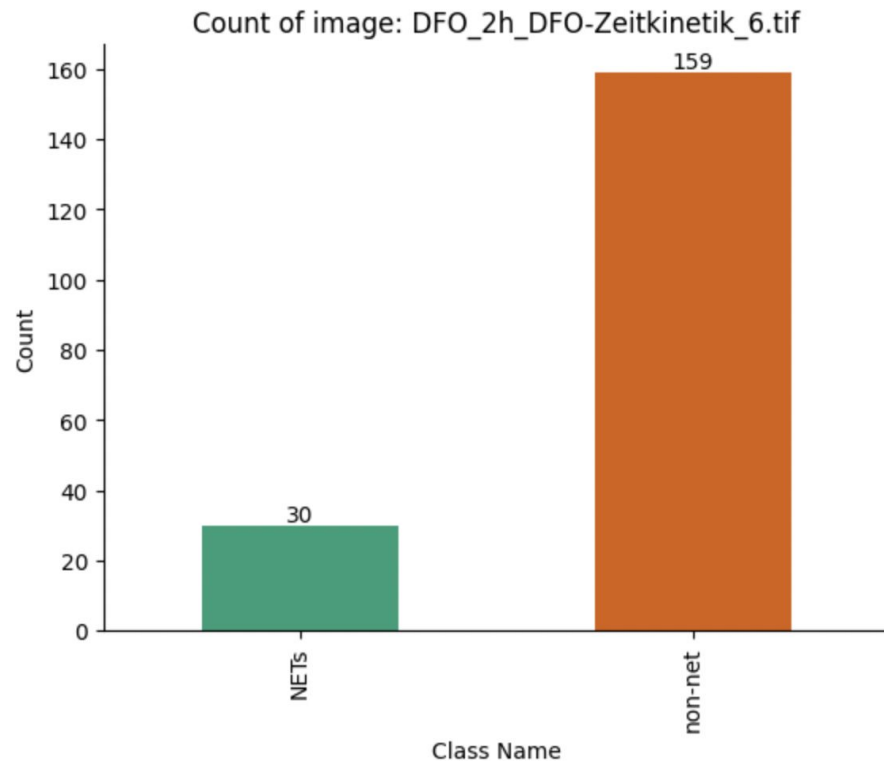
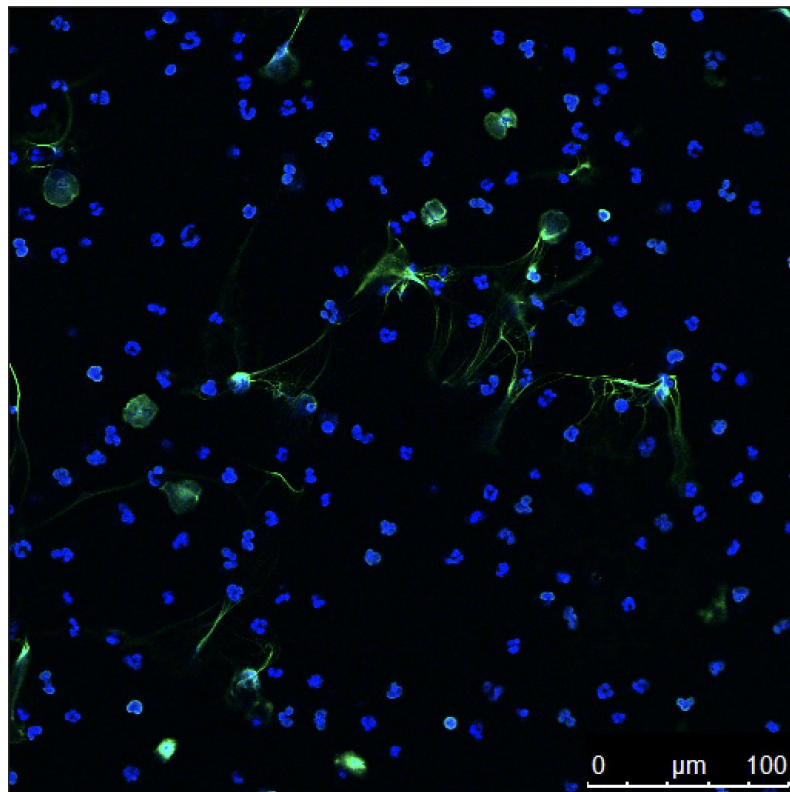
Best model: image segmentation



Instance segmentation of splitted image DFO_2h_DFO-Zeitkinetik_6

Best model: image segmentation

DFO_2h_DFO-Zeitkineitik_6



Outlook

- Model 1:
 - Work against overfitting (regularisation)
- Model 2:
 - Try yolo again with now available annotations
- Try DB scan as a third alternative
- Prepare webapp (turning into python script, streamlit)
- Potential problems:
 - Heterogeneity of the dataset and the NETs



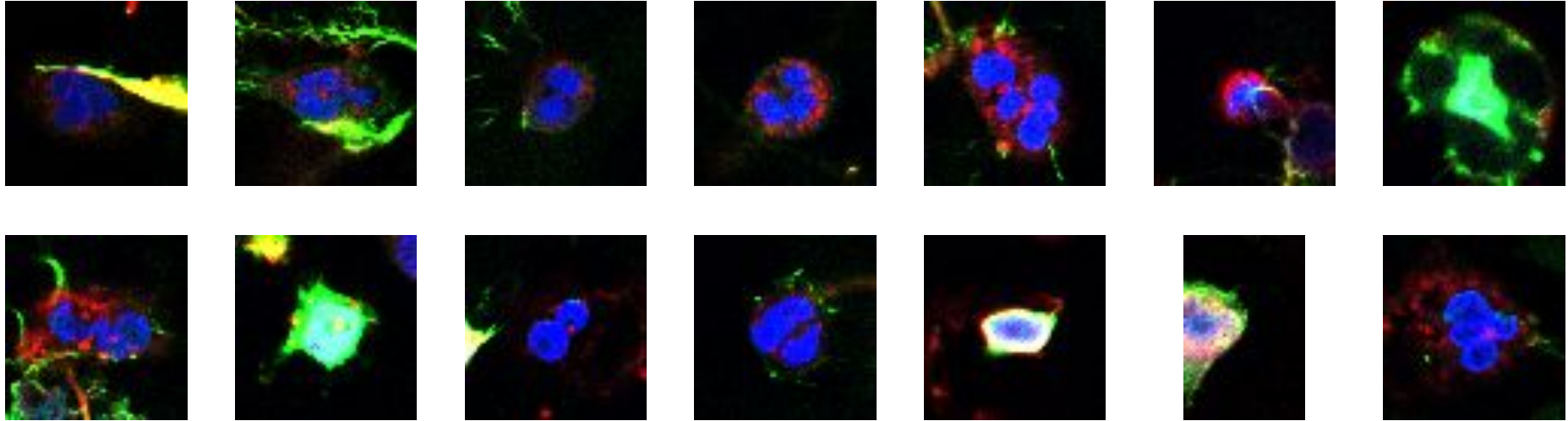
Thank you



Detailed image overview

	CTR stimulated	PMA stimulated	
RGB	12	7	19
GB	9	10	19
	21	17	38

NET heterogeneity



Problems with the NETs...

