

uc3m

# Machine learning - Deep learning

Applications to BiImage analysis

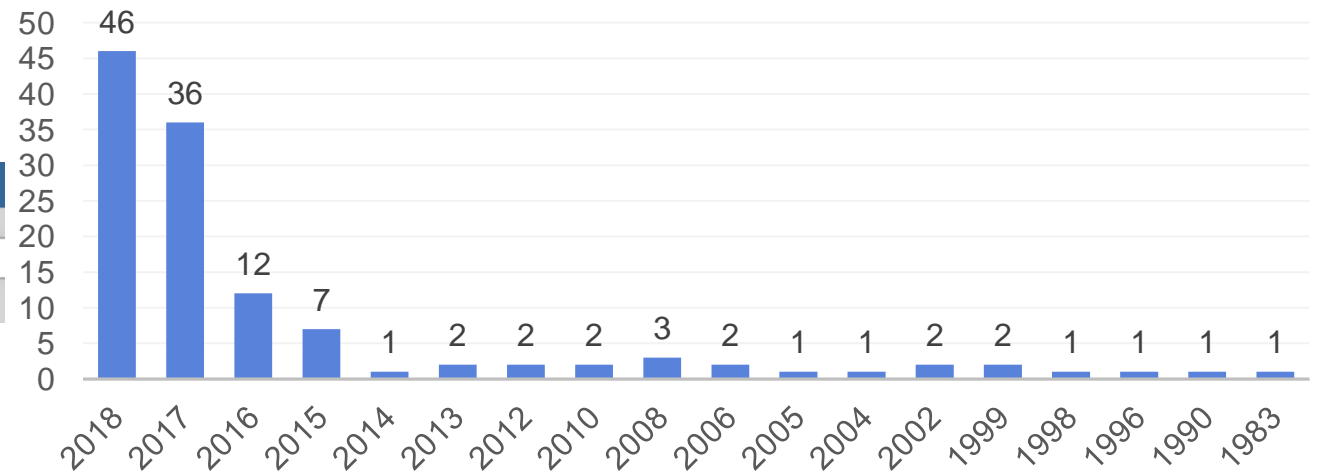
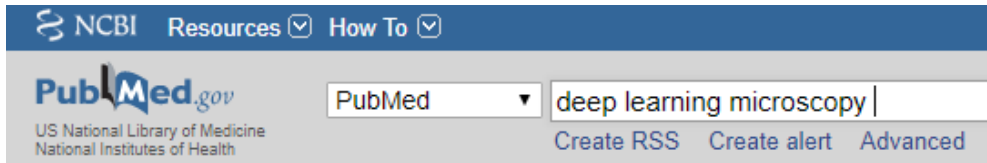
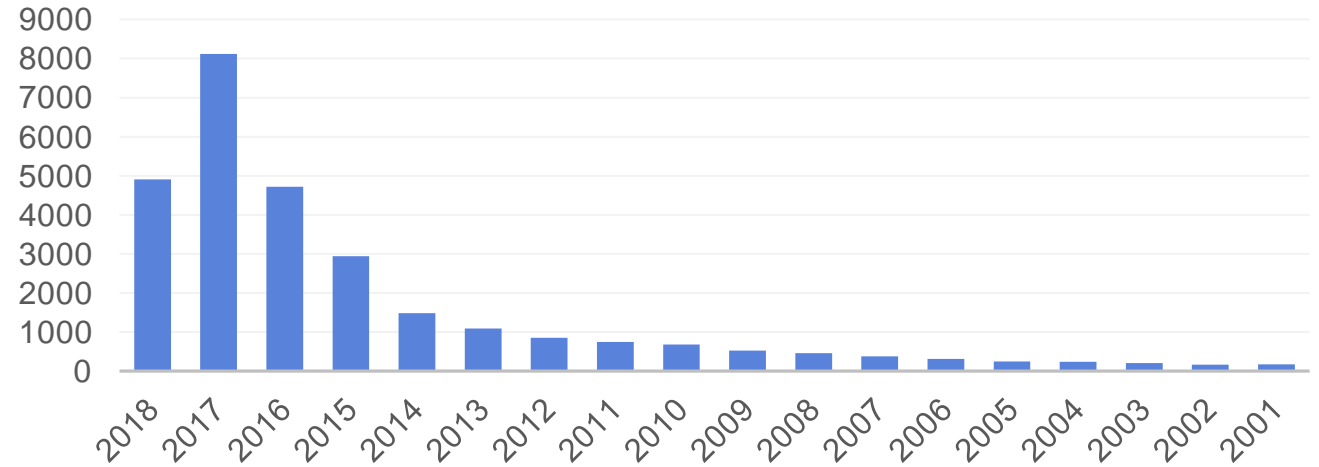
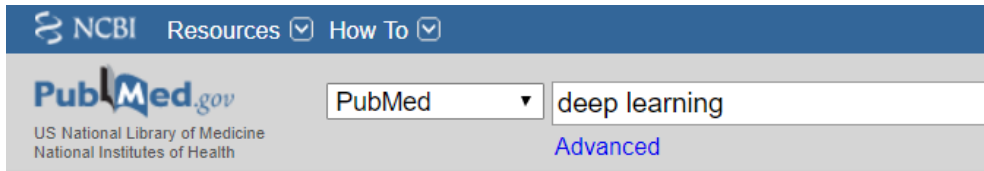
**SPAOM2018**

Estibaliz Gómez de Mariscal - Arrate Muñoz-Barrutia

Bioengineering and Aerospace Engineering department, Universidad Carlos III de Madrid, Spain

Instituto de Investigación Sanitaria Gregorio Marañón, Madrid, Spain

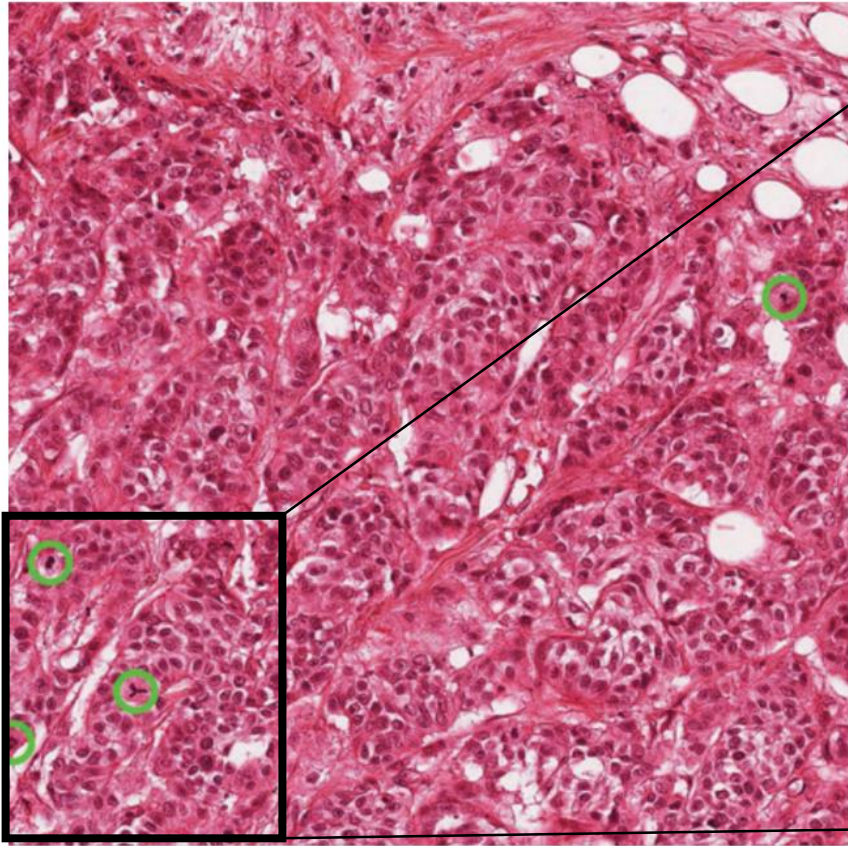




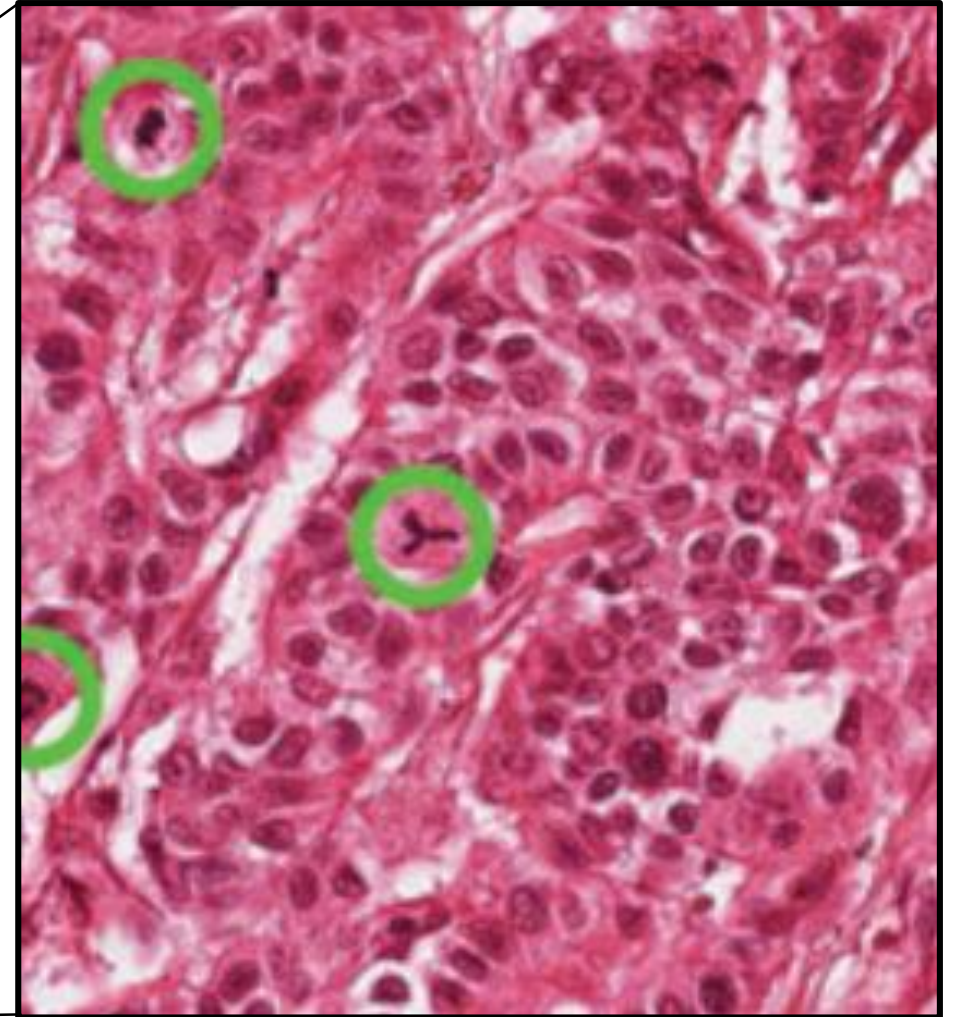
## Breakdown of published papers in Medical Image Analysis until January 2017:

- G. Litjens *et al.*, *A survey on deep learning in medical image analysis*. Medical image analysis 2017
- F. Xing, *et al.*, *Deep Learning in Microscopy Image Analysis: A Survey*. IEEE Transactions on Neural Networks and Learning Systems 2017

## Detection



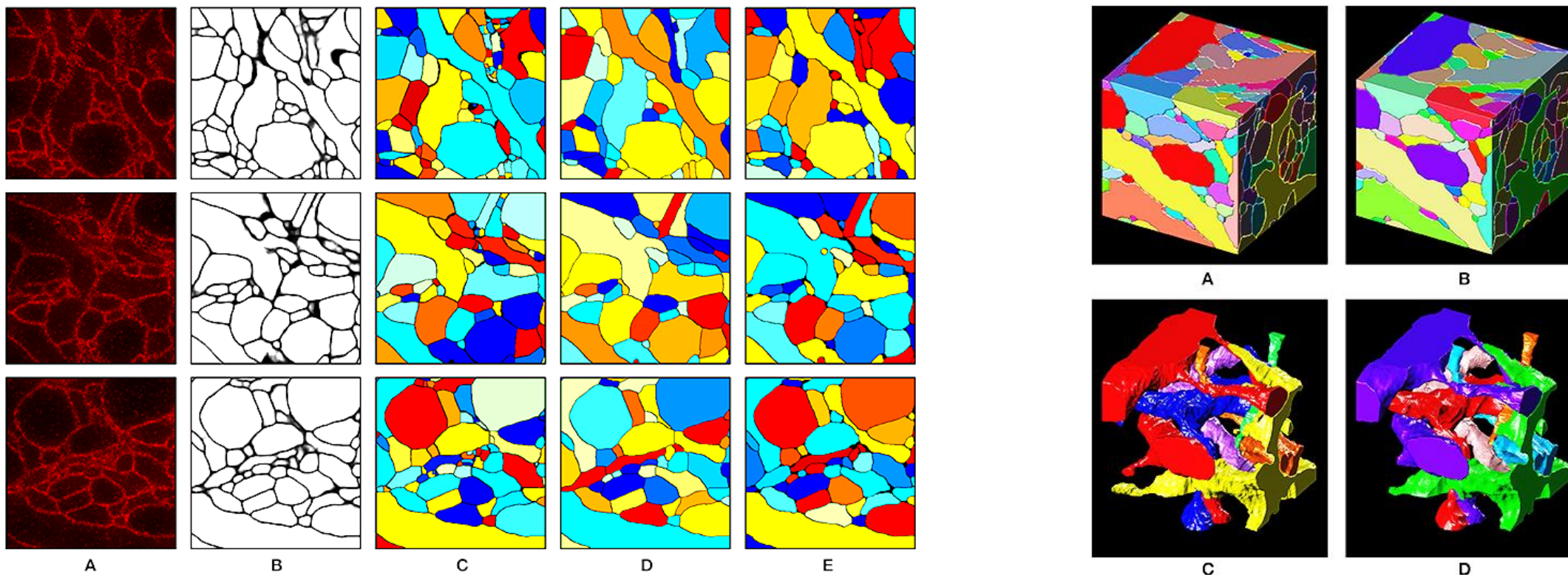
Mitosis  
detection





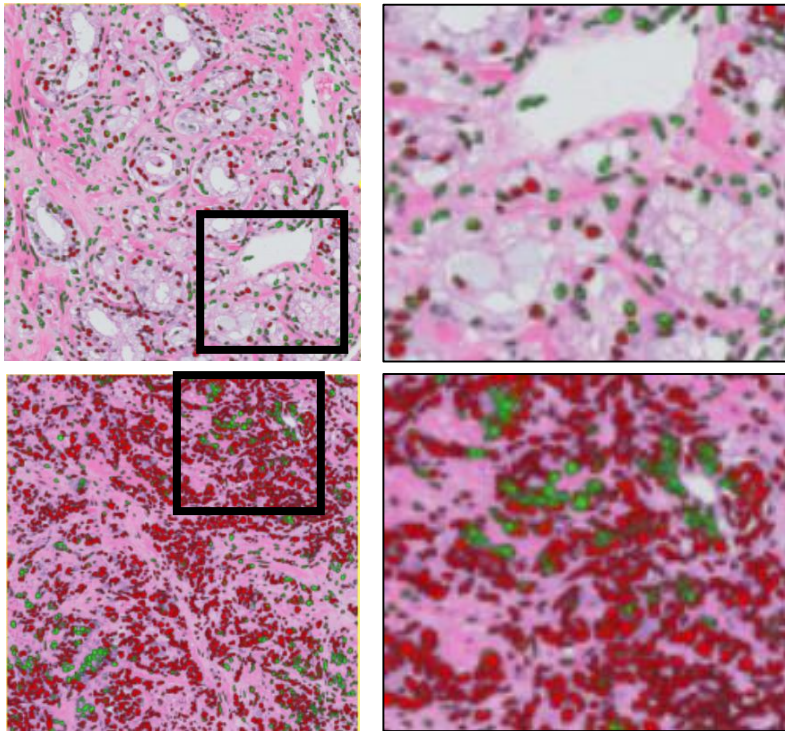
## Segmentation in 3D

Boundary prediction in 3D anisotropic Expansion Microscopy image



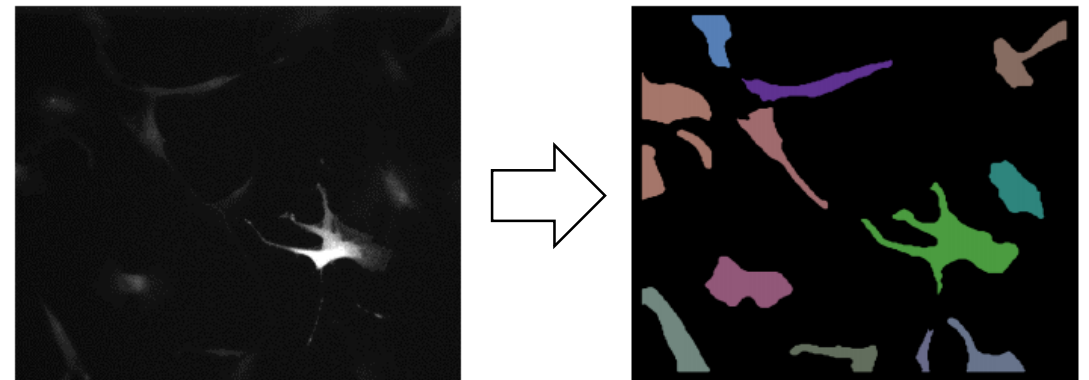
## Classification

Nucleus level segmentation and classification



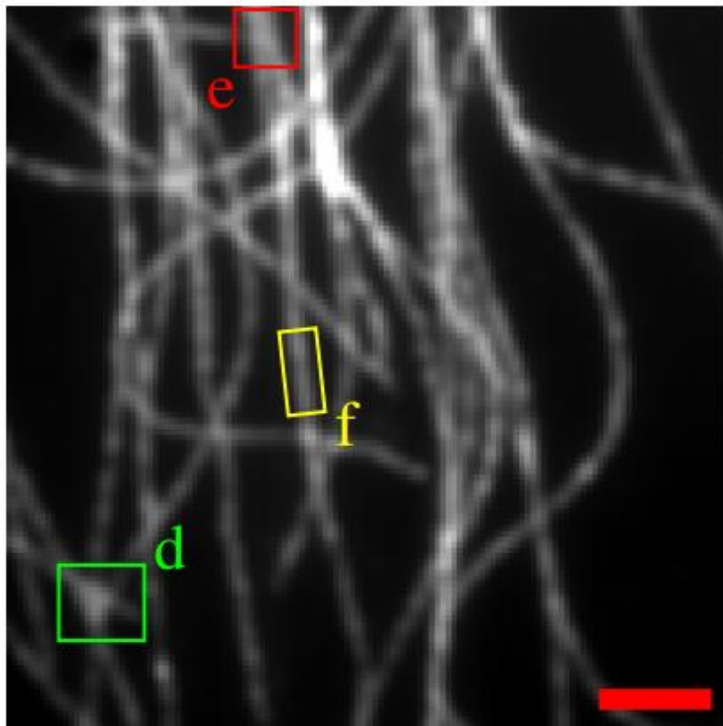
## Multi-object segmentation

Cell instance segmentation (Fluo-MS-C)

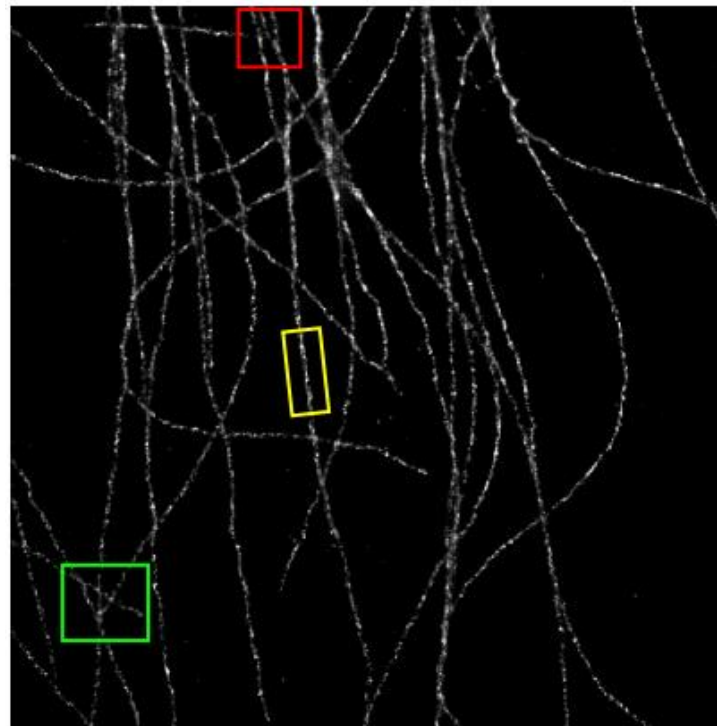
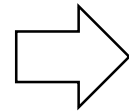


## Super-resolution imaging

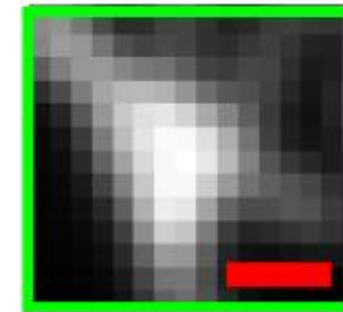
Experimentally measured microtubules.



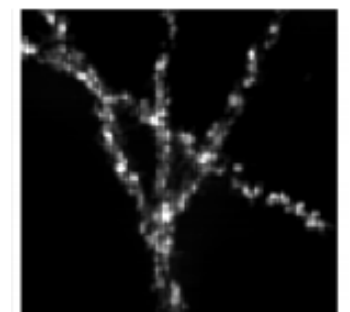
(a) Diffraction Limited



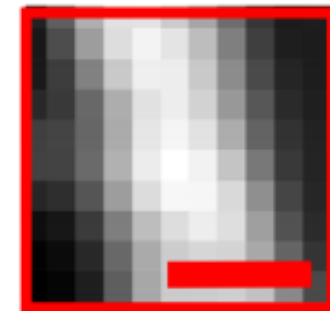
(c) Deep-STORM



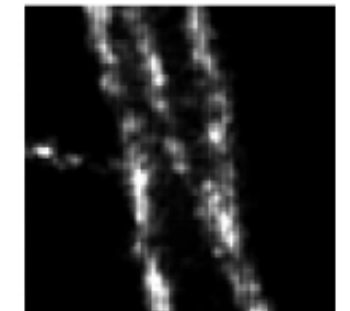
(d)



Deep-STORM



(e)

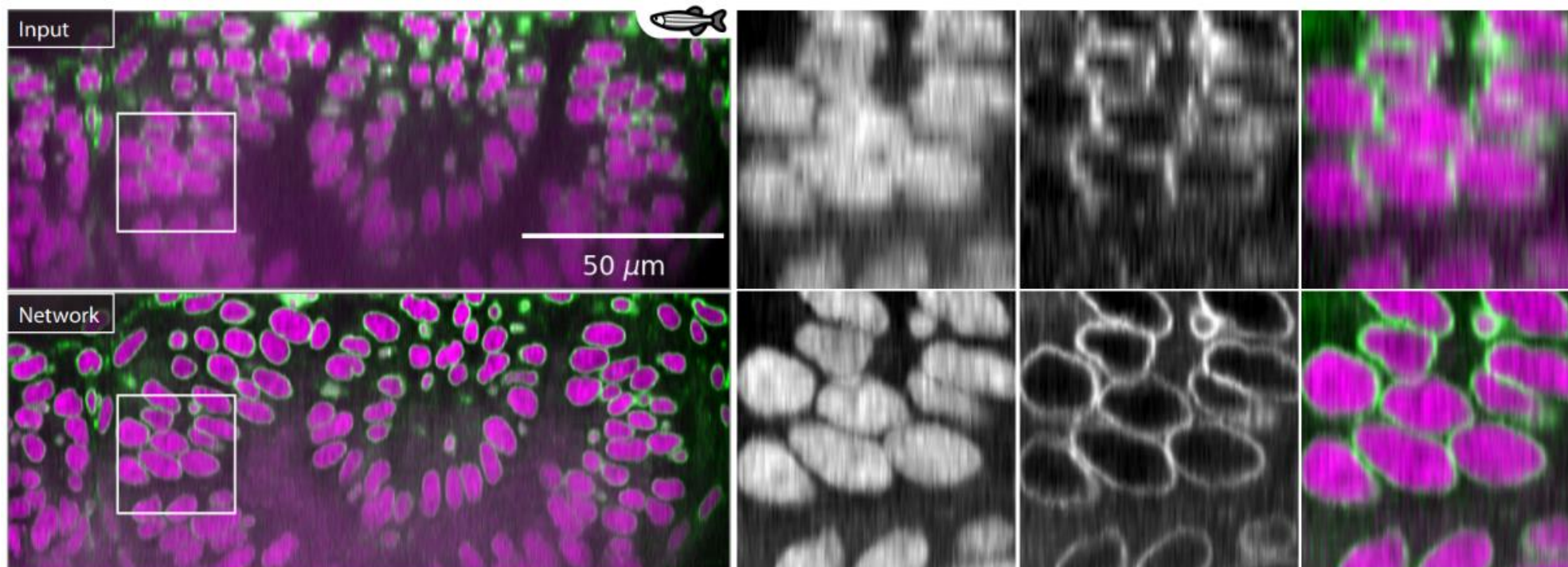


Deep-STORM



## Content-Aware image restoration

Zebrafish retina (nuclei and the nuclear envelope) in the anisotropic raw data (top row) and the isotropic restoration with deep learning.



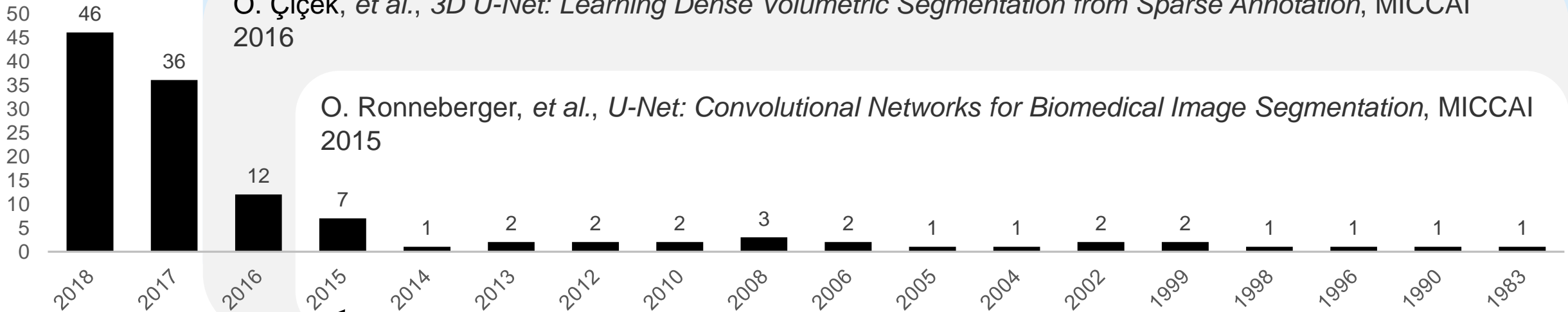
# Why deep learning?



## Effective deep learning architectures

O. Çiçek, et al., *3D U-Net: Learning Dense Volumetric Segmentation from Sparse Annotation*, MICCAI 2016

O. Ronneberger, et al., *U-Net: Convolutional Networks for Biomedical Image Segmentation*, MICCAI 2015



F. Chollet, et al., <https://keras.io> 2015

Deep Learning toolbox, Matlab 2015

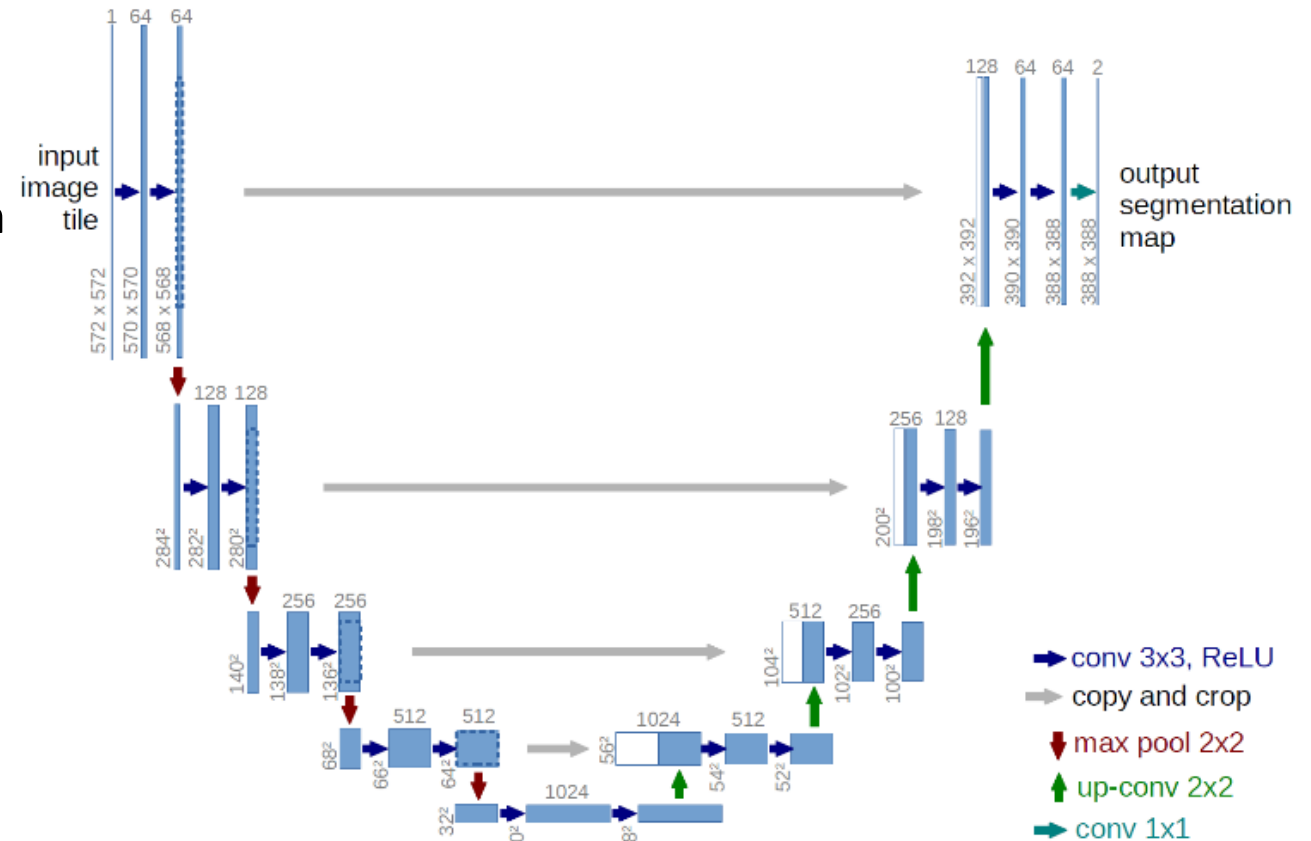
## User friendly libraries

## Advantages of U-net architecture

### I. Image processing

- Contracting path extracts high dimension features  $\rightarrow$  abstract analysis.
- Expanding path refines the processing.

### II. Data augmentation applied to medical image processing.



- ❑ DL methods fit thousands of parameters that allow to solve highly complex problems.
- ❑ Existence of sophisticated architectures ((3D)U-net, AlexNet, Mask R-CNN, Fast R-CNN, LSTM)
- ❑ Processing time after training is about seconds.
- ❑ Neural network architecture can be built easily with current software (Python, Matlab, C++, R).

O. Ronneberger, *et al.*, *U-Net: Convolutional Networks for Biomedical Image Segmentation*, MICCAI 2015

O. Çiçek, *et al.*, *3D U-Net: Learning Dense Volumetric Segmentation from Sparse Annotation*, MICCAI 2016

A. Krizhevsky, *et al.*, *ImageNet Classification with Deep Convolutional Neural Networks*. NIPS 2012

K. He, *et al.*, *Mask R-CNN*, arXiv 2018

R. Girshick, *Fast R-CNN*, IEEE ICCV 2018

S. Hochreiter, *et al.*, *Long short-term memory*, Neural computation 1997



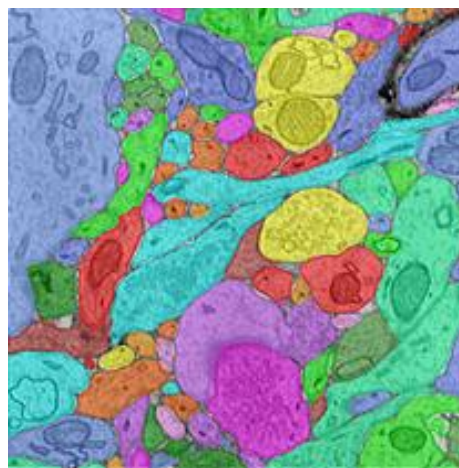
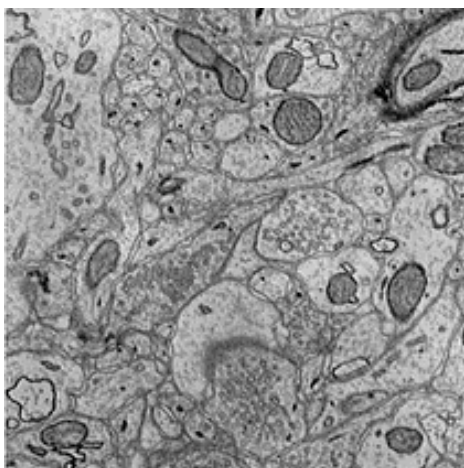
# What do you need?

The problem to solve by machine learning techniques has to be well defined.

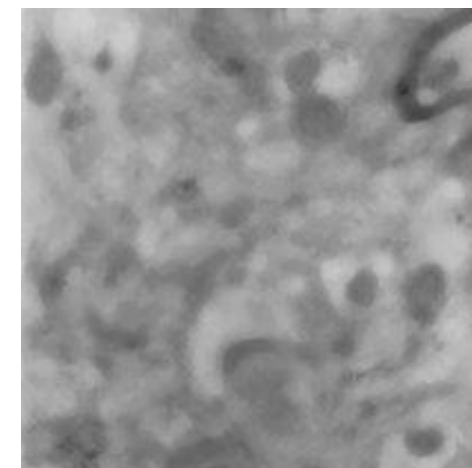
**Classification:** the number of classes has to be determined and their description cannot be ambiguous



**Segmentation:** The result of any manual annotation when performed twice by an expert, should always coincide.



High enough quality of data.



# What do you need?

To train our own model

## Technological infrastructure

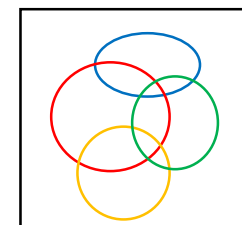
- Graphics processing units (GPU)
- Cloud computing (Google, Amazon)

**Data:** Ground Truth (GT) → manual annotations supervised by experts

- GT has to represent the real scenario of the problem.
- Large enough to train the model and evaluate it.



**NVIDIA Quadro P5000**



## Data augmentation

### Patching

### Geometrical transformations

Linear transformations (preserve shape)

- Rotation
- Translation



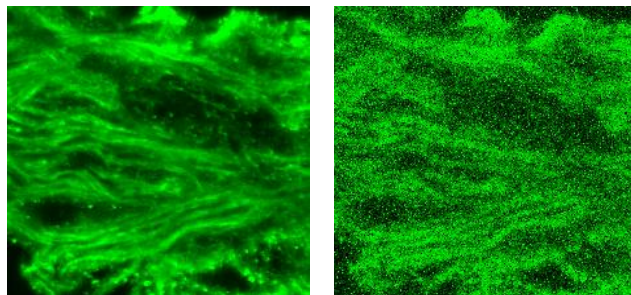
Non-linear (elastic) transformations (shape changes)

- Zooming
- Shearing



**Add artifacts: noise**

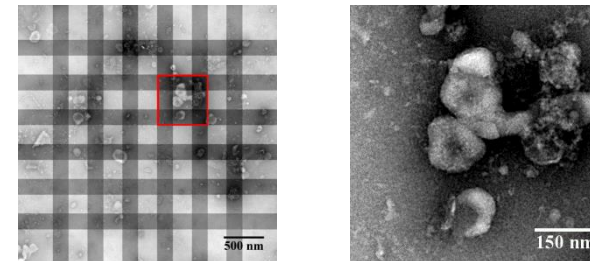
### Noising



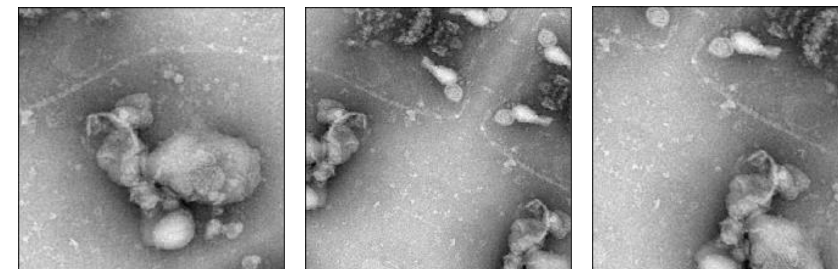
Original image

Noisy

### Patching



### Linear transformations

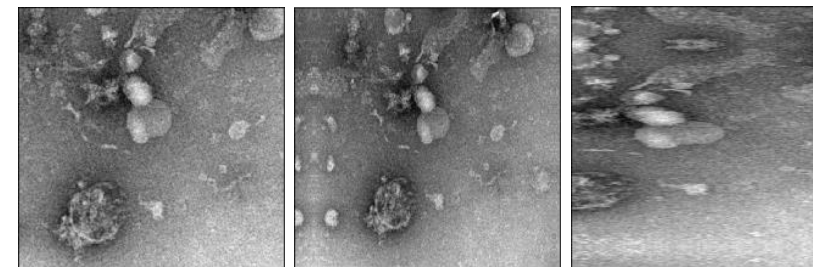


Original patch

Rotation + Shift

Rotation

### Non-linear transformations

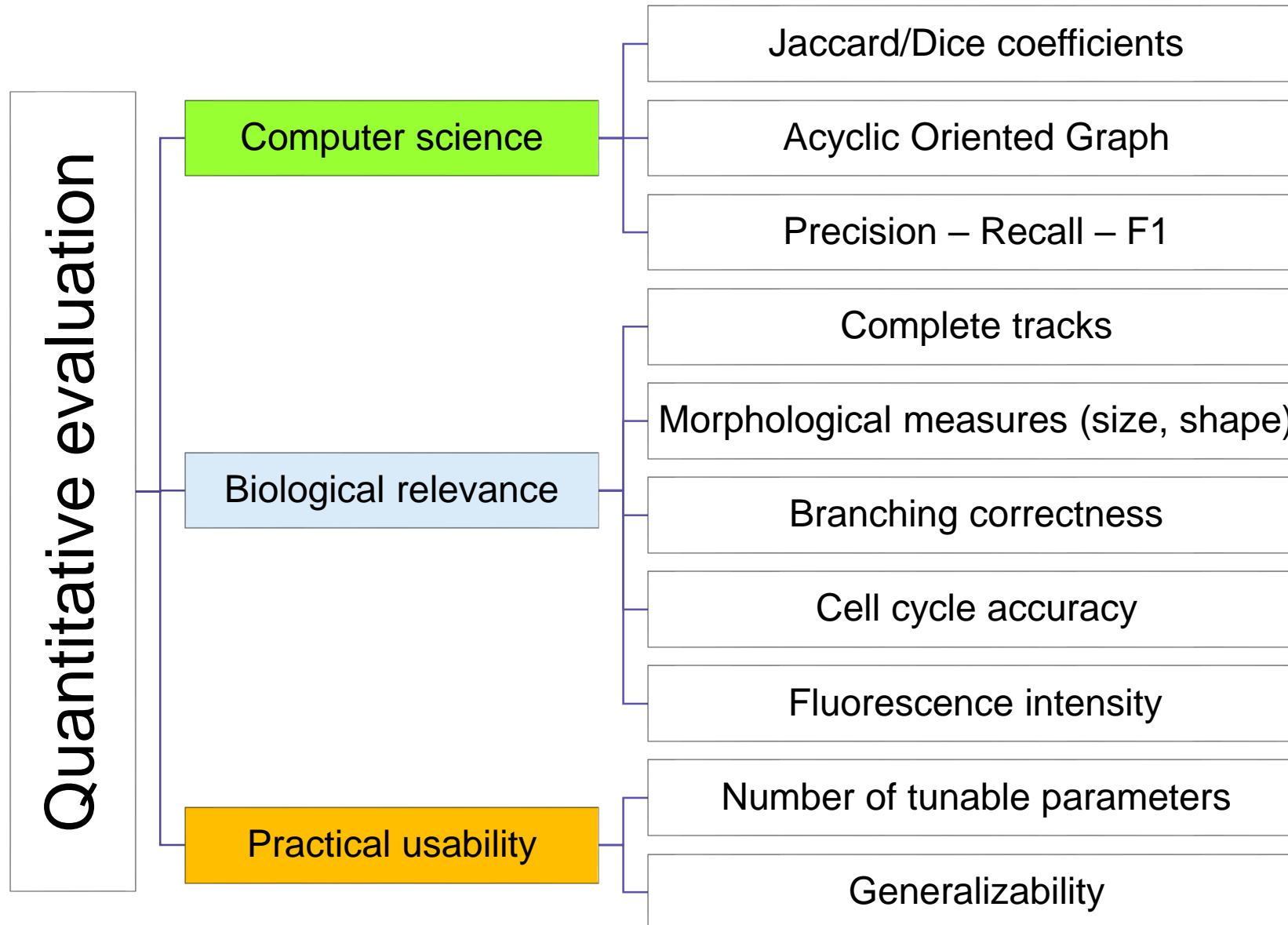


Original patch

Zoom

Shearing





## Data annotation

Slicer

Polygon-  
RNN++

ImageJ

## Data repositories

Kaggle

Cell tracking  
challenge

## Deep learning software

Python  
(Tensorflow,  
Keras)

Matlab

C++ (Caffe)

## User friendly software

ImageJ  
(U-net,  
CARE  
plugins)

Cell profiler  
upcoming

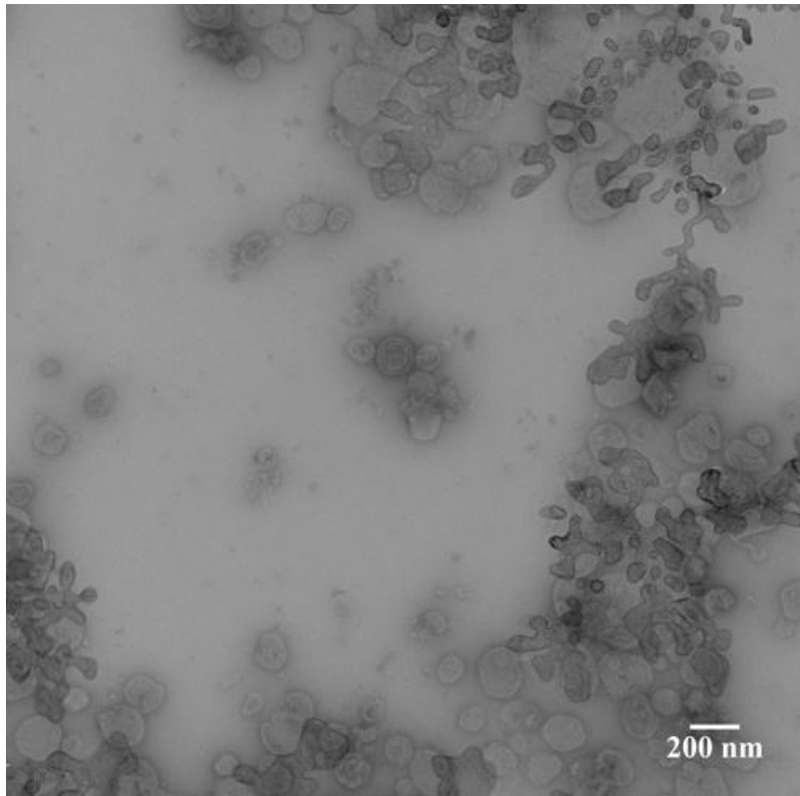
Ilastik  
upcoming

# Example of deep learning usage

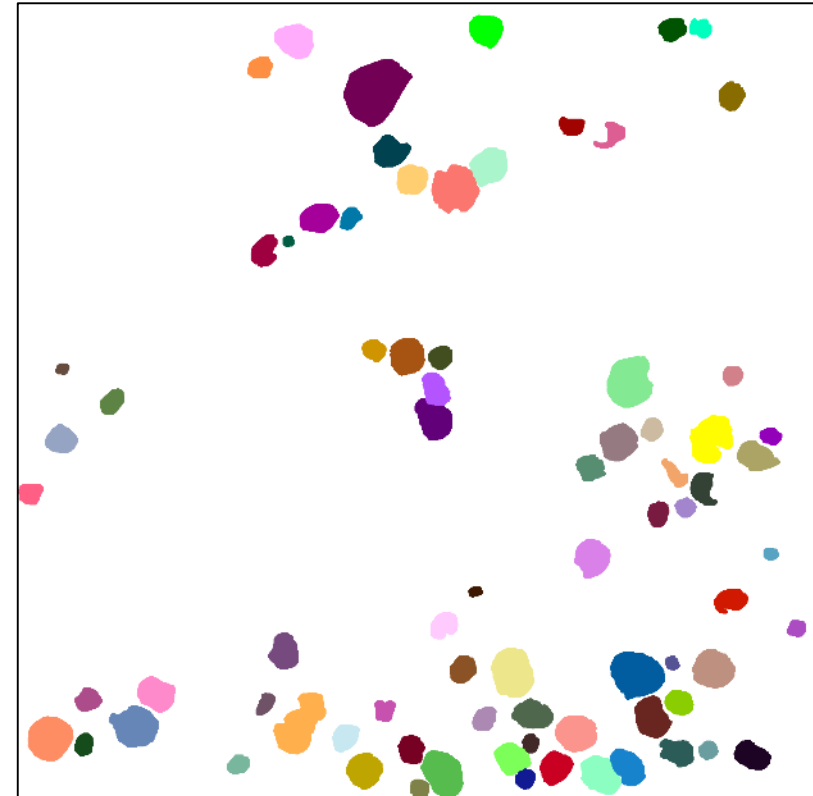


## Automatic exosomes segmentation in transmission electron microscopy images

Original image

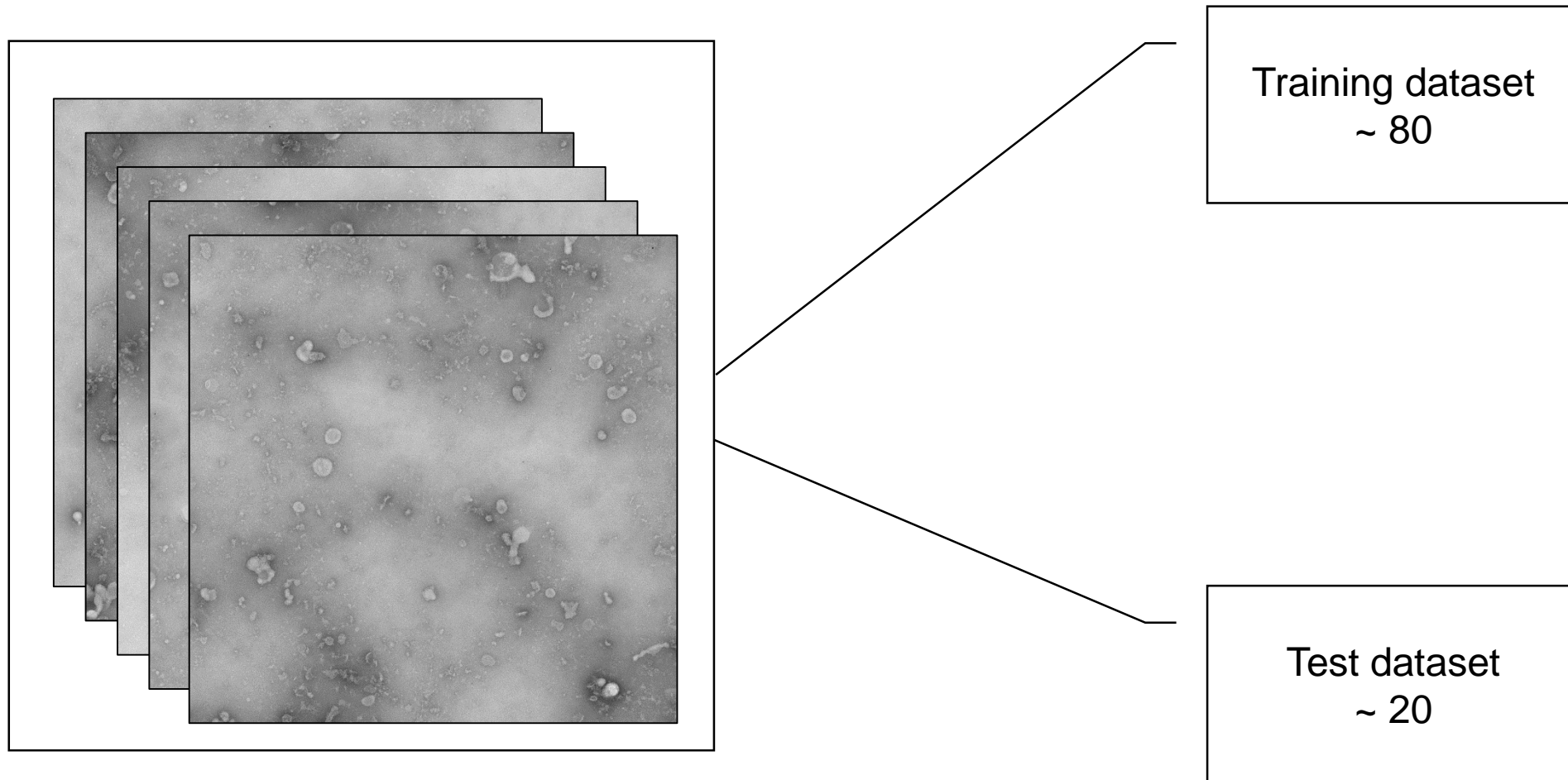


Ground truth

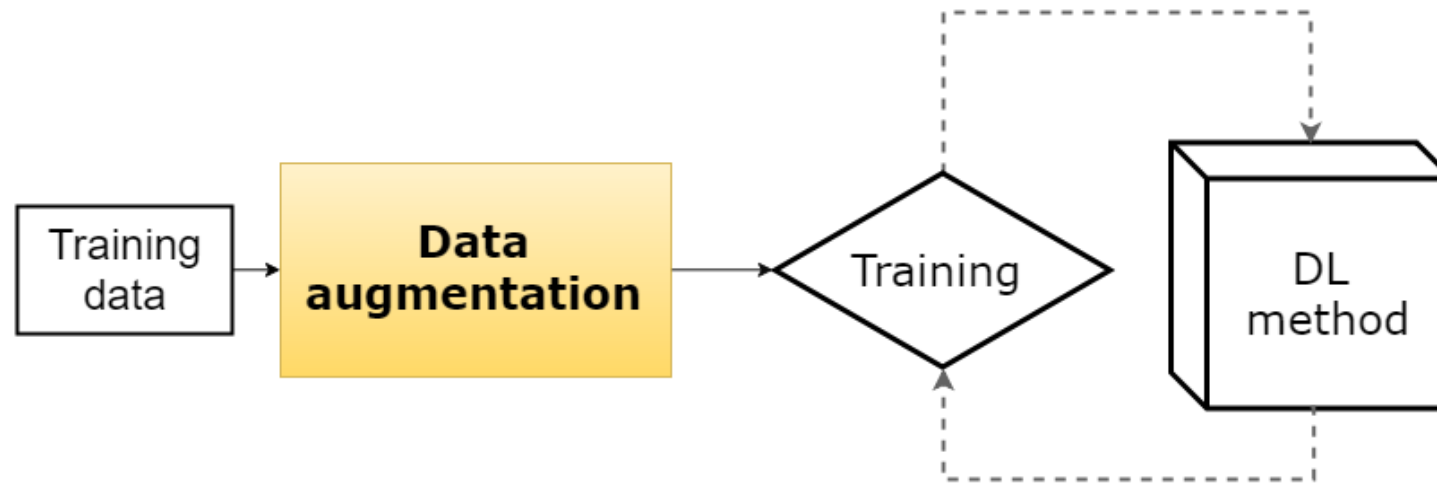


# Example of deep learning usage

Split data into training and test INDEPENDENT datasets



Increase training data and train your deep learning method



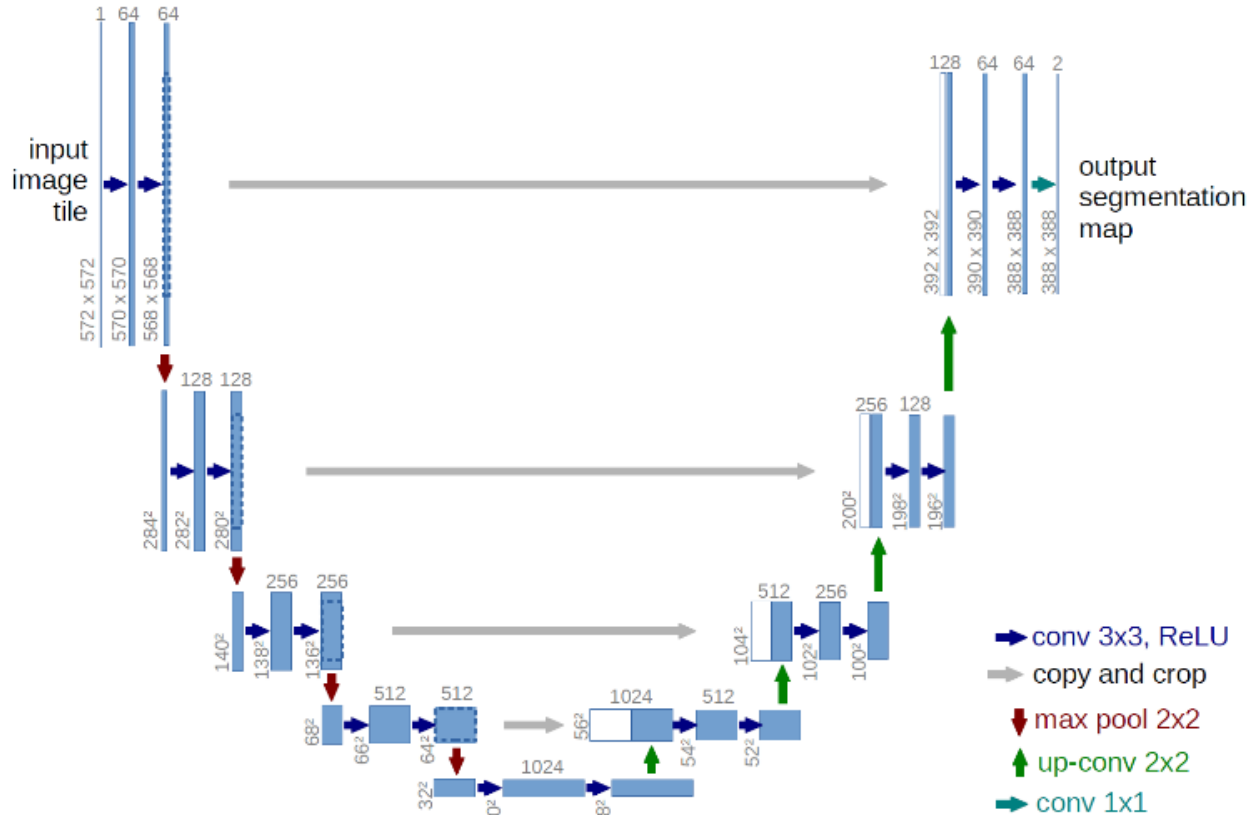
- DL architecture (number of layers and connections)
- Loss function
- Activation function
- Optimizer
- Batch size
- Batch normalization
- Drop out



Details of the method can be found in the manuscript



## Architecture



U-net convolutional neural network

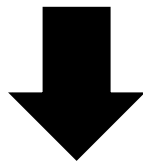
+

Residual layers

## Residual layers

Convolutional layer fits a non linear function  $G(x)$  to estimate the ground truth

$$y = G(x)$$

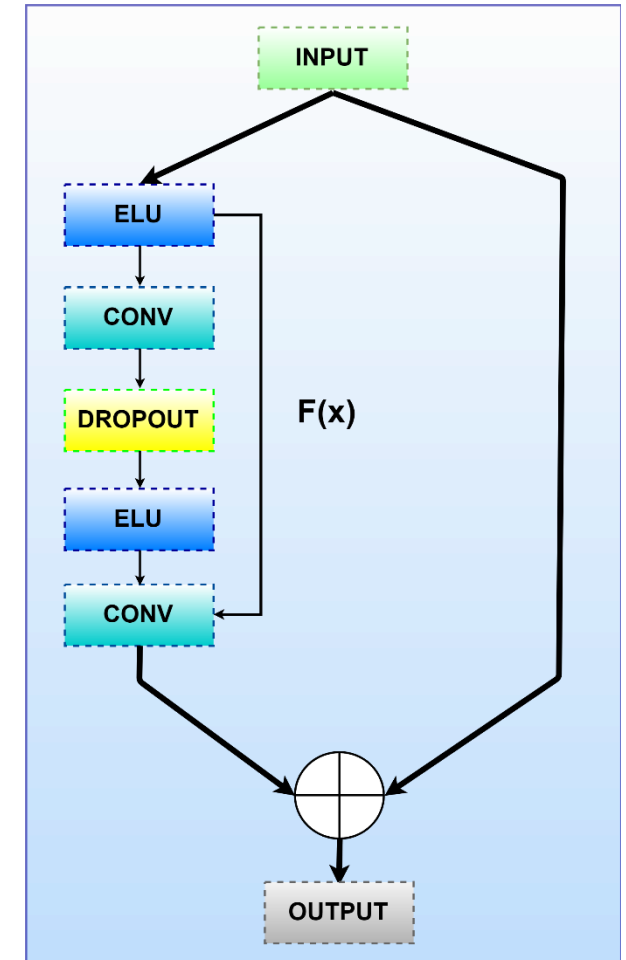


Residual layer include  $x$  as a possible solution to the problem

$$y = x + F(x)$$

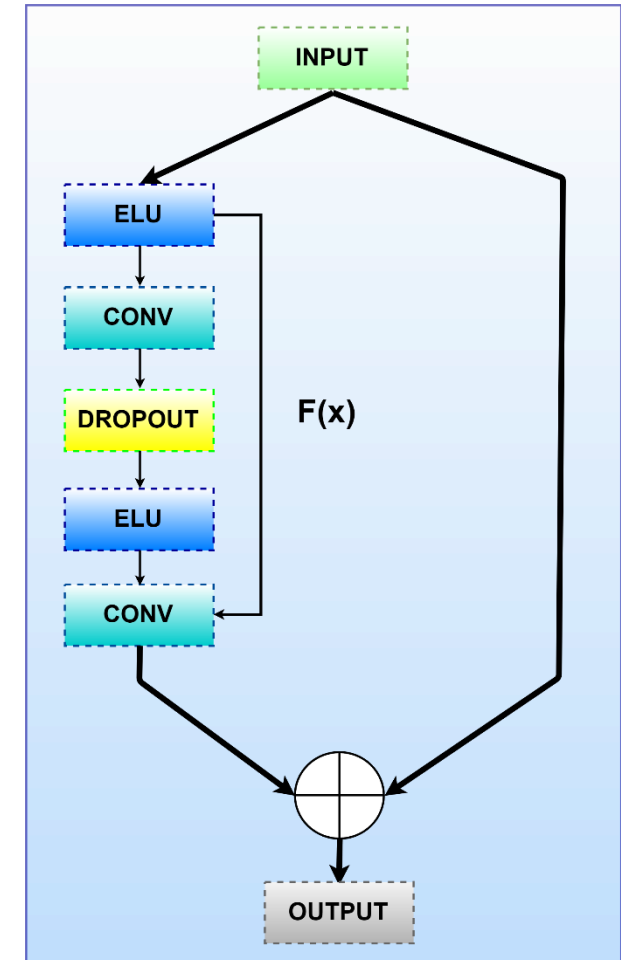
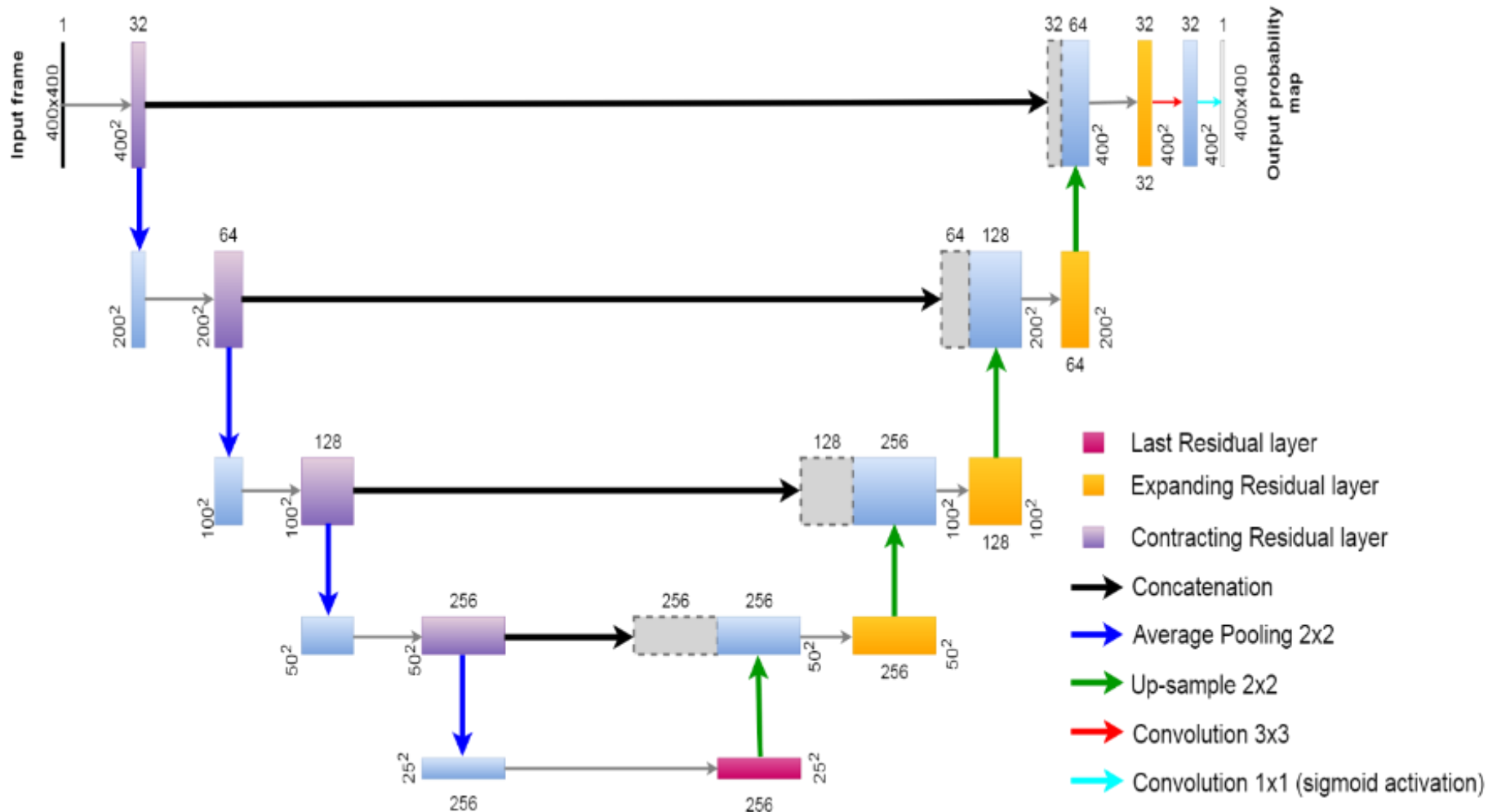
where  $F(x)$  is estimated by the residual block.

This approach limits the training error to  $x$  and prevents overfitting.





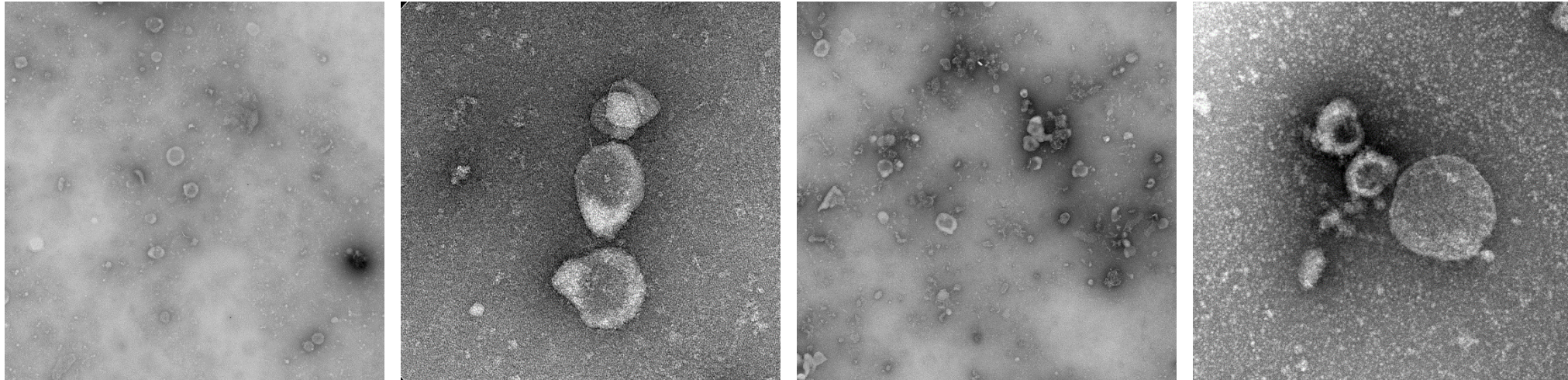
## Proposed architecture: Fully Residual U-net



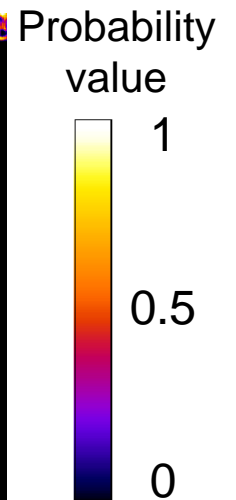
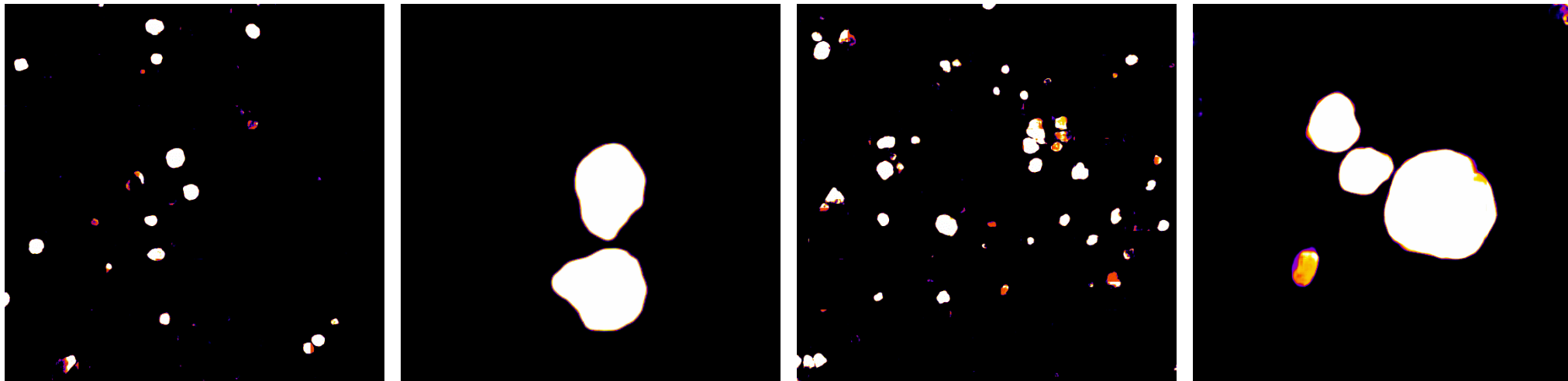
# Example of deep learning usage

Output of a convolutional neural network (CNN): probability maps

Input



Output

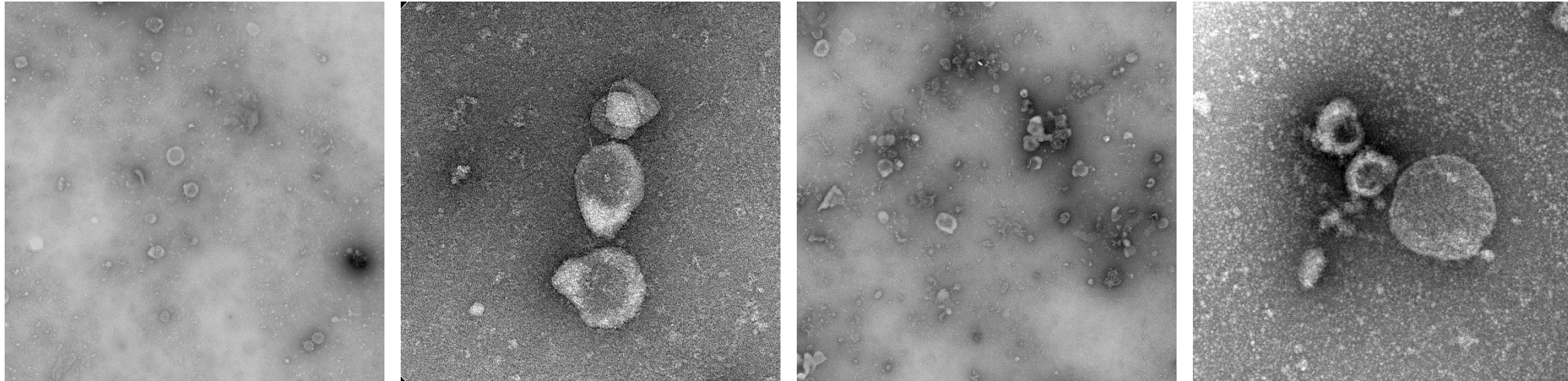




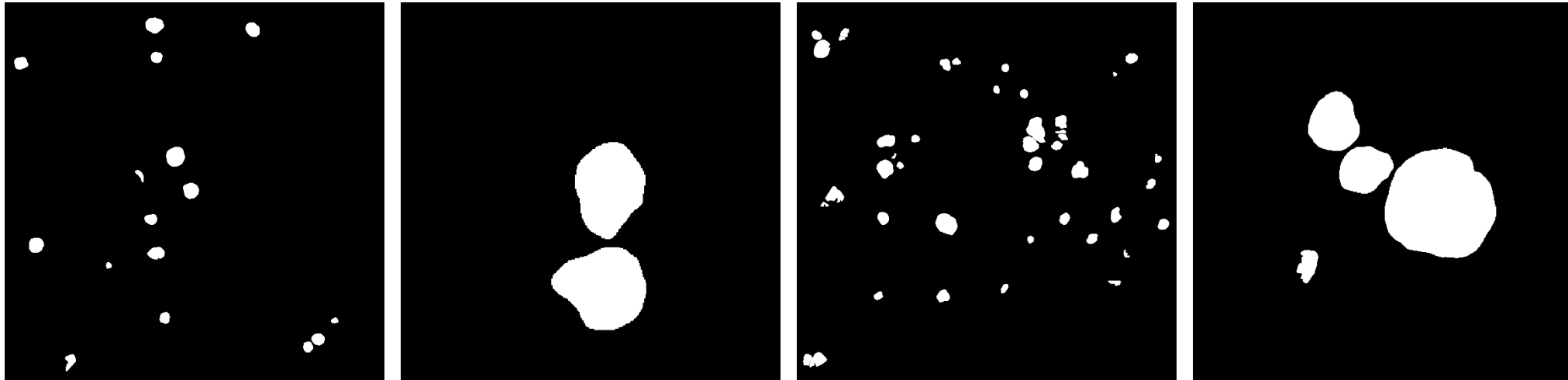
# Example of deep learning usage

Threshold probability maps to obtain binary masks

Input

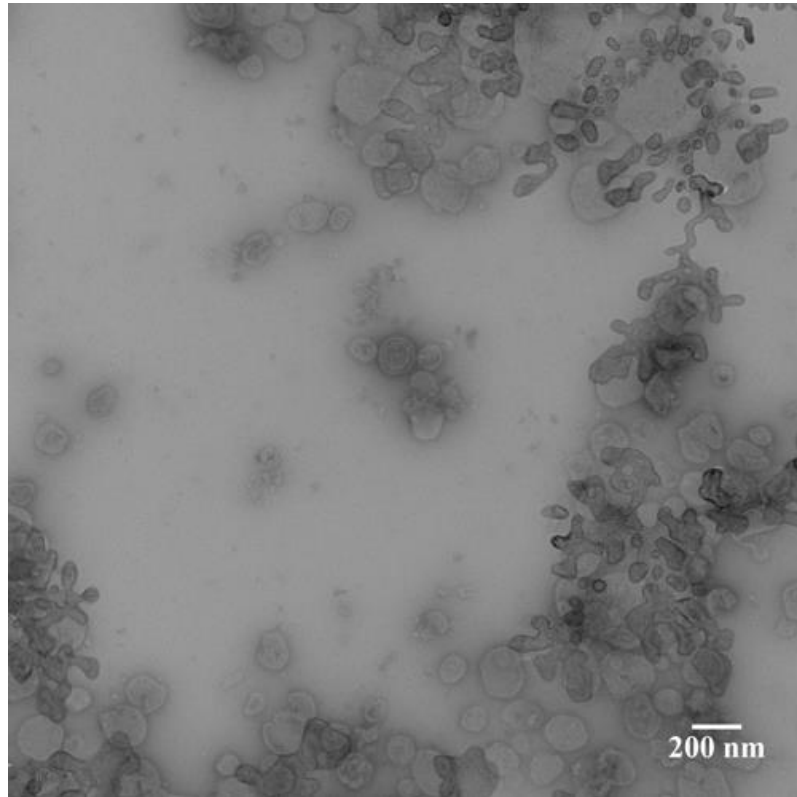


Binary  
mask

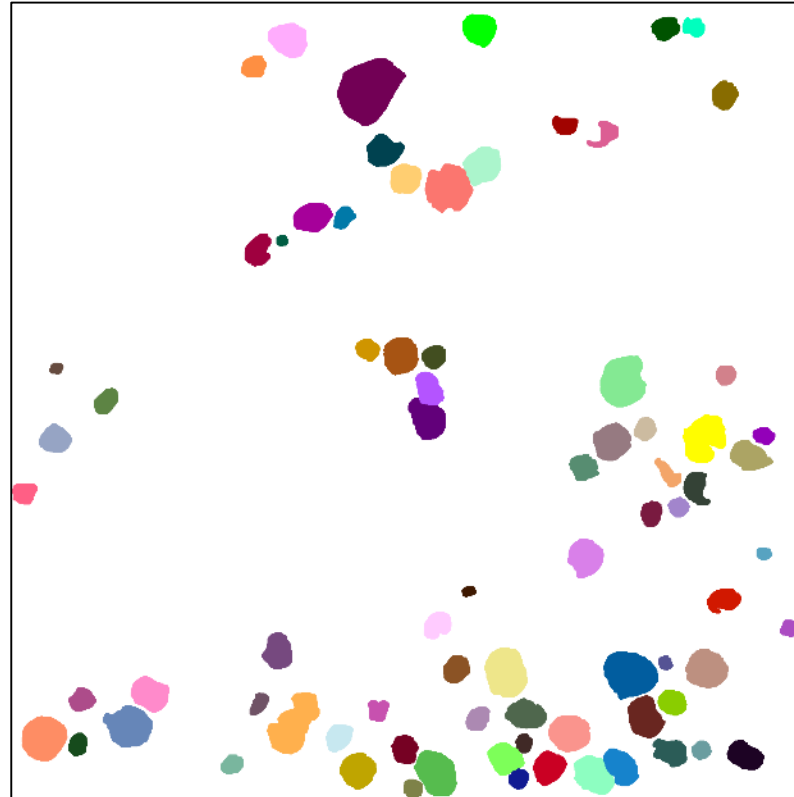


## Automatic exosomes segmentation in transmission electron microscopy images

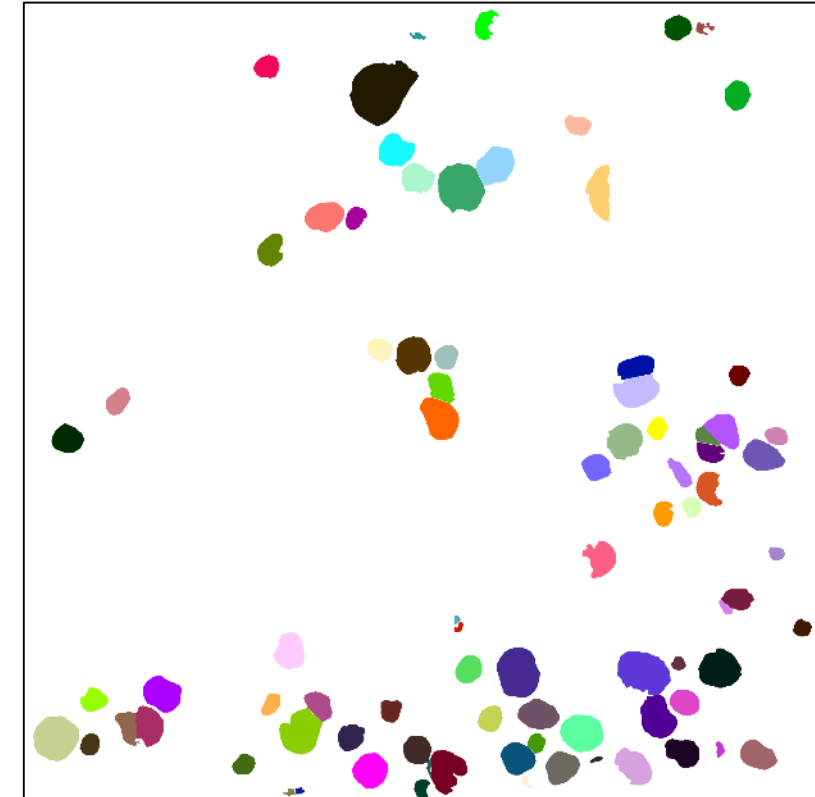
Original image



Ground truth



Fully Residual U-net result



## Accuracy measures and comparison with state of the art methods

Method	Computer Science		Biological relevance	
	Segmentation	Detection	Diameter	Roundness
	Jaccard Coeff.	Acyclic Oriented Graphs	$\delta_d$	$\delta_r$
FRU-net	<b>0.74</b>	<b>0.81</b>	<b>0.09</b>	0.11
U-net	0.68	0.77	0.11	0.13
Exosome Analyzer	0.19	0.17	0.17	<b>0.07</b>

$\delta_d$  and  $\delta_r$ : diameter and roundness similarity.

$S_i$  and  $GT_i$  correspond to the segmentation and ground truth of the exosome  $i$ .

$d$  and  $r$  to the diameter and roundness.

$$\delta_d = 1 - \frac{\min(d_{Si}, d_{GTi})}{\max(d_{Si}, d_{GTi})}$$

$$\delta_r = 1 - \frac{\min(r_{Si}, r_{GTi})}{\max(r_{Si}, r_{GTi})}$$

O. Ronnenberg, et al., *U-Net: Convolutional Networks for Biomedical Image Segmentation*, MICCAI 2015

K. Štěpka, et al., *Automatic Detection and Segmentation of Exosomes in Transmission Electron Microscopy*, ECCV 2016

P. Matula, et al., *Cell tracking accuracy measurement based on comparison of acyclic oriented graphs*, PloS one 2015



## FULLY AUTOMATIC EXOSOMES SEGMENTATION IN TRANSMISSION ELECTRON MICROSCOPY IMAGES

Gómez-de-Mariscal E.<sup>1</sup>, Maška M.<sup>2</sup>, Kotrbová A.<sup>3</sup>, Pospíchalová V.<sup>3</sup>, Matula P.<sup>2</sup>, Muñoz-Barrutia A.<sup>1</sup>

<sup>1</sup> Bioengineering and Aerospace Engineering Department, Universidad Carlos III de Madrid; Instituto de Investigación Sanitaria Gregorio Marañón, Madrid, Spain,

<sup>2</sup> Centre for Biomedical Image Analysis, Faculty of Informatics, Masaryk University, Brno, 602 00, Czech Republic and

<sup>3</sup> Department of Experimental Biology, Faculty of Science, Masaryk University, Brno, 611 37, Czech Republic

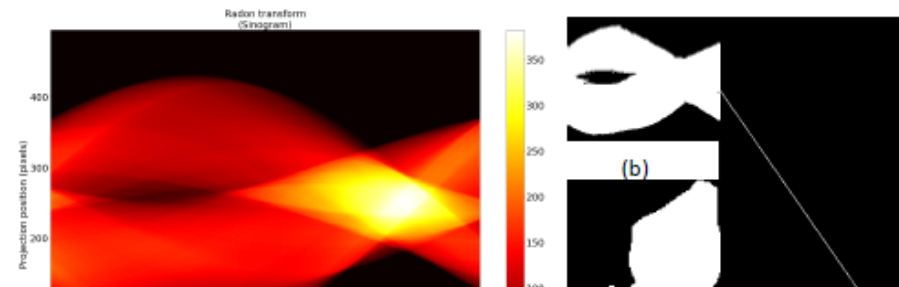
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Web: <http://image.hggm.es>

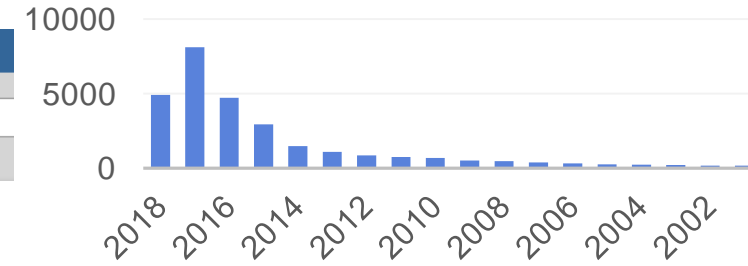
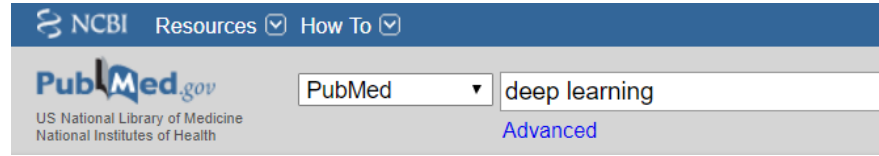
### ABSTRACT

Exosomes are nano-scale cell-derived extracellular vesicles, involved in the intercellular communication. Exosomes quantification is currently done manually by biologists and its automation will help them to remarkably progress in their research. We present the Fully Residual Unet for the segmentation of exosomes in Transmission Electron Microscope images and the Radon transform properties to separate clusters. An accuracy over 80% and 2s processing time for 2048x2048 pixels image are achieved.

### 3. Look for Radon transform local minimums to split exosomes clusters.



✓ Current situation

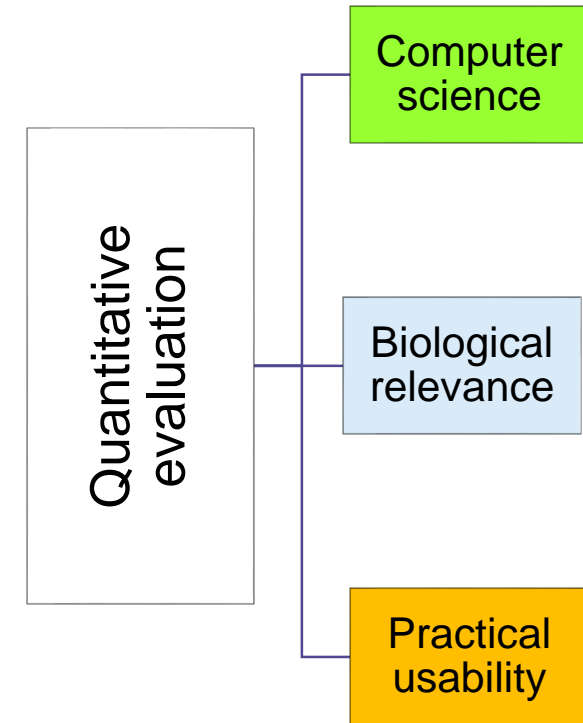


✓ Why deep learning?

✓ What do you need?



✓ Evaluation of obtained results



✓ Resources



✓ Example of deep learning usage

# Thank you!

## Machine learning - Deep learning

Applications to BioImage analysis

SPAOM2018



Estibaliz

Gómez de Mariscal

[estibaliz.gomez@uc3m.es](mailto:estibaliz.gomez@uc3m.es)



Link to slides:

<https://image.hggm.es/es/estibaliz-gomez>



Arrate

Muñoz Barrutia

[mamunozb@ing.uc3m.es](mailto:mamunozb@ing.uc3m.es)



Fundación BBVA

