

# Bio-Image Analysis, Programming, Biostatistics and Machine Learning for Computational Biology

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July 2023

# Programming

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- What is the output of these programs?

```
# we start with an empty list  
numbers = []  
  
# and add elements  
for i in range(0, 5):  
    numbers.append(i * 2)  
  
print(numbers)
```

```
numbers = [i * 2 for i in range(0, 5)]  
print(numbers)
```

```
▶ # Arrays
numbers = [0, 1, 2, 3, 4, 5, 6, 7, 8, 9]
print(numbers)
```

[0, 1, 2, 3, 4, 5, 6, 7, 8, 9]

- What do these programs output?

Starting at  
(including)

Ending at  
(excluding)

```
▶ subset = numbers[2:4]
print(subset)
```

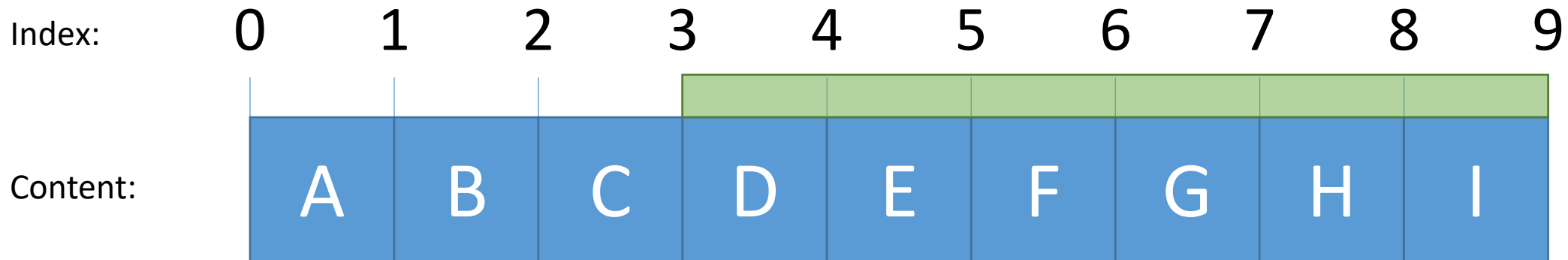
Step size

```
▶ subset_with_gaps = arr[1:8:2]
print(subset_with_gaps)
```

- What would be good comments in this code?

```
▶ #  
from skimage.io import imread  
blobs = imread("blobs.tif")  
  
#  
from skimage.filters import threshold_otsu  
threshold = threshold_otsu(blobs)  
binary_blobs = blobs > threshold  
  
#  
from skimage.measure import label  
labeled_blobs = label(binary_blobs)  
  
#  
import matplotlib.pyplot as plt  
fig, axs = plt.subplots(1, 3, figsize=(15,15))  
  
axs[0].imshow(blobs)  
axs[1].imshow(binary_blobs)  
axs[2].imshow(labeled_blobs, cmap=label_cmap)
```

```
data = ['A', 'B', 'C', 'D', 'E', 'F', 'G', 'H', 'I']
```



What's the output of

```
data[3:]
```

?

ABC

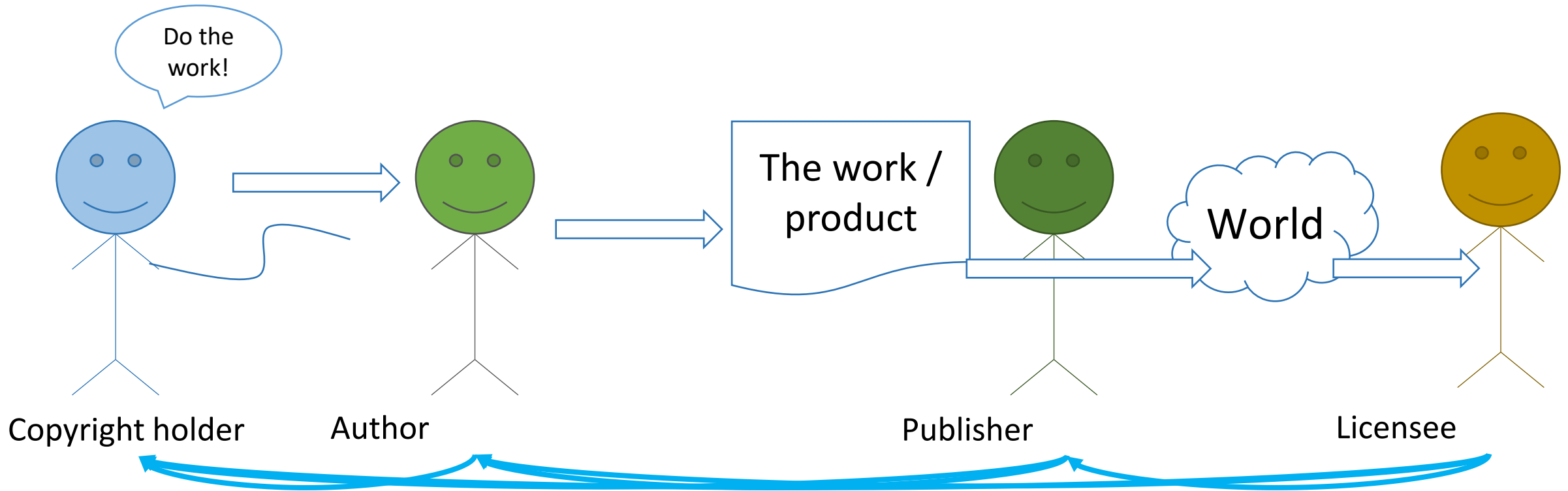
CDEFGHI

DEFGHI

# Data Management

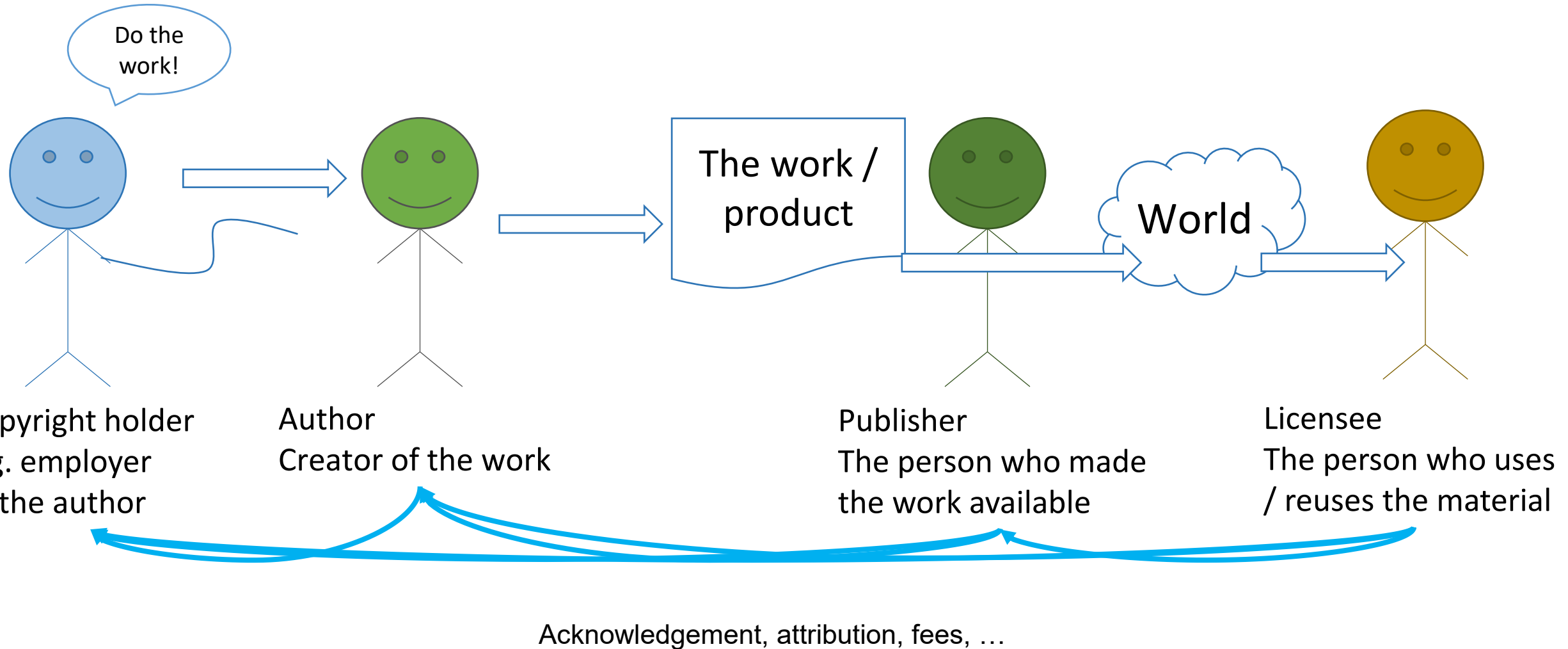
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# Terminology





# Terminology

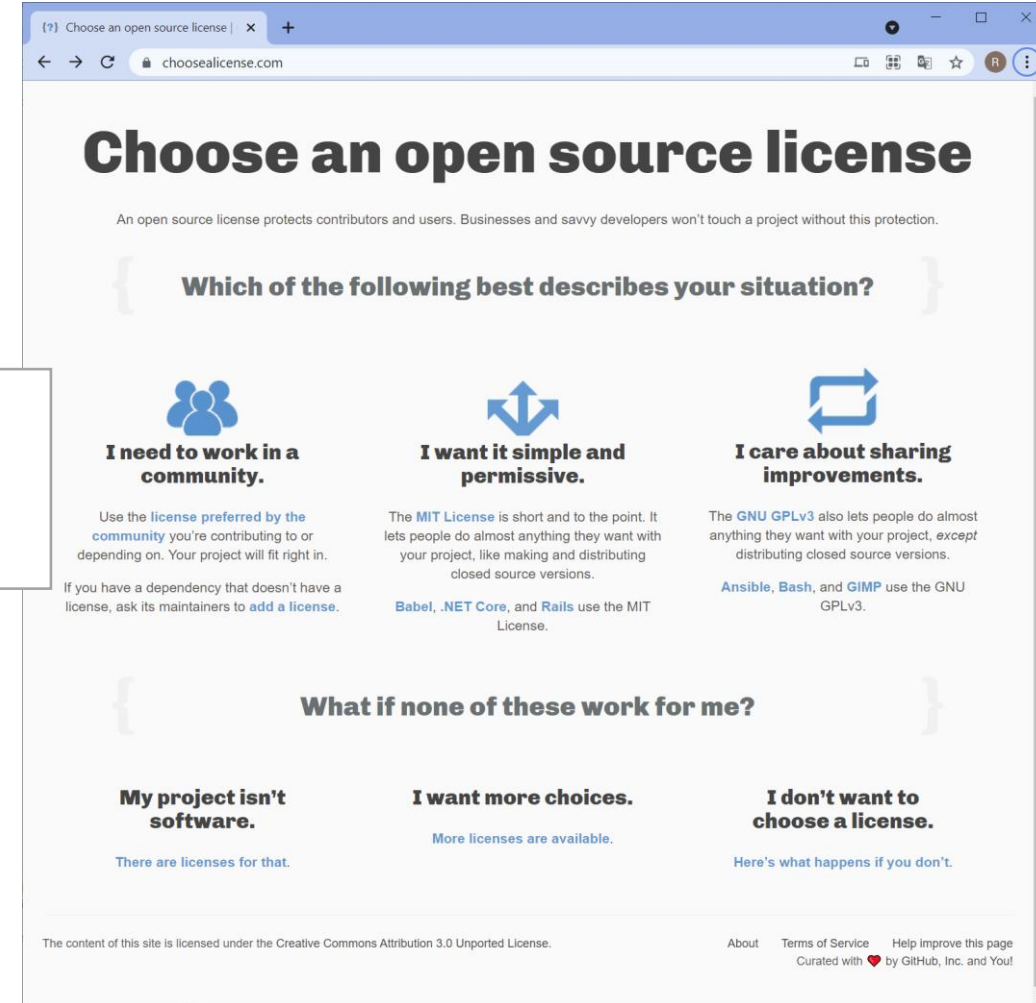
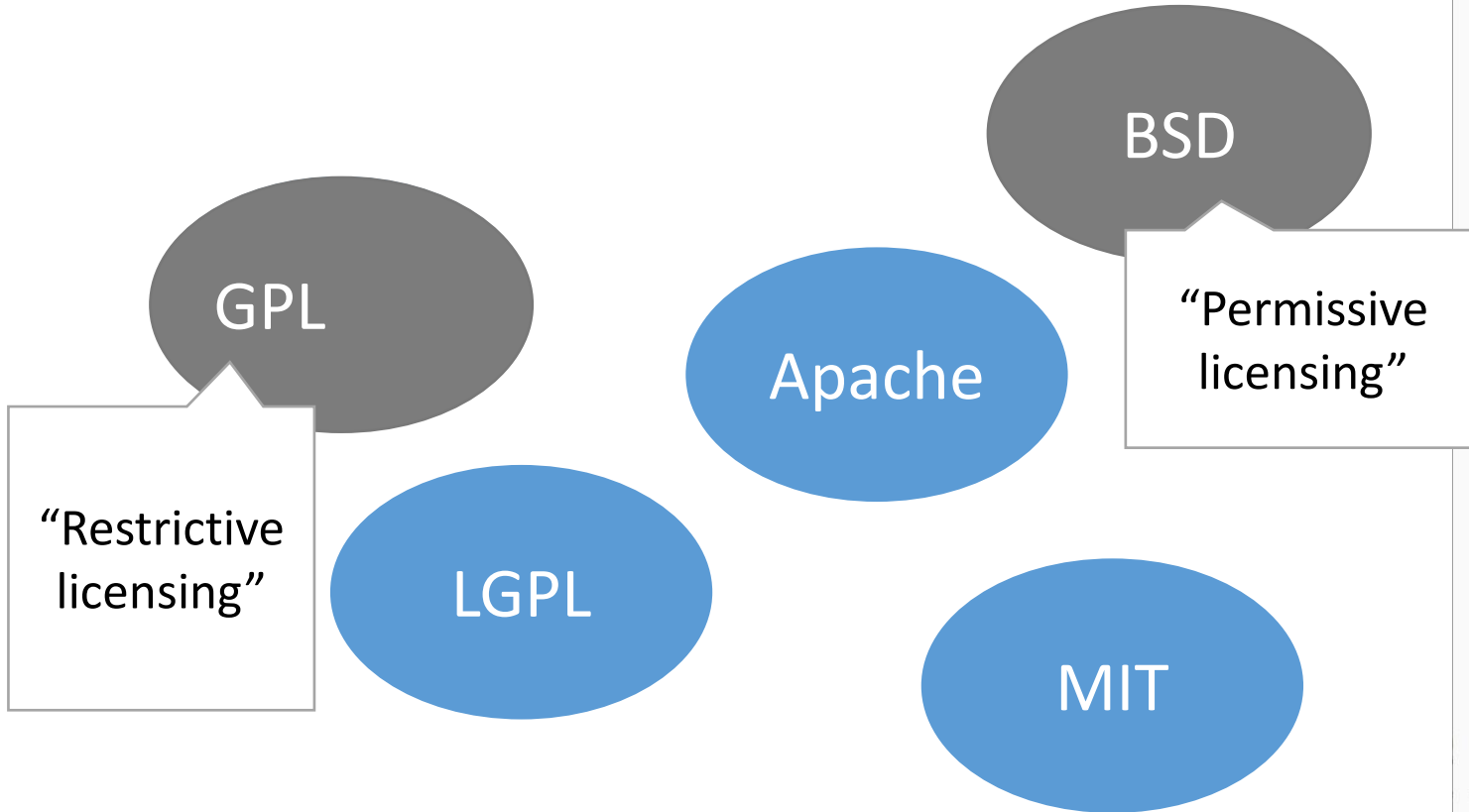


# The FAIR principles

- **Findable**
  - Meta-data, unique identifiers, searchable databases
- **Accessible**
  - Open & free protocols
  - Authentication / authorization
- **Interoperable**
  - [meta] data use a formal, accessible, shared, broadly accessible language
  - References to other [meta]data
- **Reusable**
  - Properly licensed, associated relevant meta data, community-relevant community standards

# Licensing software

- In the software world, other licenses are more popular, historically grown.



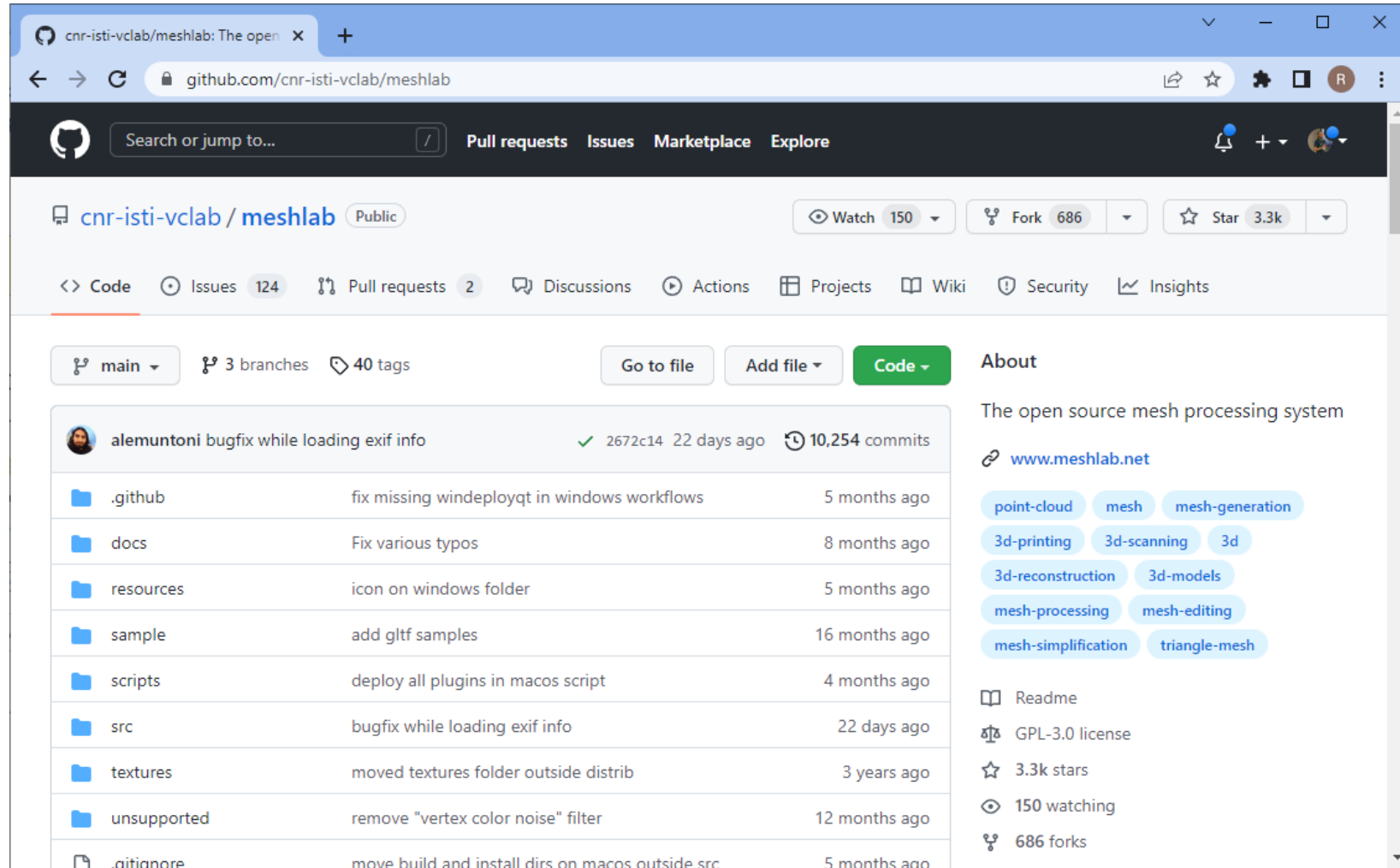
# Quiz

- May I reuse code from this repository in my own BSD-licensed work?

Yes



No



The screenshot shows the GitHub repository page for `cnr-isti-vclab/meshlab`. The repository is public and has 150 watchers, 686 forks, and 3.3k stars. It contains 124 issues, 2 pull requests, and 40 tags. The main branch is selected. The repository description is "The open source mesh processing system" with the website `www.meshlab.net`. The license is GPL-3.0. The repository includes a list of folders and their associated commit messages and dates:

Folder	Commit Message	Time Ago
<code>.github</code>	fix missing windeployqt in windows workflows	5 months ago
<code>docs</code>	Fix various typos	8 months ago
<code>resources</code>	icon on windows folder	5 months ago
<code>sample</code>	add gltf samples	16 months ago
<code>scripts</code>	deploy all plugins in macos script	4 months ago
<code>src</code>	bugfix while loading exif info	22 days ago
<code>textures</code>	moved textures folder outside distrib	3 years ago
<code>unsupported</code>	remove "vertex color noise" filter	12 months ago
<code>.gitignore</code>	move build and install dirs on macos outside src	5 months ago

# Quiz: Digital Object Identifiers

- Which of these is a *unique* digital object identifier?

<https://twitter.com/haesleinhuepf/status/891596662782779392>

<https://doi.org/10.5281/zenodo.28325>

<https://github.com/haesleinhuepf/devbio-napari>

<https://napari.org/>

# Documenting dependencies

- Maintain a document with the dependencies (and versions) you need in your project!
  - The conda way

```
environment.yml - Notepad
File Edit Format View Help
name: cellpose-test
channels:
  - conda-forge
  - pytorch
dependencies:
  - python=3.8
  - pytorch=1.12
  - devbio-napari
  - cellpose-napari
Ln 3, C 100% Windows (CRLF) UTF-8
```

In case your environment is screwed up, you can rebuild it any time.

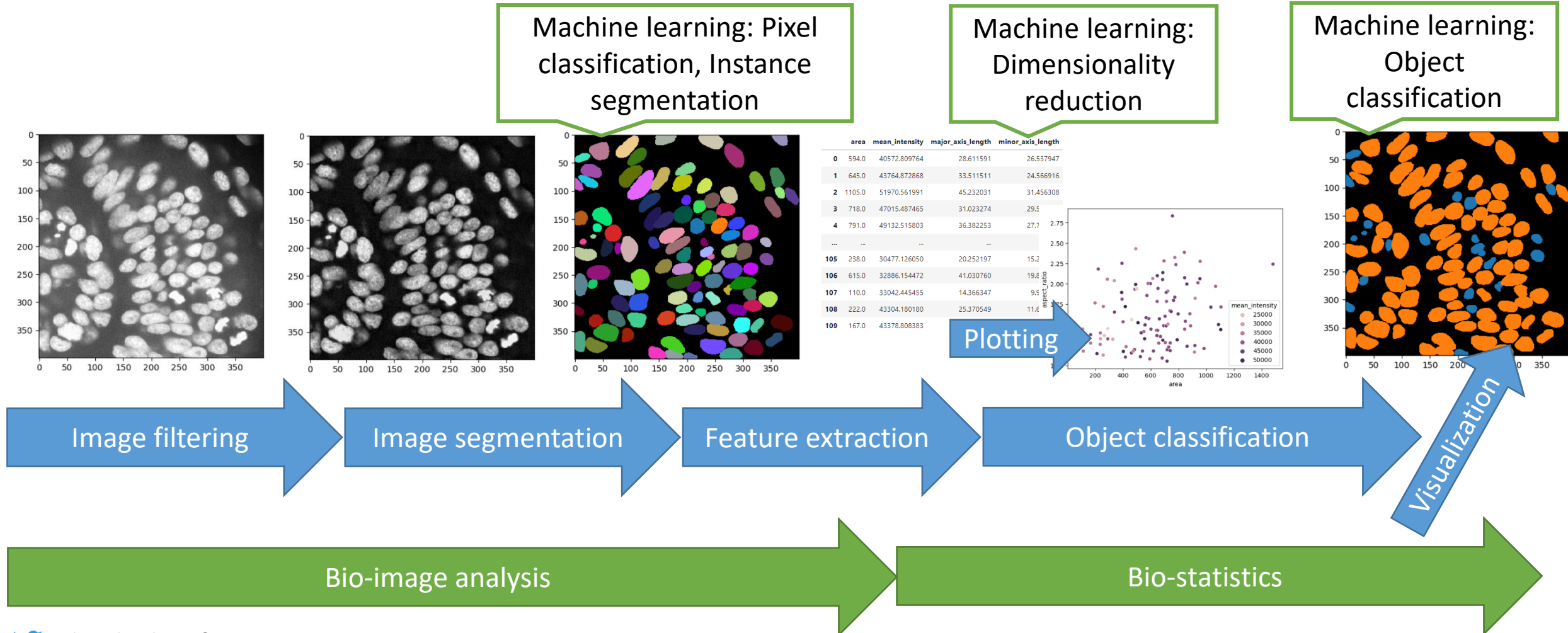
```
conda env create -f environment.yml
```

# Bio-image Analysis

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# Lecture overview: Bio-image Analysis

- Image Data Analysis workflows
- Goal: **Quantify observations, substantiate conclusions with numbers**

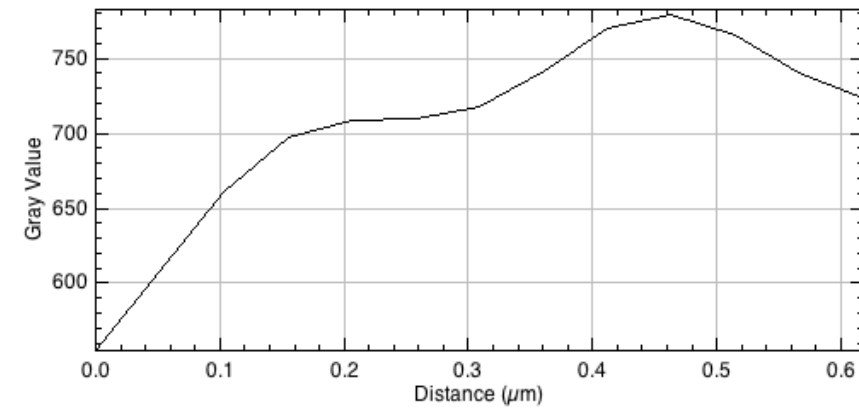
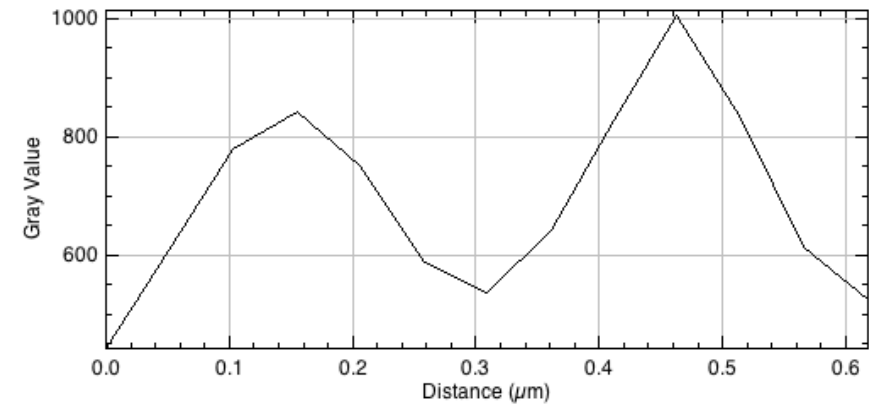
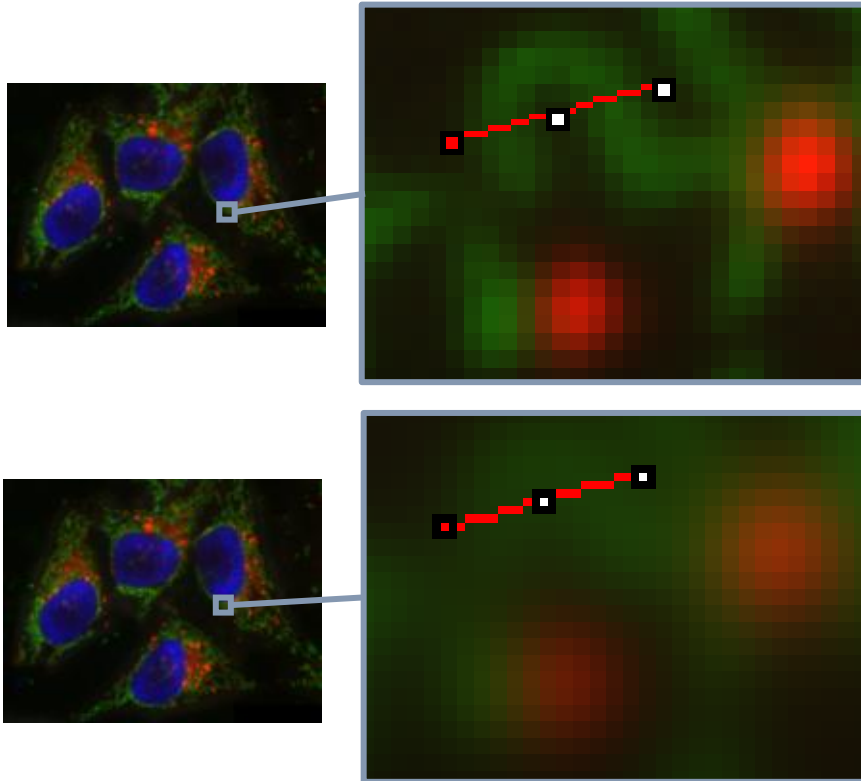




- Bio-image analysis is supposed to be
  - **Quantitative**
    - We derive numbers from images which describe physical properties of the observed sample.
  - **Objective**
    - The derived measurement does not depend on who did the measurement. The measurement is free of interpretation.
  - **Reliable (trustworthy / validated)**
    - We are confident that the measurement is describing what it is supposed to describe.
  - **Reproducible**
    - Enabling others to re-do the experiment. For this, documentation is crucial!
  - **Replicability**
    - Others *do* execute the same analysis, potentially on other data, and see consistent results.
  - **Repeatable**
    - We can do the same experiment twice under the *same conditions* and get the same measurements.

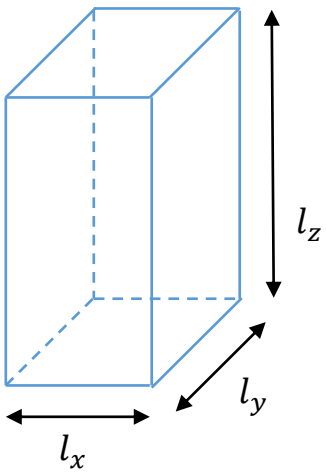
# Pixel size versus resolution

- How is the resolution of an imaging system defined?
- How is the pixel size of an image defined?

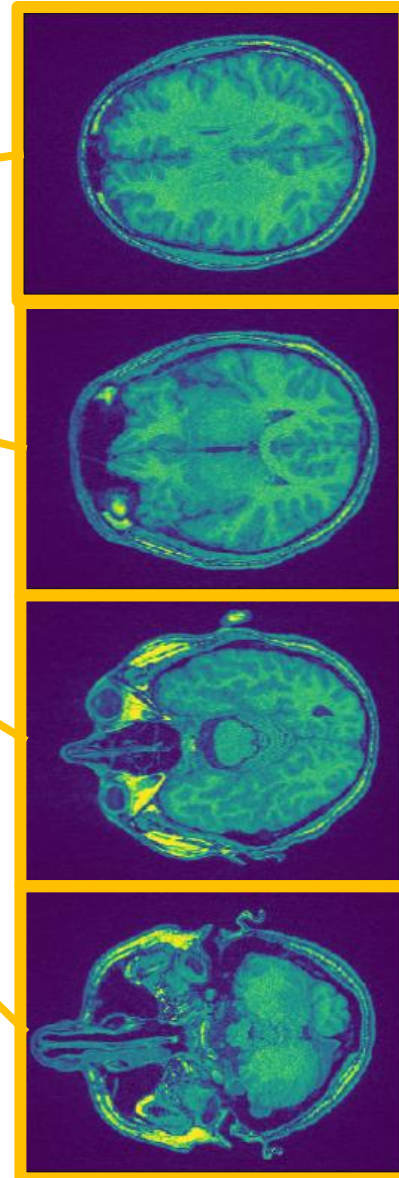
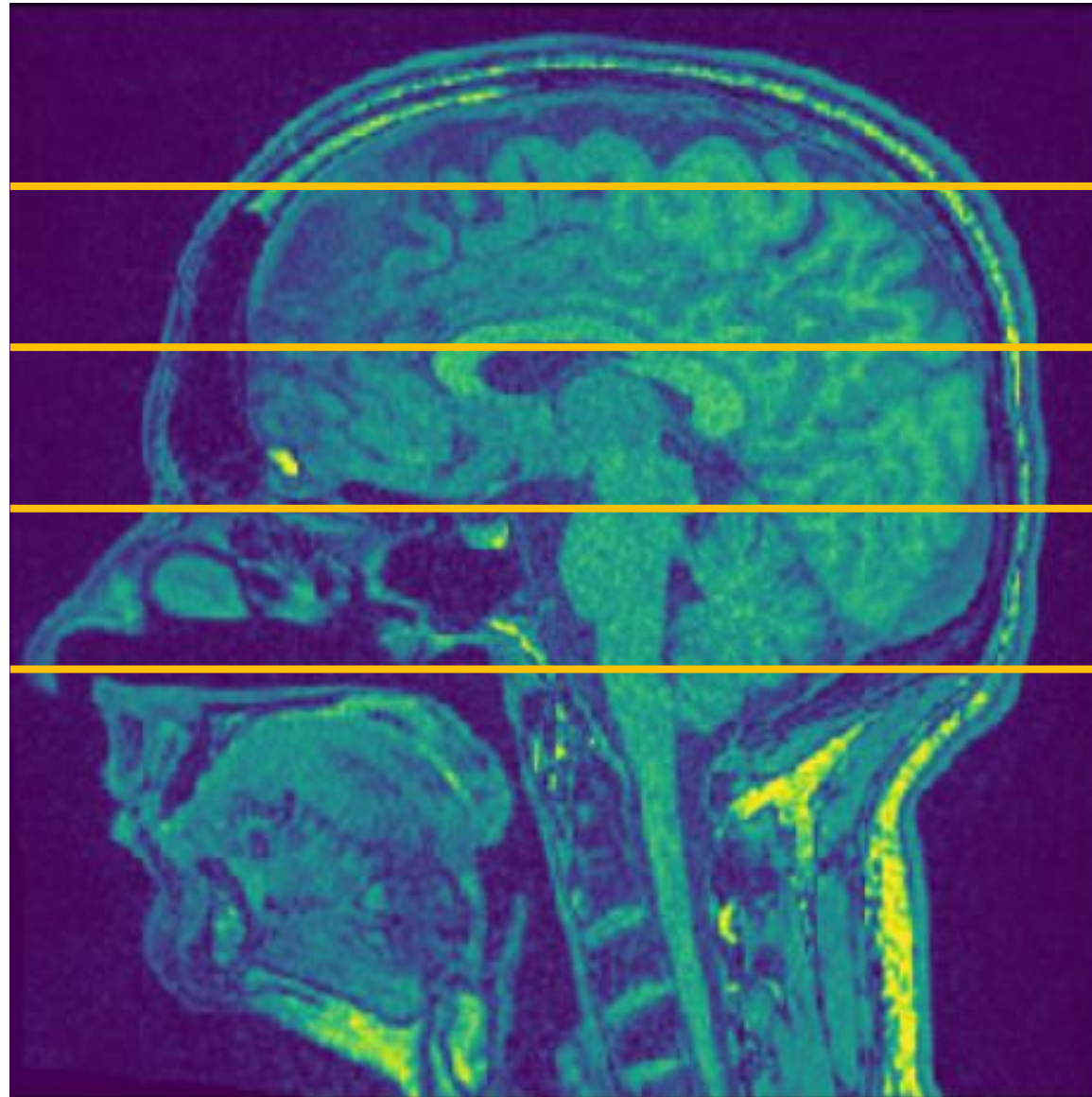


# Image stacks and voxels

- 3-dimensional images consisting of voxels
- “Image stack”
- Often anisotropic



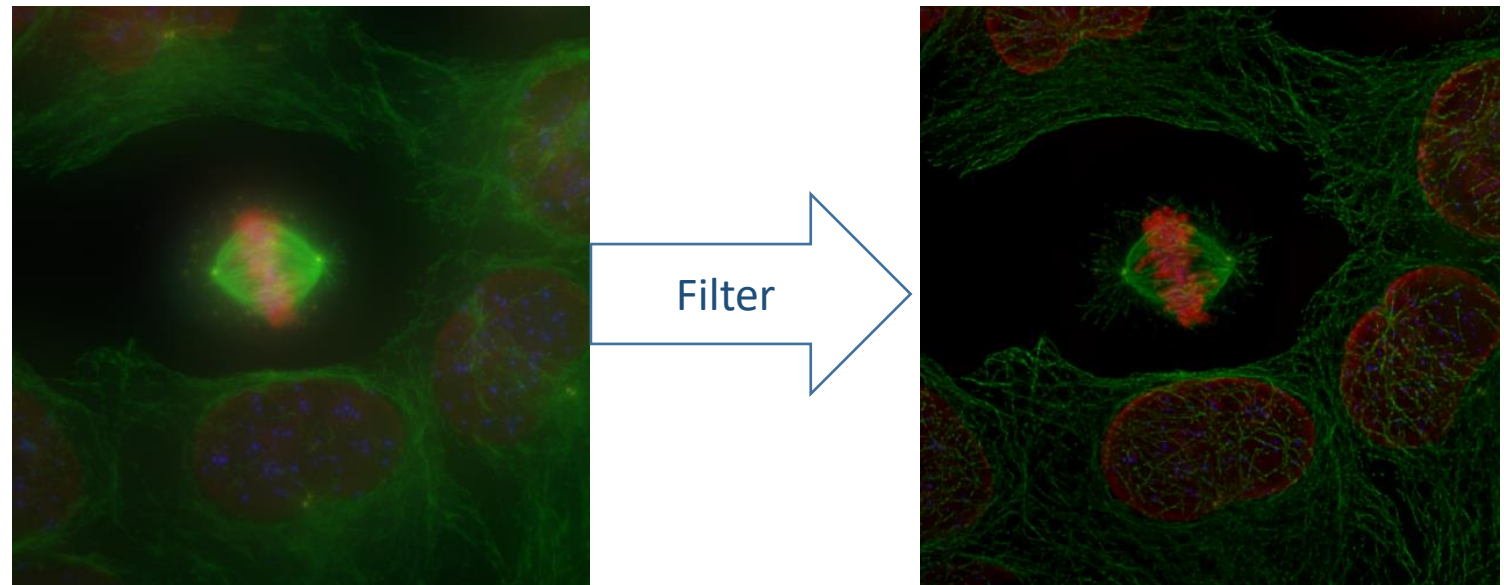
$$l_x = l_y \neq l_z$$



# Image Processing Filters

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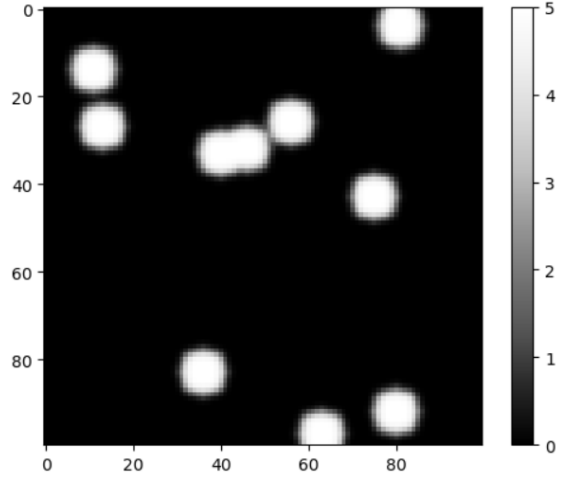
- An image processing filter is an operation on an image.
- It takes an image and produces a new image out of it.
- Filters change pixel values.
  
- There is no “best” filter. Which filter fits your needs, depends on the context.
- Filters do not do magic. They can not make things visible which are not in the image.
  
- Application examples
  - Noise-reduction
  - Background removal
  - Artefact-removal
  - Contrast enhancement
  - Correct uneven illumination



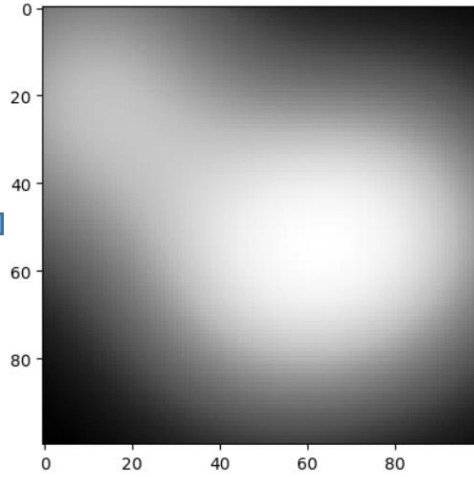
# Effects harming image quality

- Image formation (simulated)

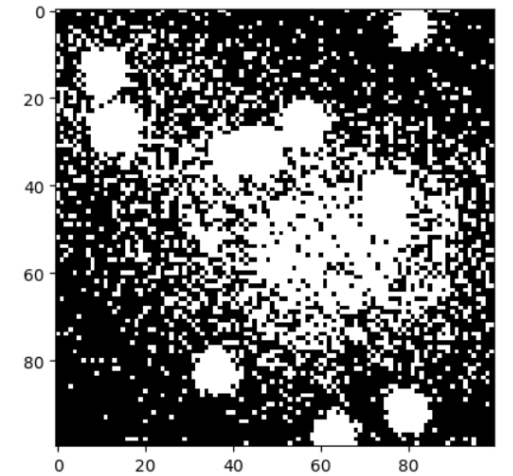
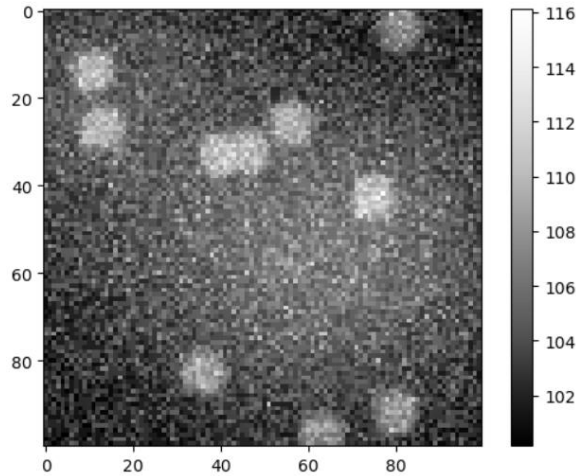
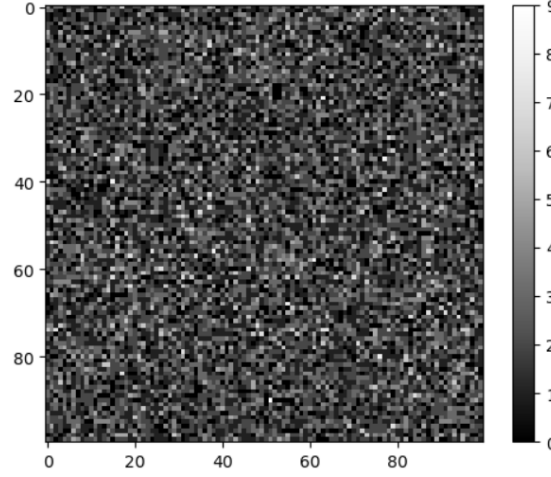
“nuclei”



“background”

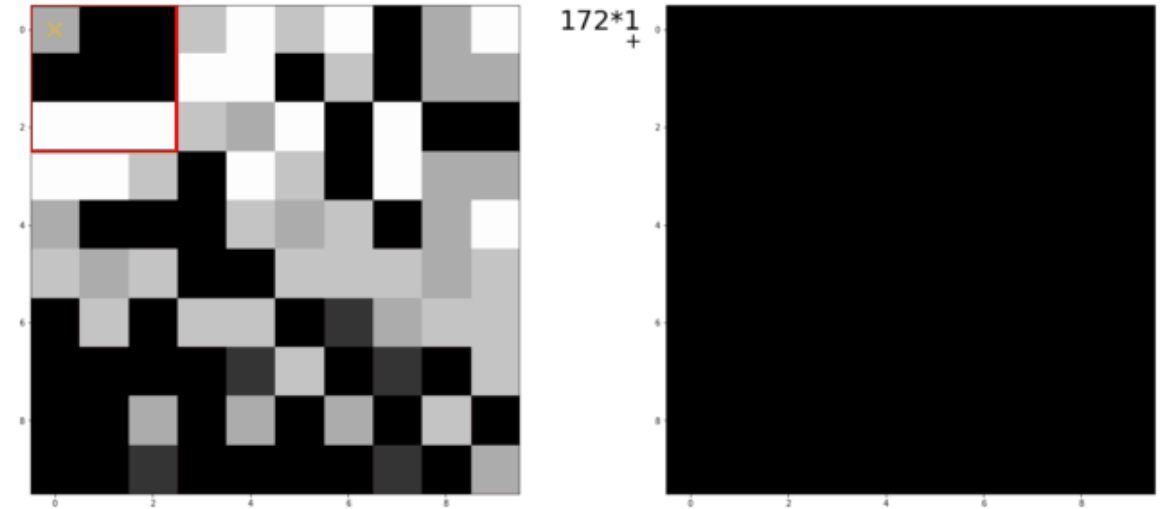


“noise”



# Linear Filters: Convolution

- What is an image processing filter?
- What is a filter kernel?
- How does convolution work?

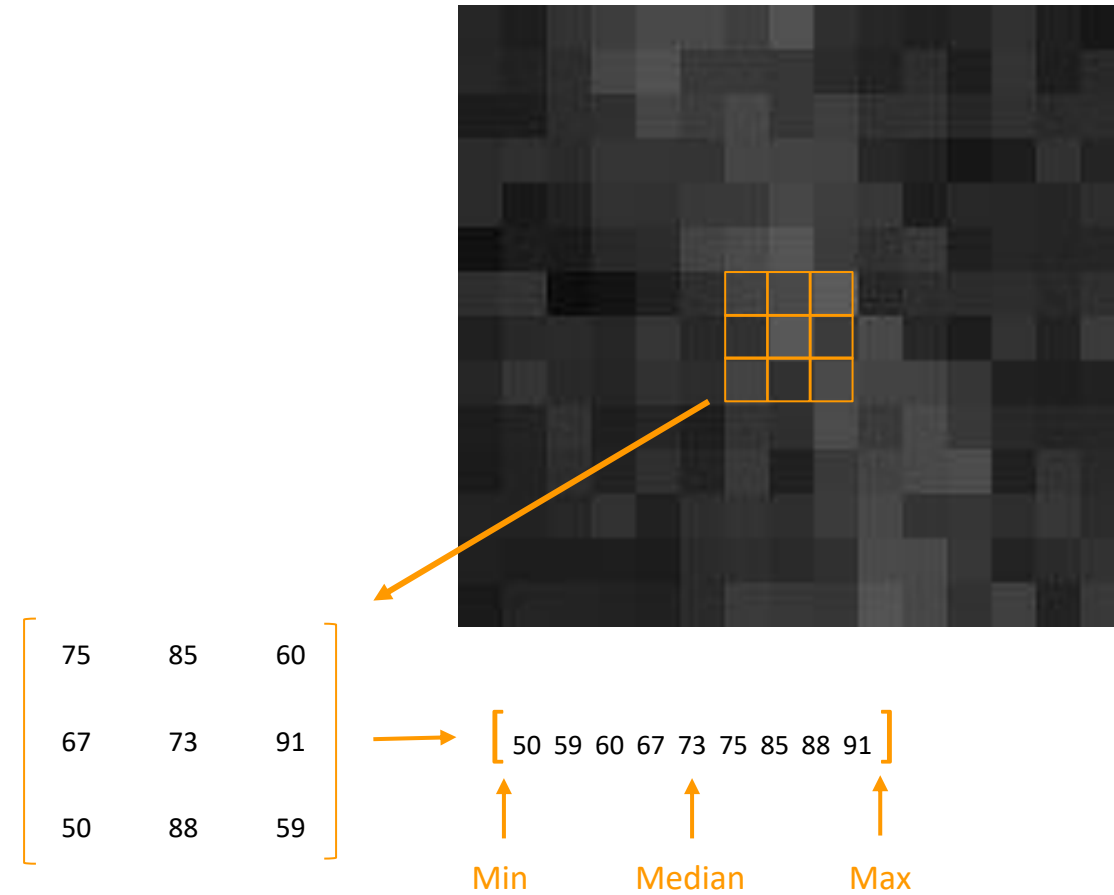


Mean filter

$$\begin{bmatrix} 1/9 & 1/9 & 1/9 \\ 1/9 & 1/9 & 1/9 \\ 1/9 & 1/9 & 1/9 \end{bmatrix}$$

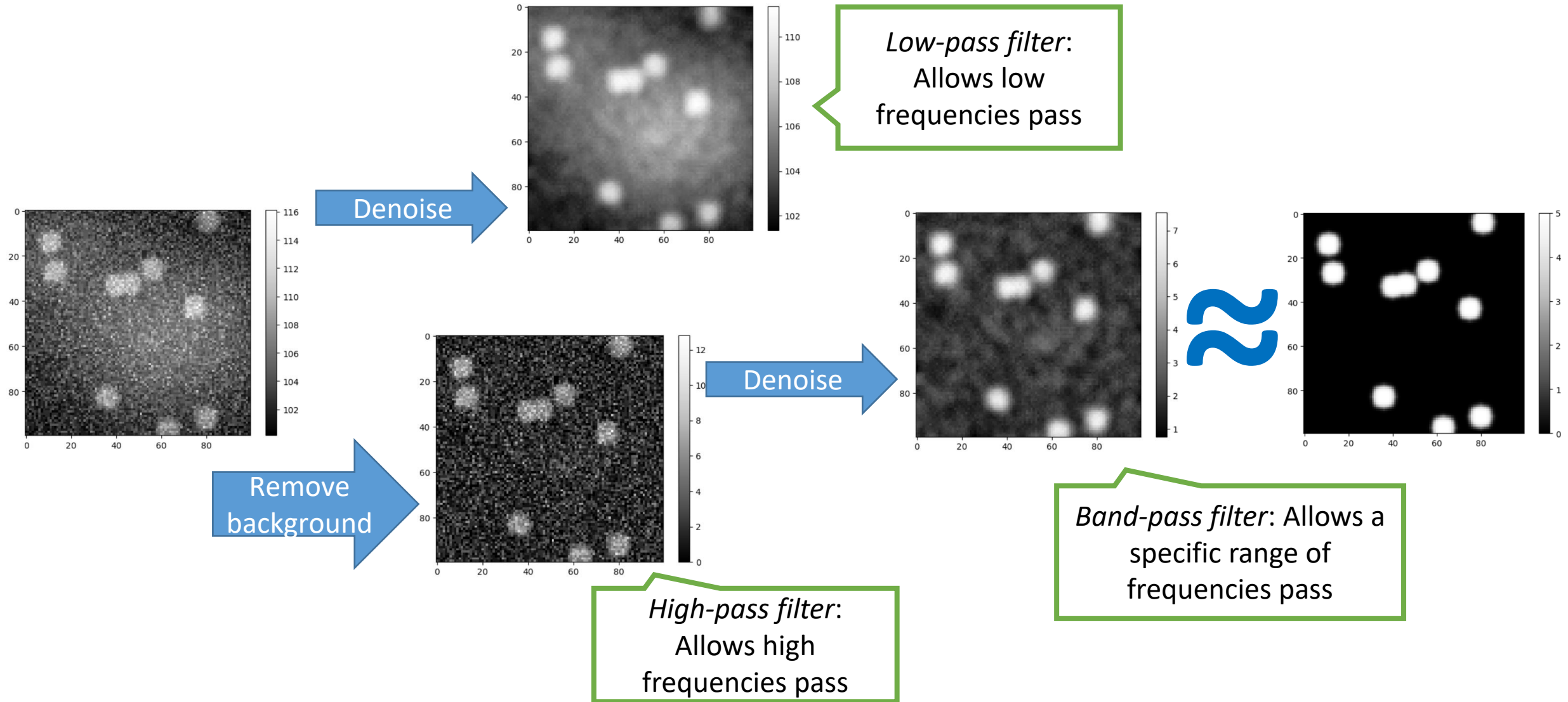
Animation source: Dominic Waithe, Oxford University  
[https://github.com/dwaithe/generalMacros/tree/master/convolution\\_animation](https://github.com/dwaithe/generalMacros/tree/master/convolution_animation)

- What differentiates linear and non-linear filters?





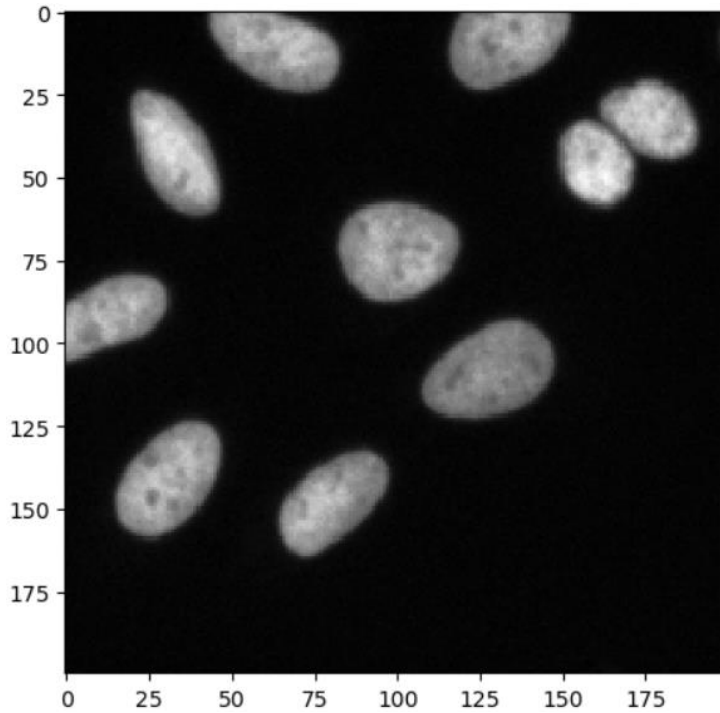
- High-pass versus low-pass filters



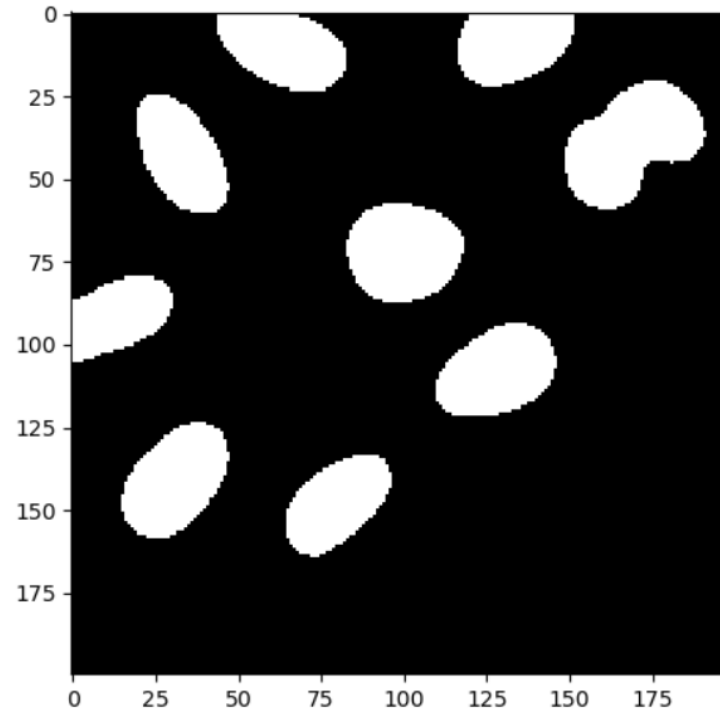
# Image Segmentation

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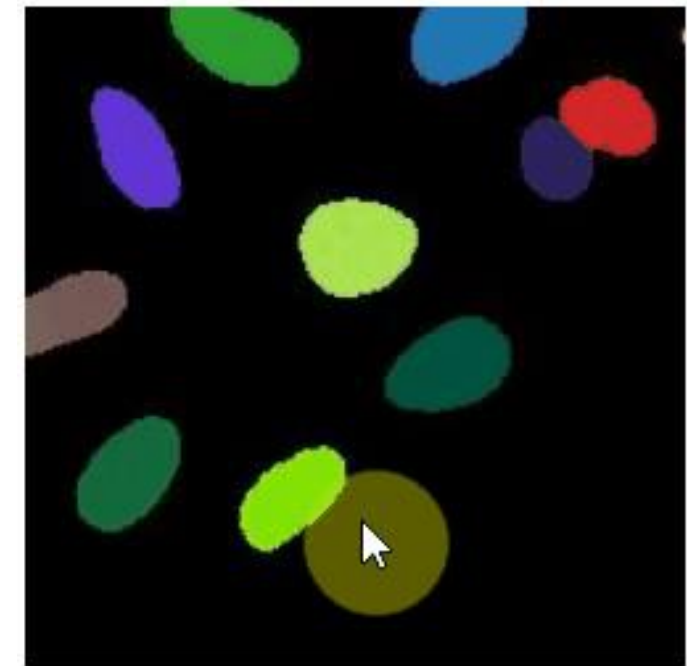
## Intensity image



## Binary image

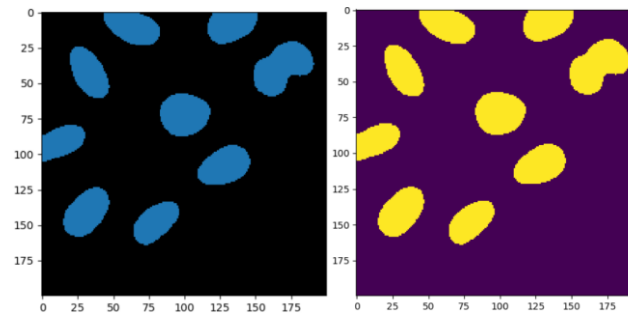


## Label image



[y=152, x=92] = 0

No matter how they are displayed



- Annotations are typically drawn by humans (e.g. to train machine learning models)

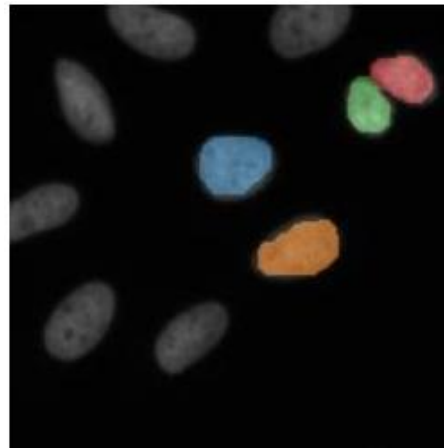
Instance  
segmentation



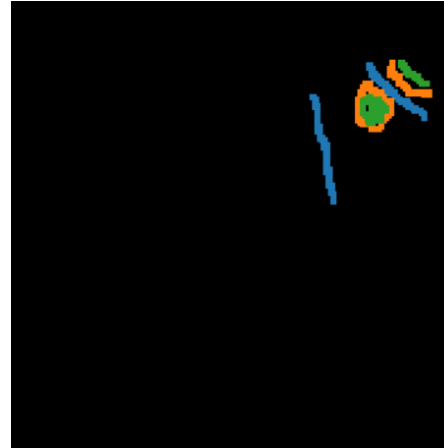
Semantic  
segmentation

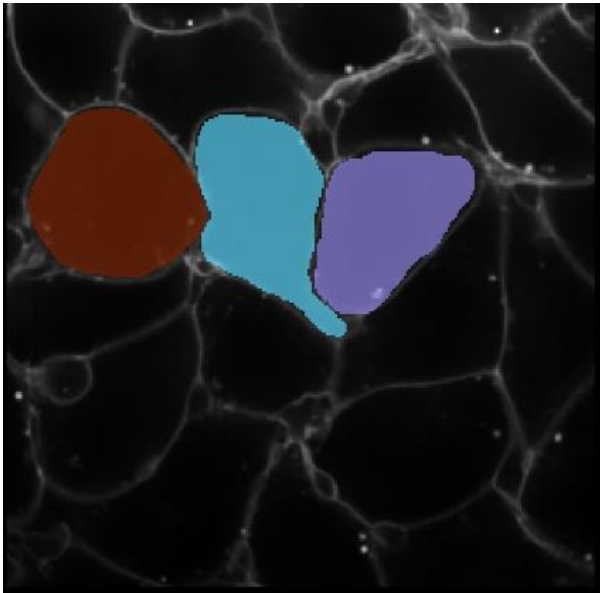


Sparse instance  
annotation



Sparse semantic  
annotation





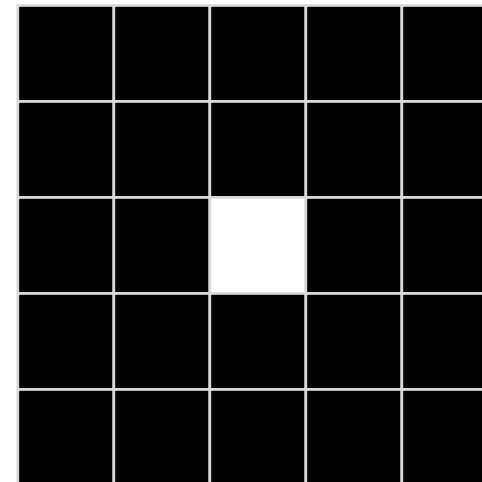
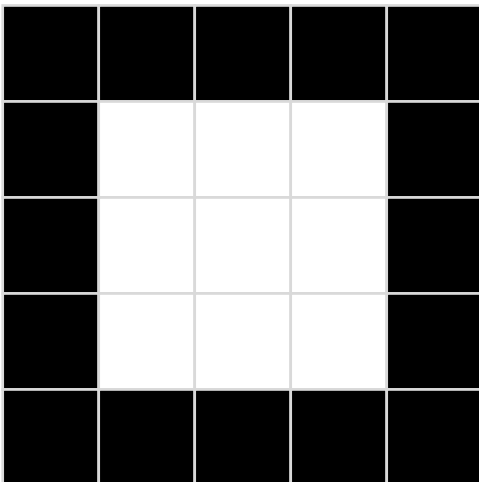
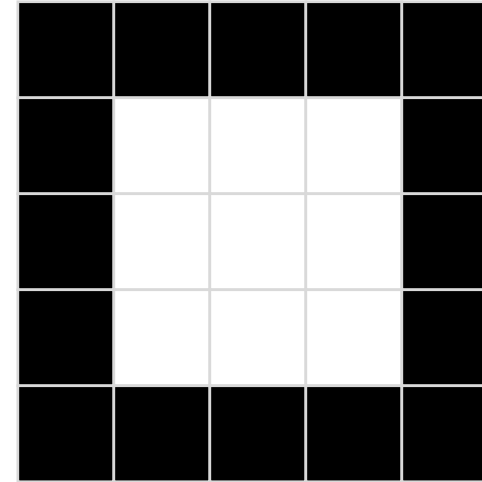
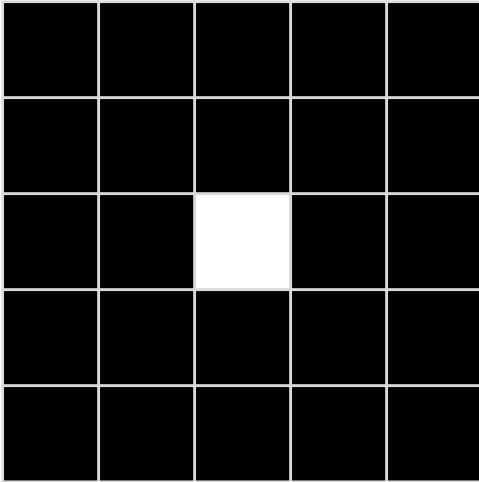
This is a ...

Sparse  
instance  
segmentation

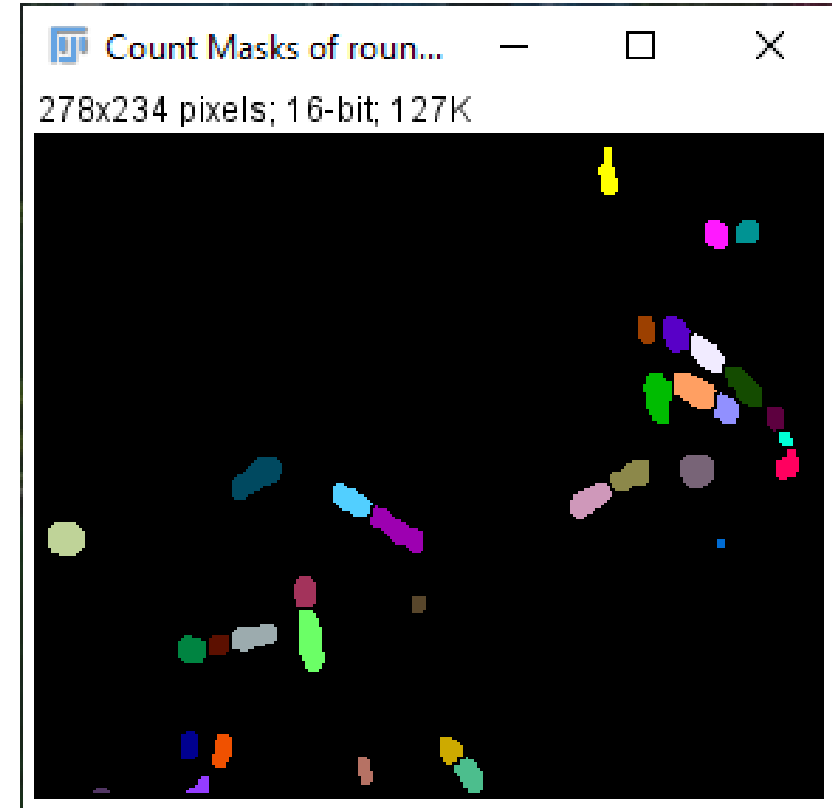
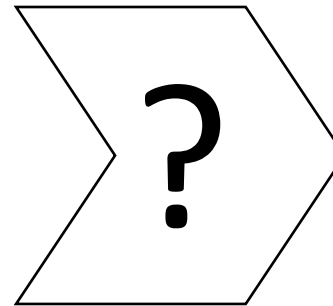
Sparse  
semantic  
segmentation

# Refining masks: Dilation and Erosion

- Dilation: Every pixel with at least one white neighbor becomes white.



- What's the name of the operation applied here?



Thresholding

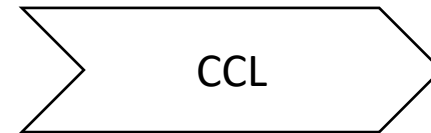
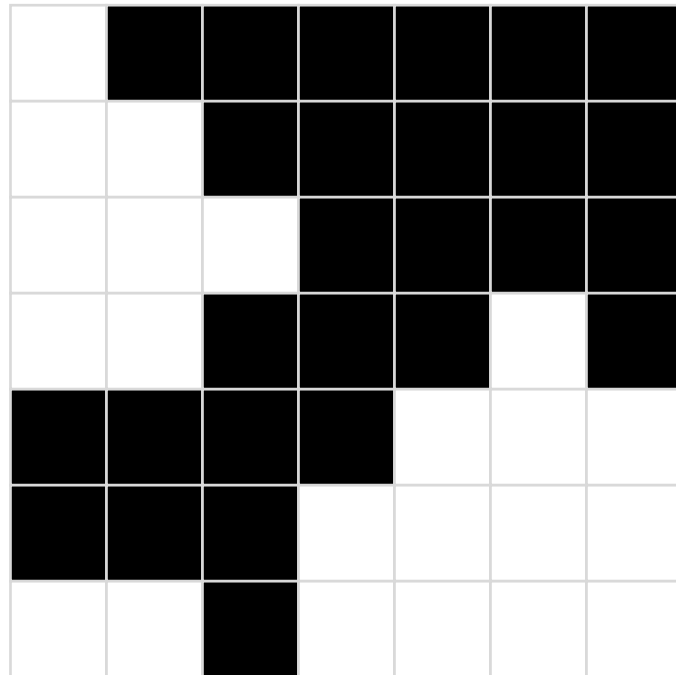
Binary closing

Connected components analysis

Watershed

# Connected components labelling

- In order to allow the computer differentiating objects, connected components analysis (CCA) is used to mark pixels belonging to different objects with different numbers
- Background pixels are marked with 0.
- The maximum intensity of a labelled map corresponds to the number of objects.

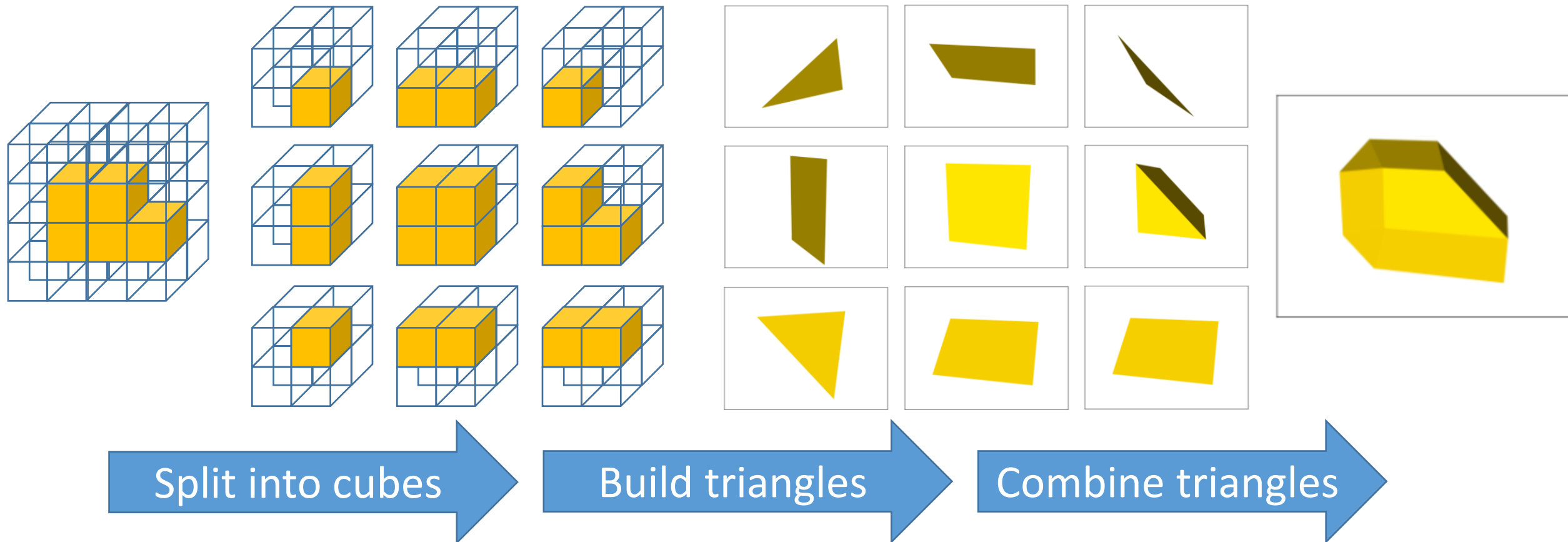


1	0	0	0	0	0	0
1	1	0	0	0	0	0
1	1	1	0	0	0	0
1	1	0	0	0	3	0
0	0	0	0	3	3	3
0	0	0	3	3	3	3
2	2	0	3	3	3	3

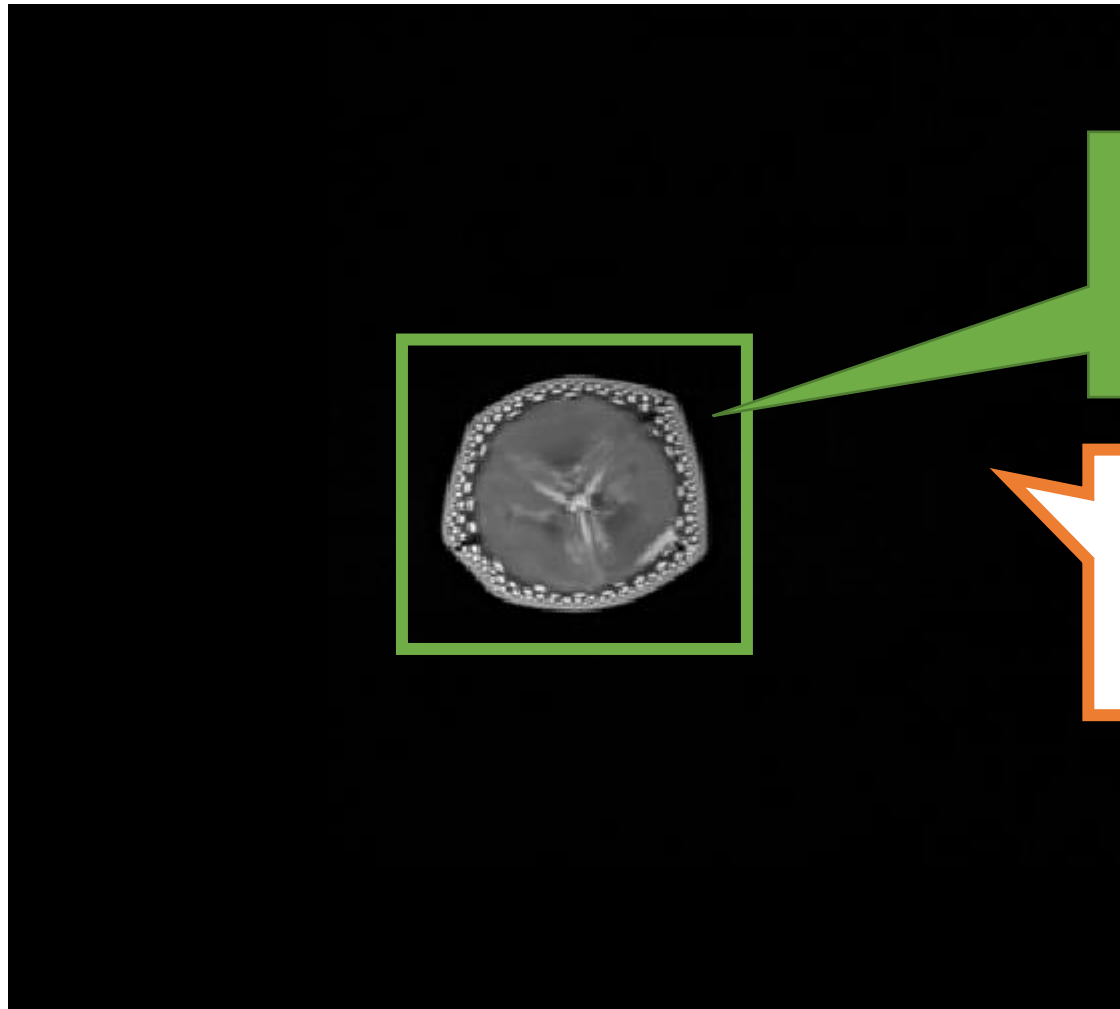


# Marching cubes algorithm

- Starting point: 3D binary image
- Cuts the image in small cubes and iterates over them



- Crop out the region you're interested in



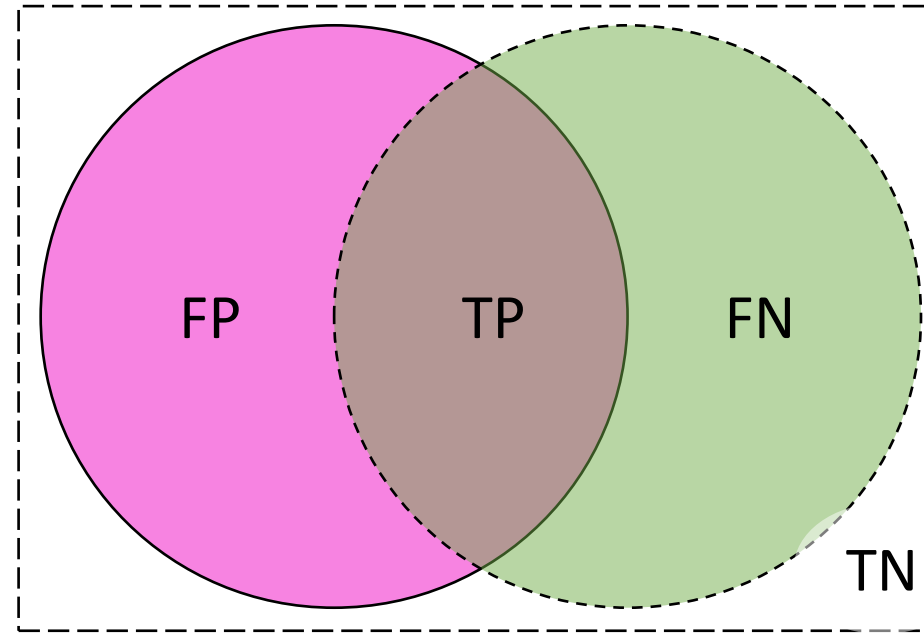
Interesting

Not  
interesting

In this case  
you can spare  
8/9 compute time for  
following processing steps

# Segmentation quality estimation

- In general
  - Define what's positive and what's negative.
  - Compare with a reference to figure out what was true and false
- Welcome to the Theory of Sets



- A Prediction A
- B Reference B (ground truth)
- ROI Region of interest
- TP True-positive
- FN False-negative
- FP False-positive
- TN True-negative

Overlap  
(a.k.a. Jaccard index)  $\frac{TP}{TP + FN + FP}$

How much do A and B overlap?

Precision  $\frac{TP}{TP + FP}$

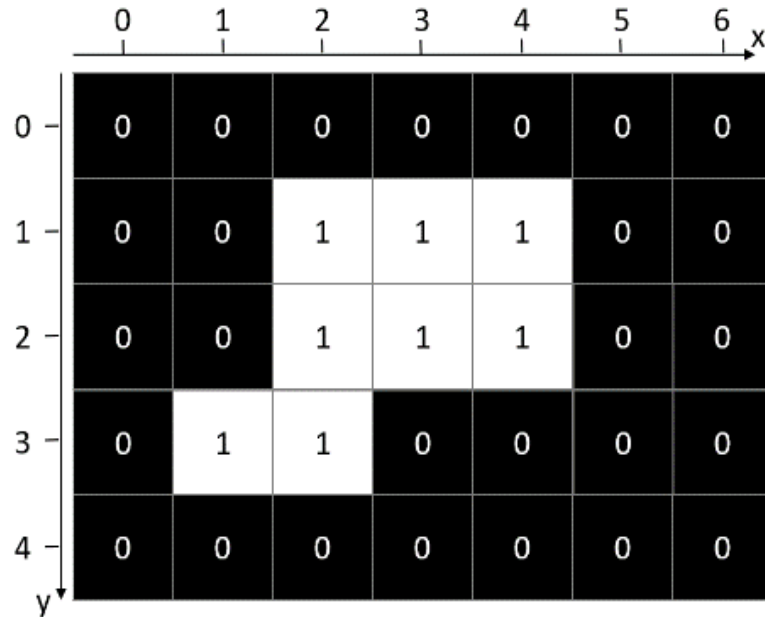
What fraction of points that were predicted as positives were really positive?

Recall  
(a.k.a. sensitivity)  $\frac{TP}{TP + FN}$

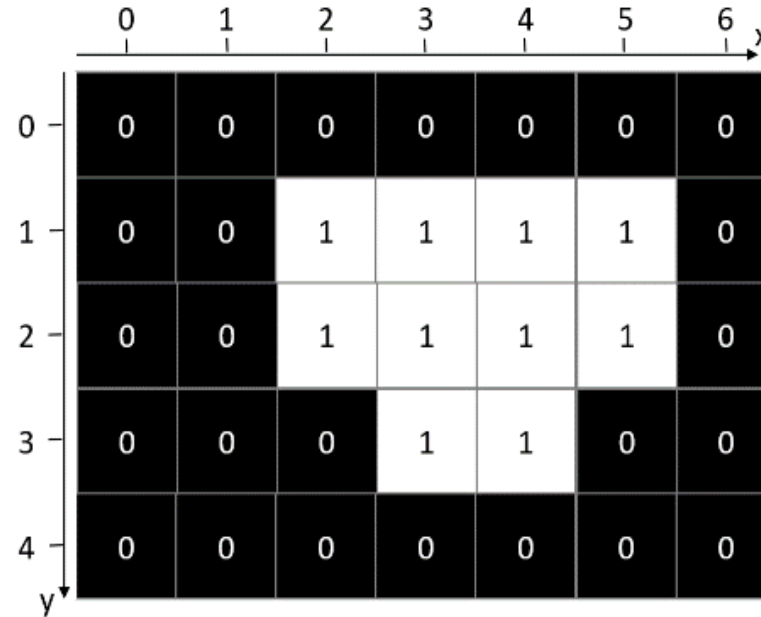
What fraction of positives points were predicted as positives?

- Assume you are evaluating a binary segmentation algorithm by comparing its result to a given ground

Segmentation result



Ground truth



TP = 6  
FP = 2  
FN = 4  
J = 6 / 12  
P = 6 / 8  
R = 6 / 10

Jaccard index  $\frac{TP}{TP + FN + FP}$

Precision  $\frac{TP}{TP + FP}$

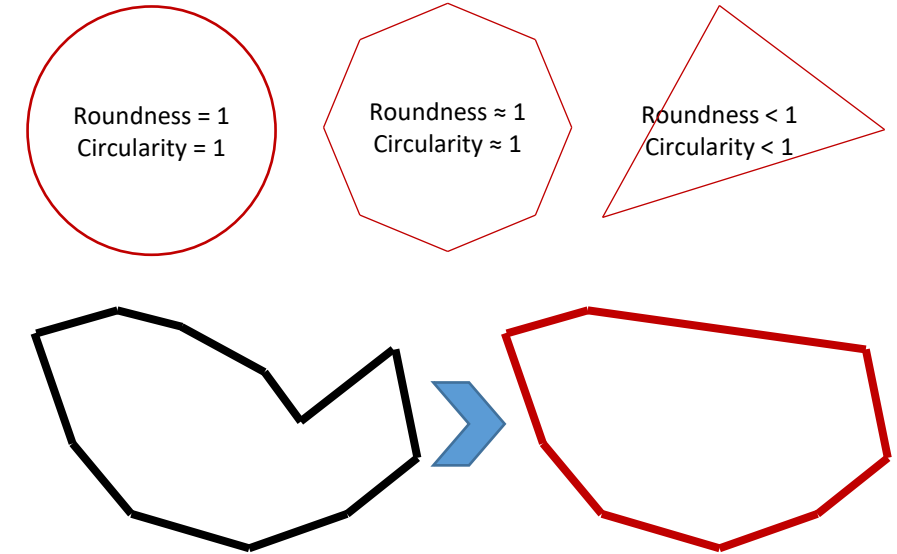
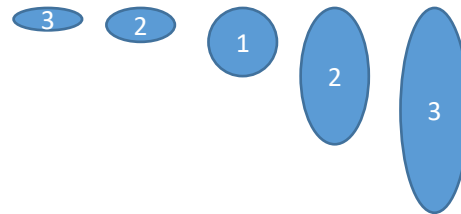
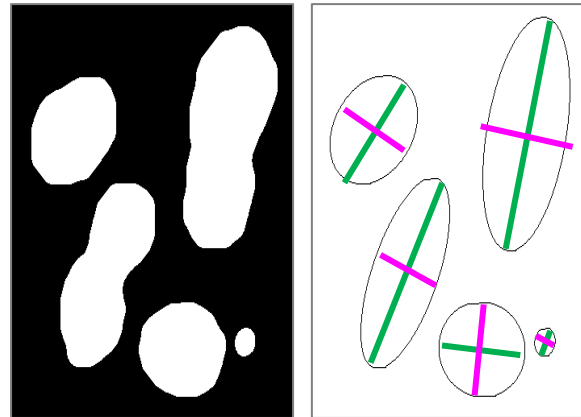
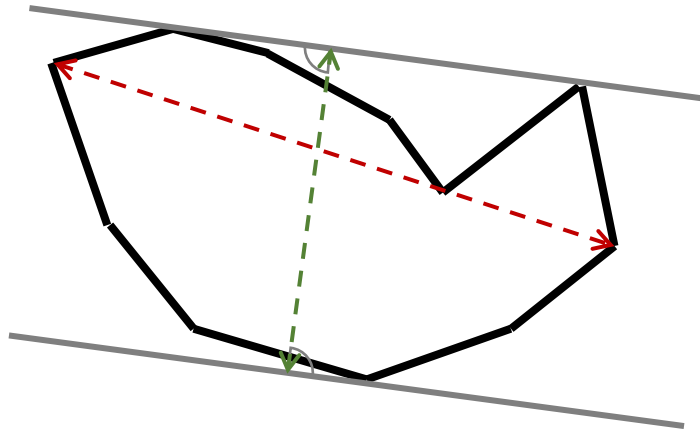
Recall  $\frac{TP}{TP + FN}$

# Feature extraction

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- A *feature* is a countable or measurable property of an image or object.
- Goal of feature extraction is finding a minimal set of features to describe an object well enough to differentiate it from other objects.
- Intensity based features
  - Mean intensity
  - Standard deviation
  - Total intensity
  - Textures
  - ...
- Shape based / spatial features
  - Area / Volume
  - Roundness
  - Solidity
  - Circularity / Sphericity
  - Elongation
  - Centroid
  - Bounding box
  - ...
- Spatio-temporal features
  - Displacement,
  - Speed,
  - Acceleration,
  - ...
- Others
  - Overlap
  - Colocalisation
  - Network-analysis
  - ...
- Mixed features
  - Center of mass
  - Local minima / maxima
  - ...

- How are objects shaped?



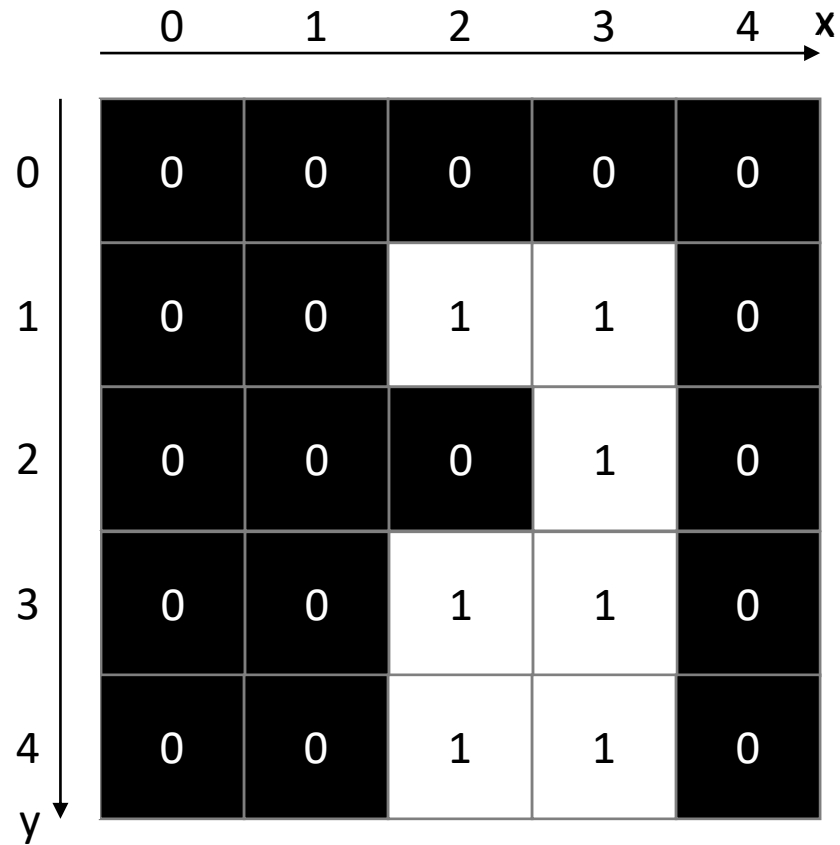
- **Feret's diameter**
- The **minimum caliper** ("Minimum Feret")

- Fit ellipse
  - Major axis ... long diameter
  - Minor axis ... short diameter
- Aspect ratio

- Roundness
- Circularity
- Solidity

# Quiz: solidity

- What is the solidity of the white object in this image?
  - Hint: Area of the convex hull



0.875

1

5

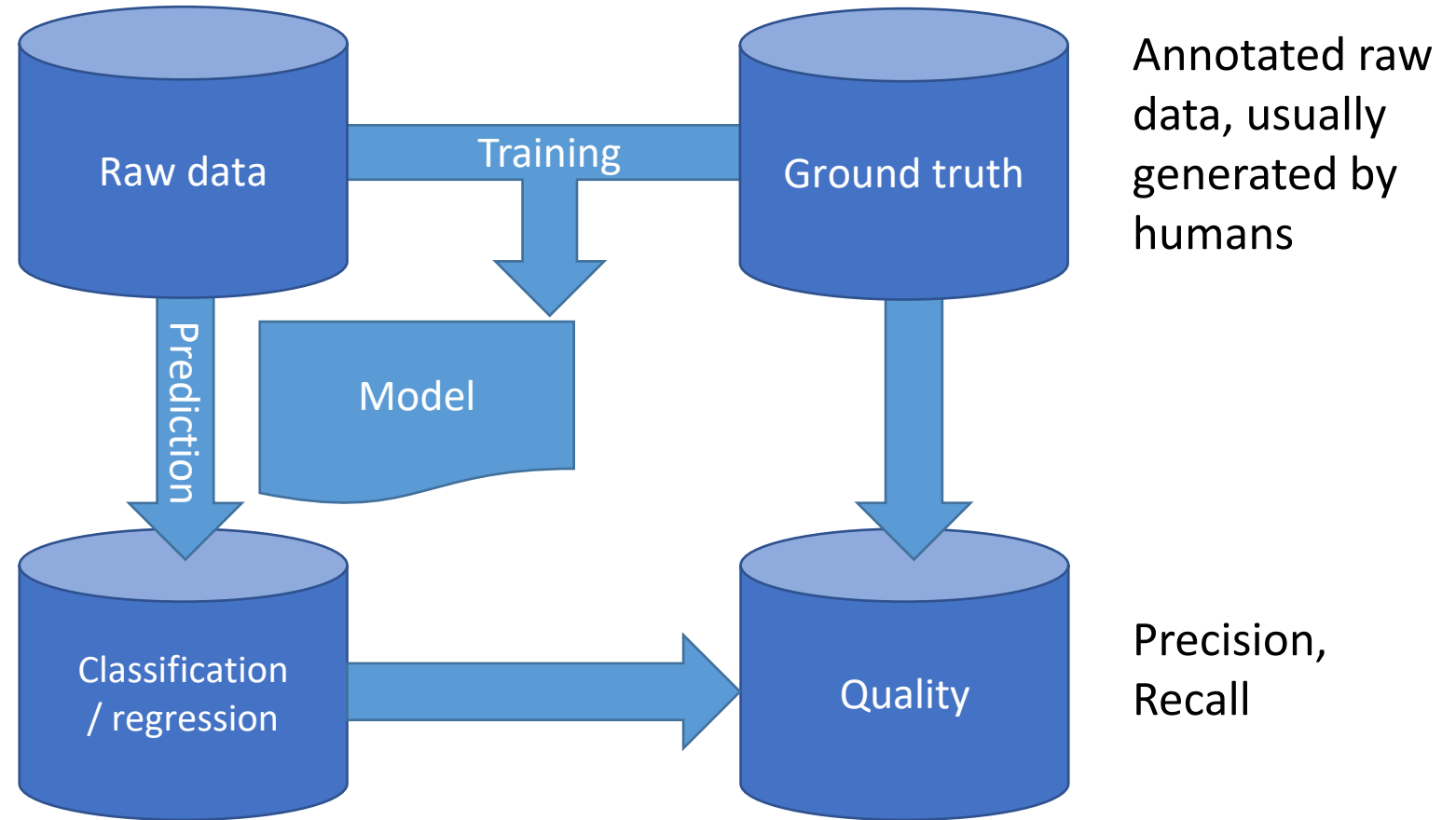
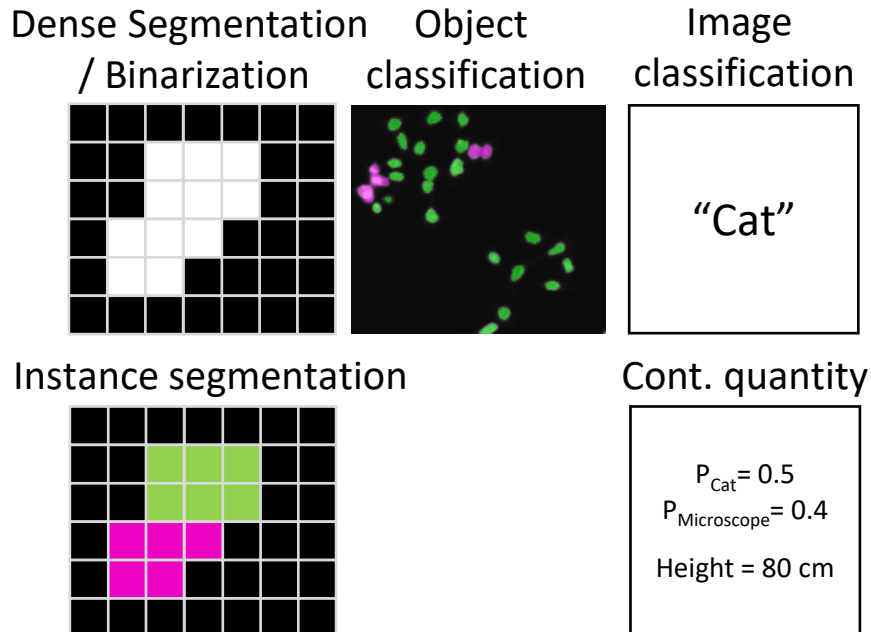
1.2



# Machine learning

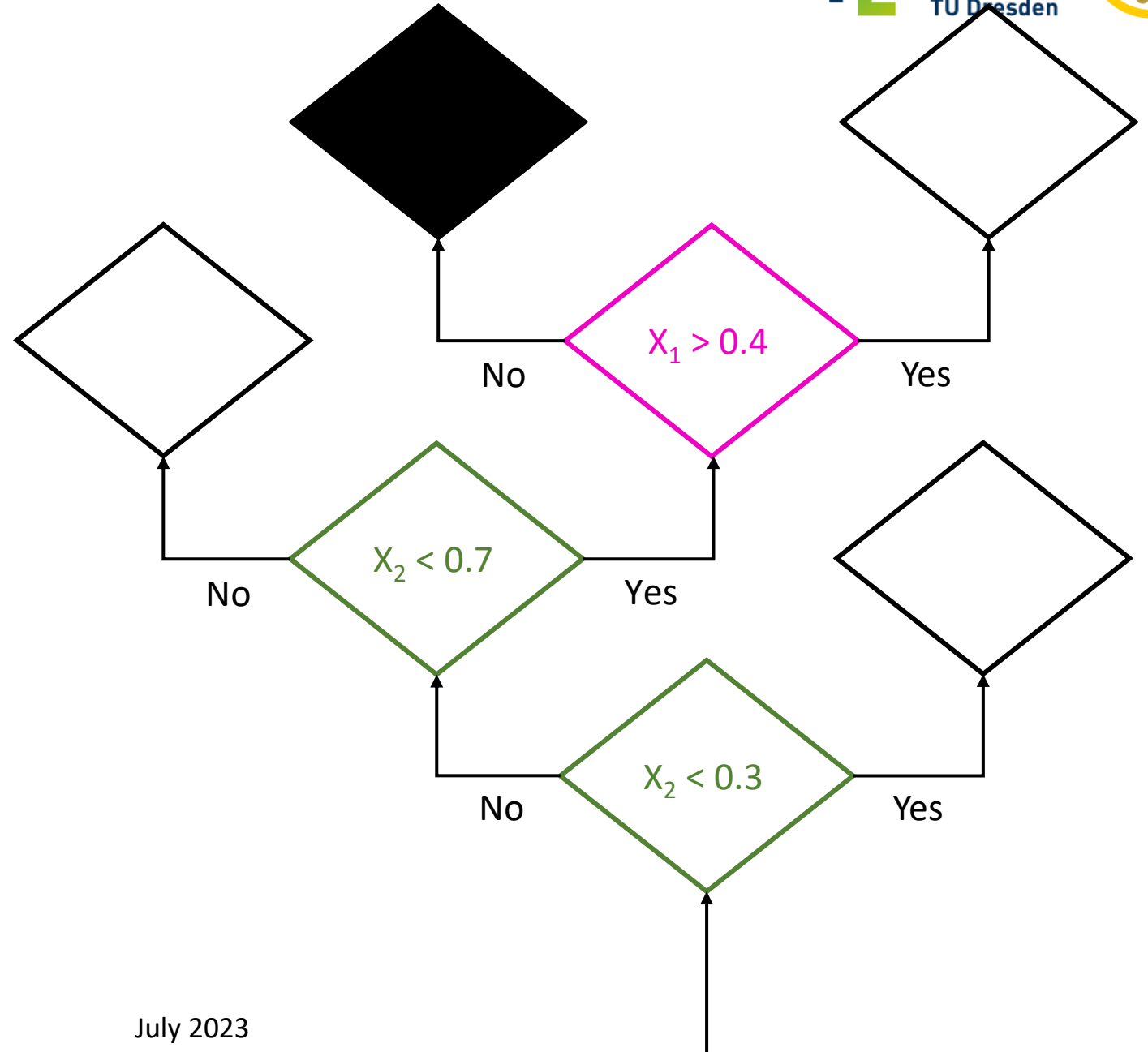
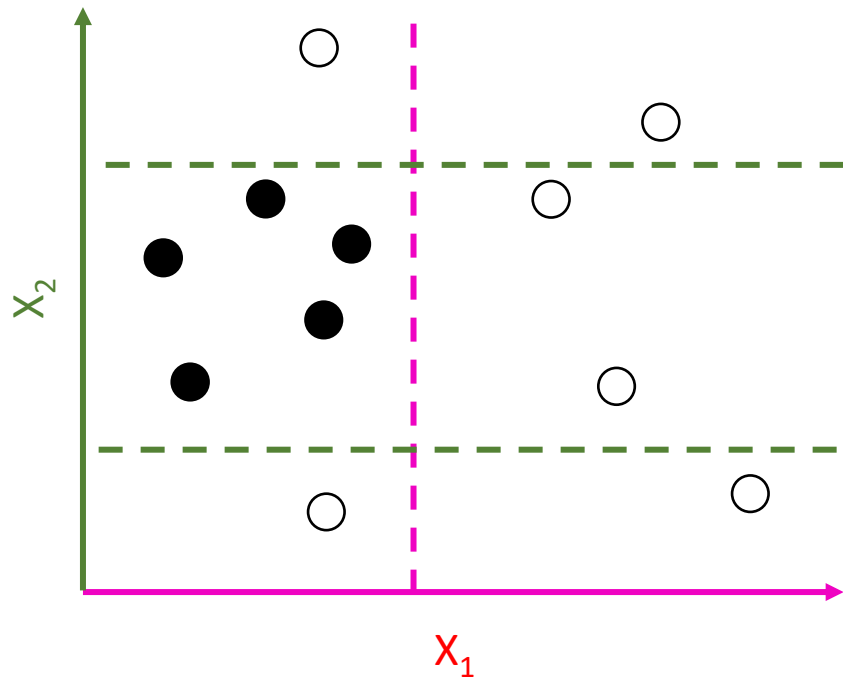
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- Automatic construction of predictive models from given data



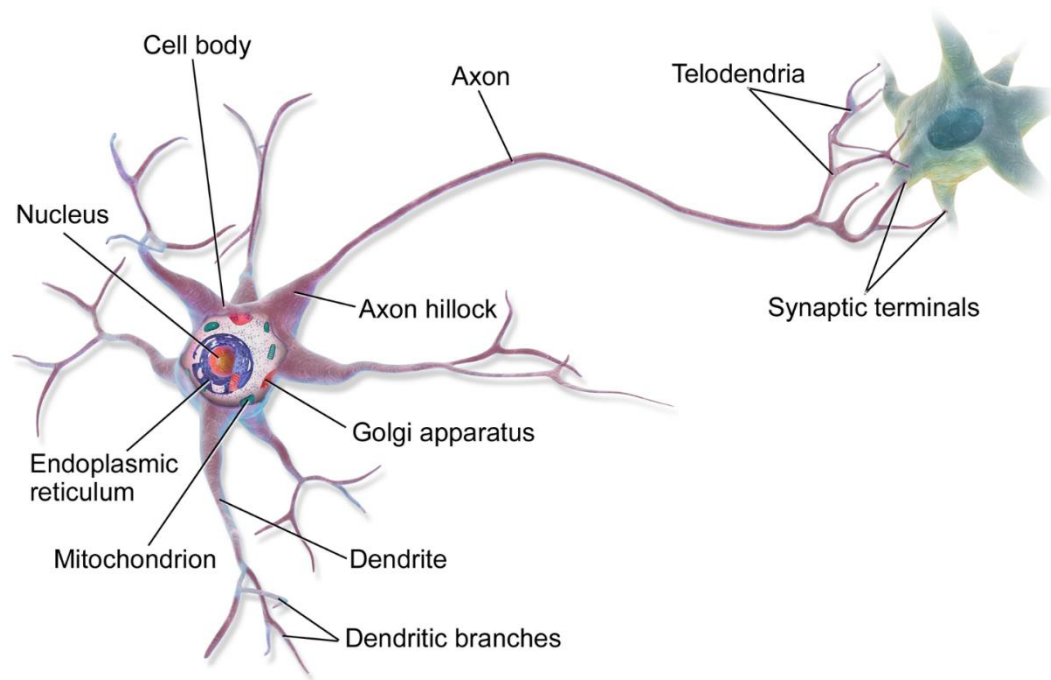
# Deriving random decision trees

- Depending on sampling, the decision trees are different

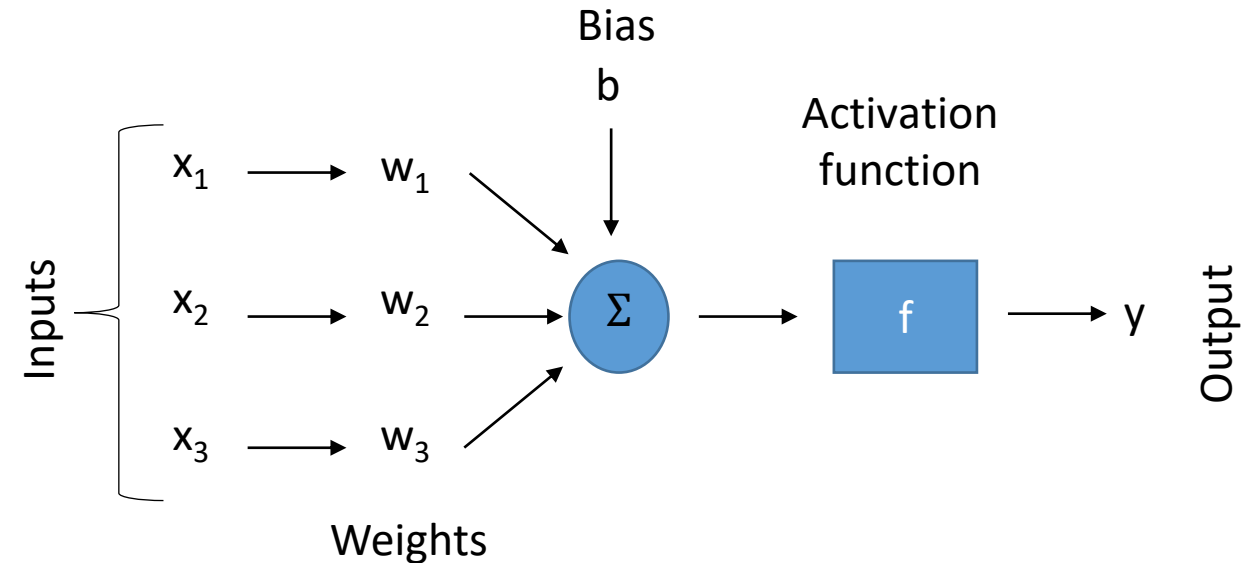


# Neural networks

- How biologists see neurons



- How computer scientists see neurons  
“perceptron”



$$y = f(w_1x_1 + w_2x_2 + w_3x_3 + b)$$

Neuron image source:

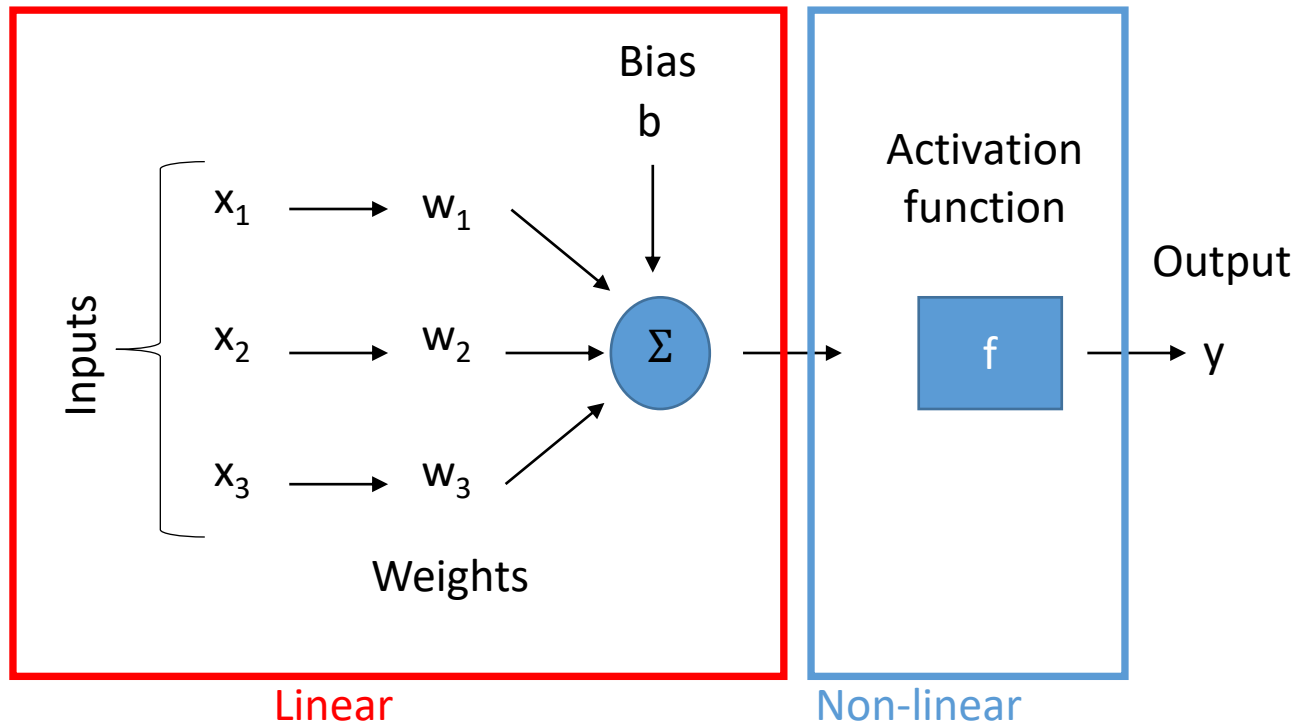
[https://commons.wikimedia.org/wiki/File:Blausen\\_0657\\_MultipolarNeuron.png](https://commons.wikimedia.org/wiki/File:Blausen_0657_MultipolarNeuron.png)

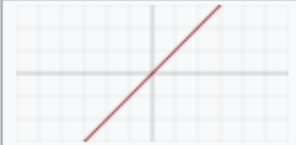



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# Activation functions

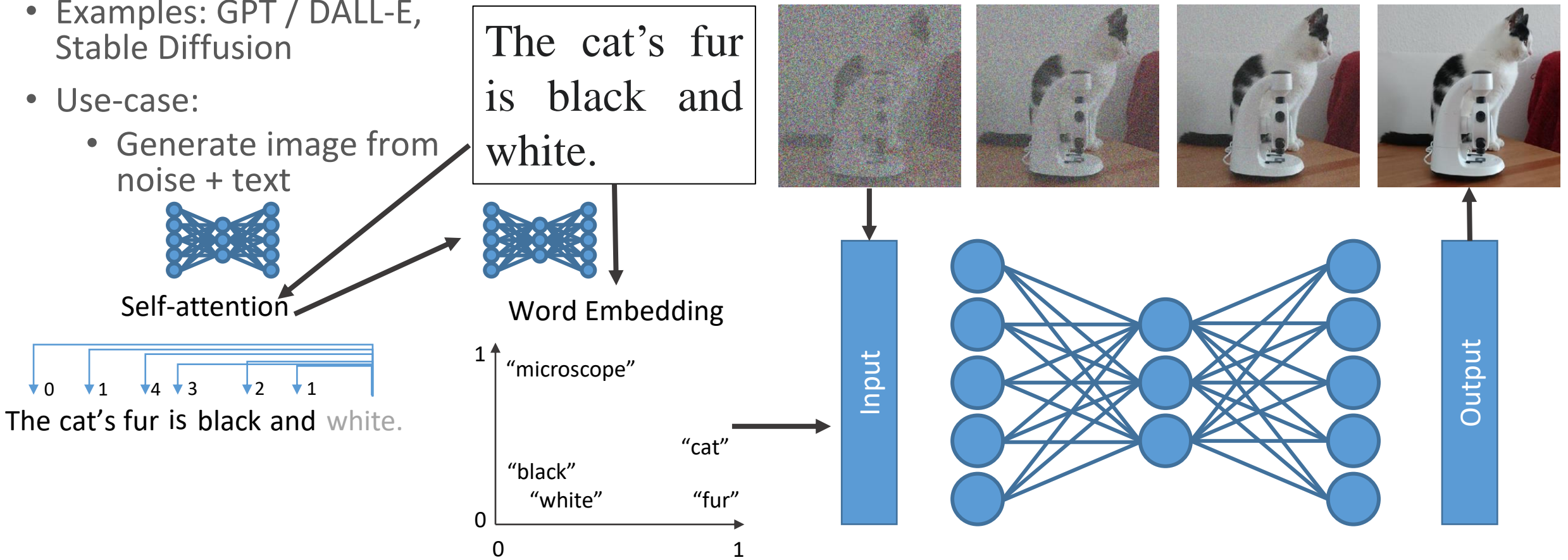
- Introduction of *non-linearity* and *activation functions* enabled what we call *deep-learning* today.



Identity		$x$
Binary step		$\begin{cases} 0 & \text{if } x < 0 \\ 1 & \text{if } x \geq 0 \end{cases}$
Logistic, sigmoid, or soft step		$\sigma(x) \doteq \frac{1}{1 + e^{-x}}$
Rectified linear unit (ReLU) <sup>[8]</sup>		$(x)^+ \doteq \begin{cases} 0 & \text{if } x \leq 0 \\ x & \text{if } x > 0 \end{cases}$ $= \max(0, x) = x \mathbf{1}_{x>0}$

# Generative AI / Large Language Models

- Combination of neural networks + other elements + various data sources
- Examples: GPT / DALL-E, Stable Diffusion
- Use-case:
  - Generate image from noise + text



## Train dataset (e.g. 80% of the data)

- Used for training directly

## Validation dataset (10% of the data)

- After every iteration see if the model overfits

## Test dataset (10% of the data)

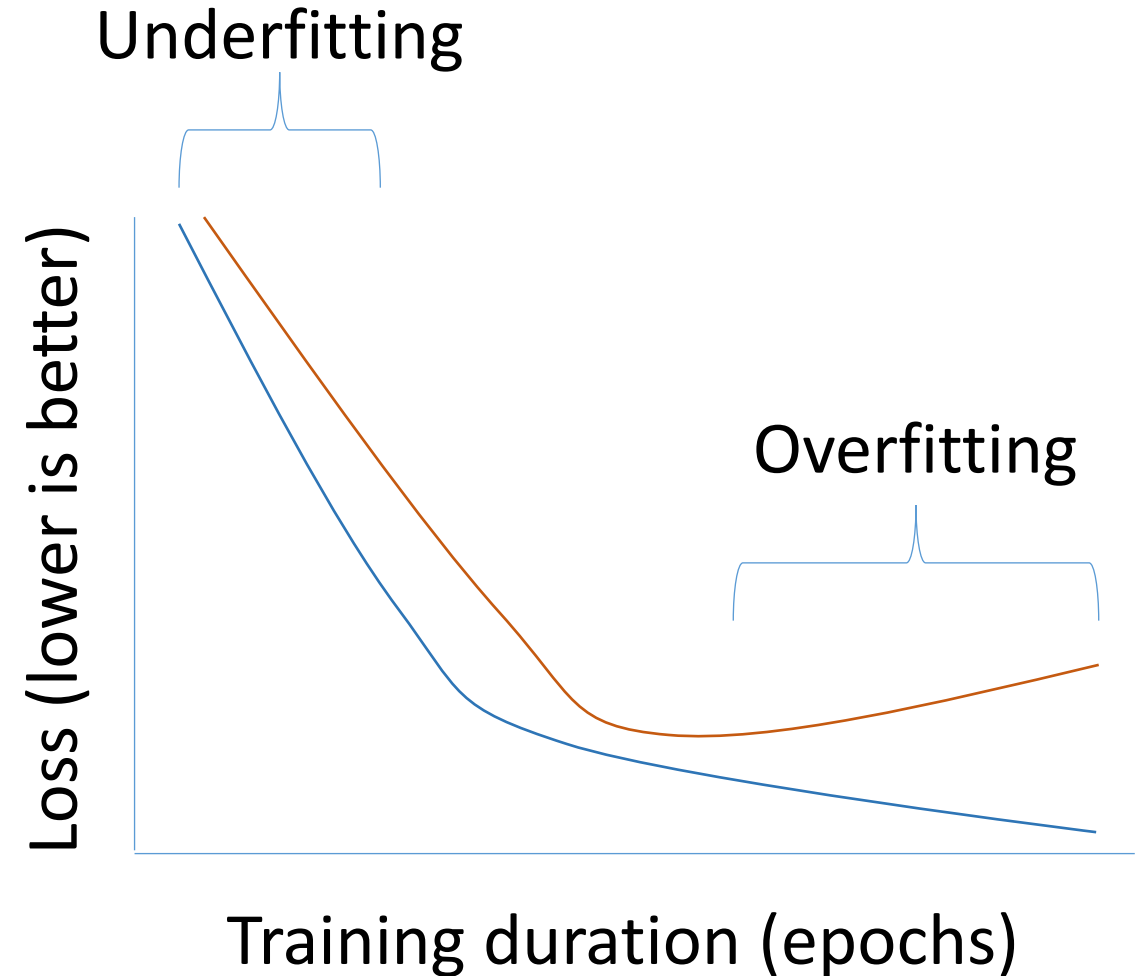
- Final evaluation after training is finished (once)

## Underfitting

- A trained model that is not even able to properly process the data it was trained on

## Overfitting

- A model that is able to process data it was trained on well
- It processes other data poorly



<https://towardsdatascience.com/how-to-split-data-into-three-sets-train-validation-and-test-and-why-e50d22d3e54c>

# Generative Artificial Intelligence

- Challenges
  - Data safety / security
  - Computational cost of training neural networks
    - CO2-footprint/climate change
    - Accessibility
  - Bias: “a nice photo of a human”
  - Hallucinations
  - Glitch tokens
  - False information / fake news
  - Debugging is hard



Model: Default (GPT-3.5)

R

Who developed Gradoop?



Gradoop is an open-source framework for scalable graph analytics. It was developed by the Hasso Plattner Institute (HPI) in Potsdam, Germany. The main contributors to Gradoop are researchers and developers from the HPI's Database Systems and Information Management (DIMA) group, led by Prof. Dr. Felix Naumann. The project started in 2015 as part of the HPI's research on large-scale graph data management and analysis. Since then, it has gained popularity among researchers and industry professionals working with graph data.

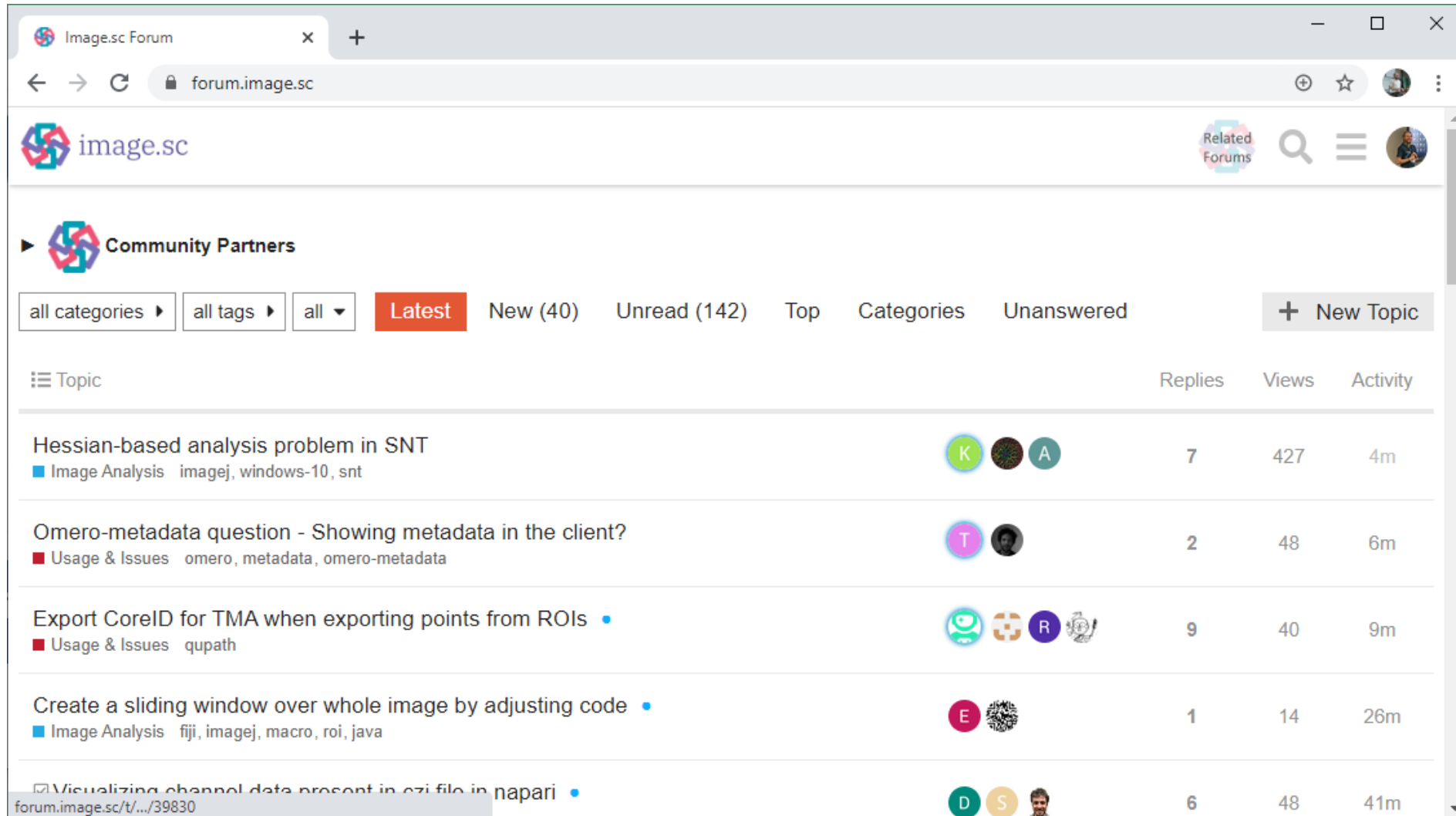
Sources: <https://replicate.com/stability-ai/stable-diffusion>  
<https://chat.openai.com/>



# Final remarks

July 2023

- Visit <http://forum.image.sc> !

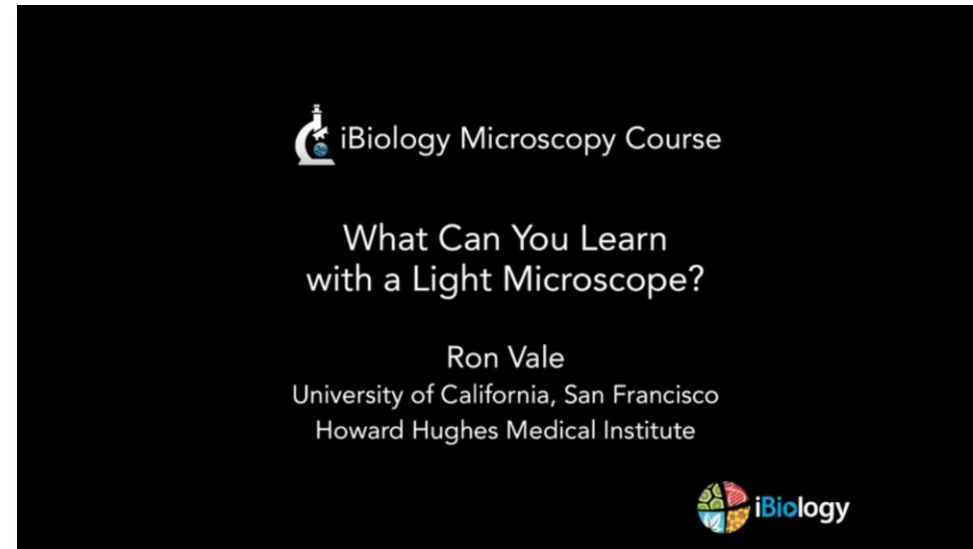
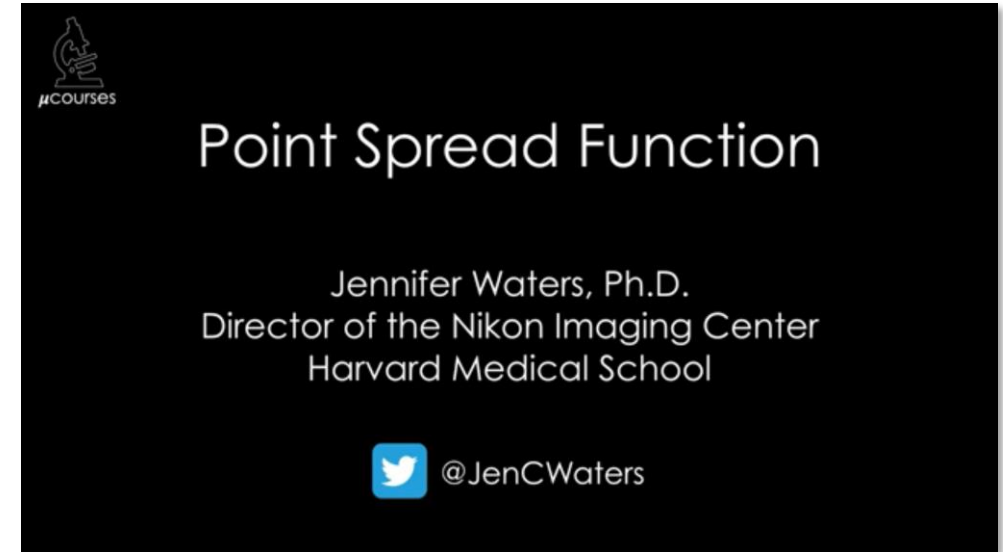


The screenshot shows the Image.sc Forum website. The browser address bar displays "forum.image.sc". The page header includes the "image.sc" logo and navigation icons. Below the header, there is a "Community Partners" section. The main content area features a navigation bar with filters: "all categories", "all tags", "all", "Latest" (highlighted in red), "New (40)", "Unread (142)", "Top", "Categories", "Unanswered", and a "+ New Topic" button. Below the navigation bar is a table of forum topics with columns for "Topic", "Replies", "Views", and "Activity".

Topic	Replies	Views	Activity
<a href="#">Hessian-based analysis problem in SNT</a> Image Analysis imagej, windows-10, snt	7	427	4m
<a href="#">Omero-metadata question - Showing metadata in the client?</a> Usage & Issues omero, metadata, omero-metadata	2	48	6m
<a href="#">Export CoreID for TMA when exporting points from ROIs</a> Usage & Issues qupath	9	40	9m
<a href="#">Create a sliding window over whole image by adjusting code</a> Image Analysis fiji, imagej, macro, roi, java	1	14	26m
<a href="#">Visualizing channel data present in ozi file in napari</a> forum.image.sc/t/.../39830	6	48	41m

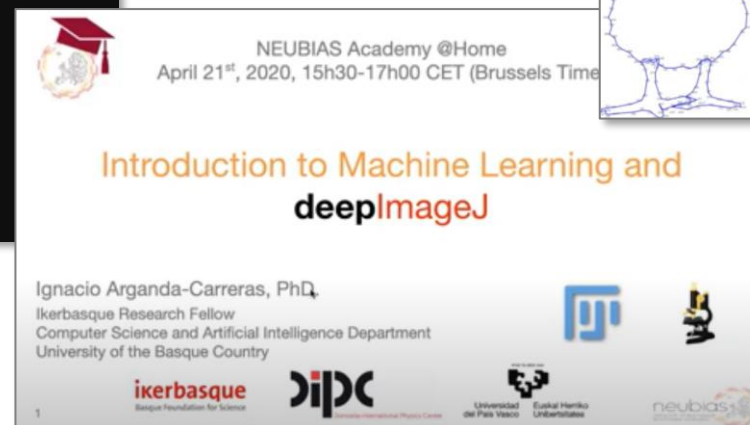
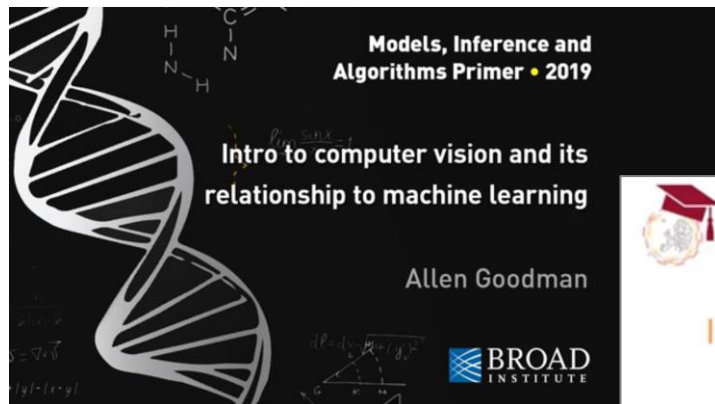
## Microscopy

- BioDIP Dresden Light Microscopy Course: [https://youtu.be/60\\_jgZtyR6U](https://youtu.be/60_jgZtyR6U)
- Microcourses: [https://youtu.be/Tkc\\_GOCjx7E](https://youtu.be/Tkc_GOCjx7E)
- iBiology Microscopy Course: <https://youtu.be/4c5ILWQmqRY>



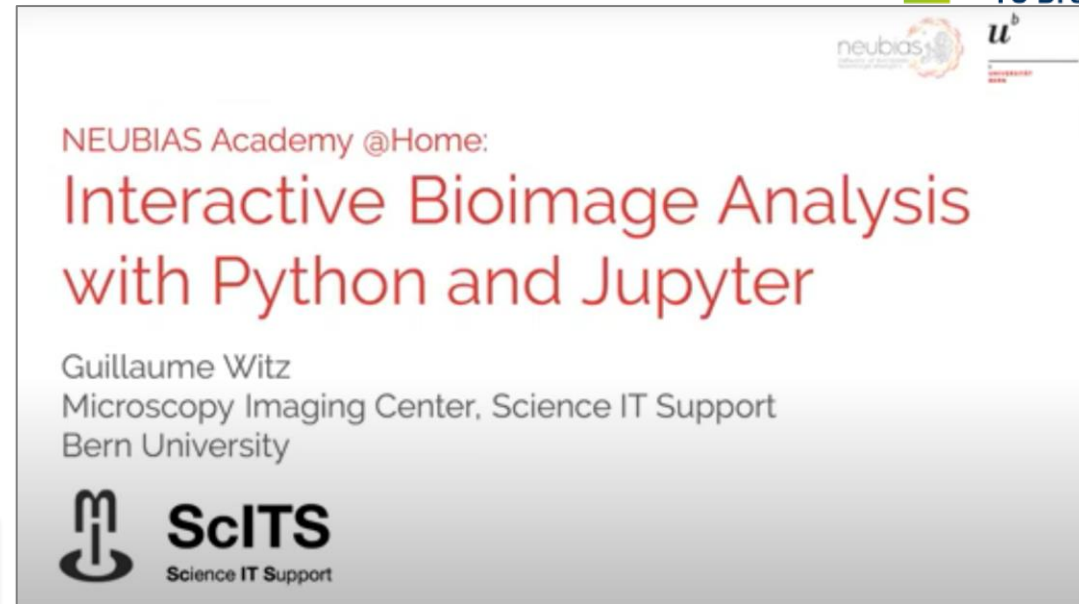
# What's next?

- More machine learning for Bio-Image analysis?
  - Computer Vision / machine learning: <https://youtu.be/Kzb5vTpvDBM>
  - Computer vision: <https://youtu.be/Smw3suzynho>
  - DeepImageJ: <https://youtu.be/0vTbsO8Vnuo>
  - CSBDeep: <https://youtu.be/ipp0mxfhwY>
  - StarDist: [https://youtu.be/Amn\\_eHRGX5M](https://youtu.be/Amn_eHRGX5M)
  - ilastik: <https://www.youtube.com/ilastikTeam>



# What's next?


- Image Analysis with Python
  - Python & Jupyter
    - <https://youtu.be/2KF8vBrp3Zw>
    - <https://youtu.be/Y3pB3wnOivE>
  - Scikit-image
    - [https://youtu.be/pZATswy\\_IsQ](https://youtu.be/pZATswy_IsQ)
    - <https://youtu.be/d1CIV9irQAY>
  - Napari
    - <https://youtu.be/VgvDSq5aCDQ>



neubias  
u<sup>b</sup>

NEUBIAS Academy @Home:  
**Interactive Bioimage Analysis  
with Python and Jupyter**

Guillaume Witz  
Microscopy Imaging Center, Science IT Support  
Bern University

 **ScITS**  
Science IT Support

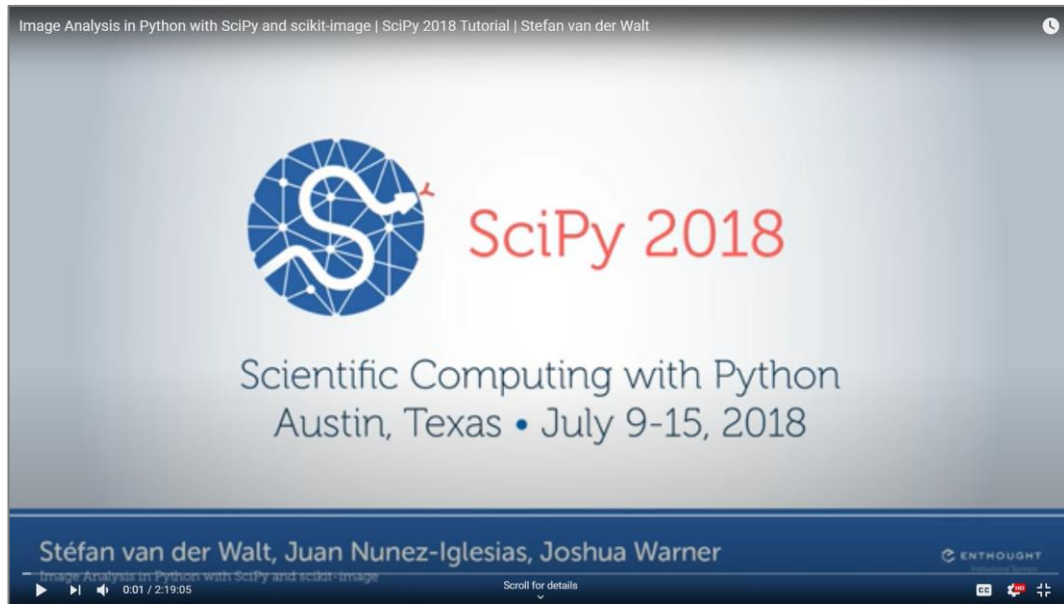



Image Analysis in Python with SciPy and scikit-image | SciPy 2018 Tutorial | Stefan van der Walt

 **SciPy 2018**

Scientific Computing with Python  
Austin, Texas • July 9-15, 2018

Stéfan van der Walt, Juan Nunez-Iglesias, Joshua Warner

ENTHOUGHT  
Image Analysis in Python with SciPy and scikit-image  
0:01 / 2:19:05





**napari**  
a multi-dimensional image viewer for Python

June 4th 2020 - NEUBIAS Academy @Home  
Nicholas Sofroniew

Chan  
Zuckerberg  
Initiative

neubias  
cost  
EUROPEAN COOPERATION  
IN SCIENCE & TECHNOLOGY

# What's next?

## More general

- Python for Microscopists + beyond: <https://youtube.com/digitalsreeni>
- Image data integrity: [https://youtu.be/c\\_Oi2HKom\\_Y](https://youtu.be/c_Oi2HKom_Y)
- Coloc: <https://youtu.be/cOrCz4qc8DI>
- Automated microscopy: <https://youtu.be/w0ERCrKx4gk>

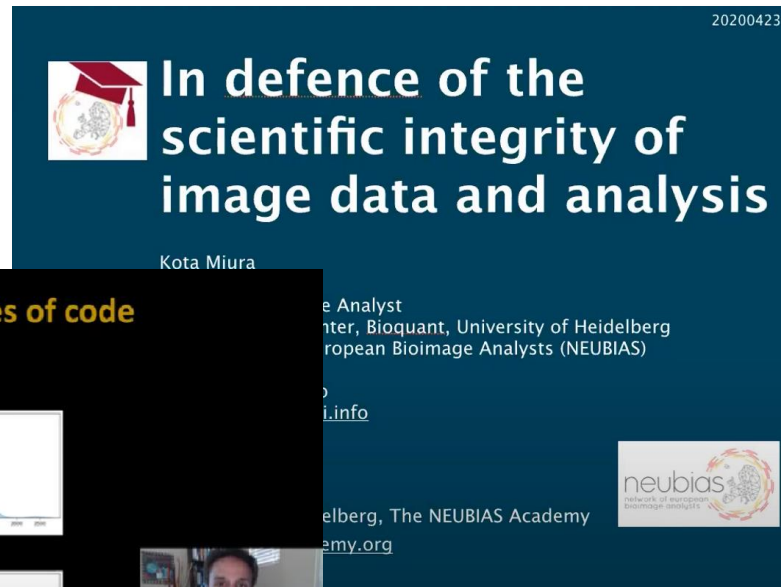


Automating microscopy acquisition with deep learning and augmented reality.

By Dr. Dominic Waithe UKRI Innovation Fellow  
Weatherall Institute of Molecular Medicine, University of Oxford.

Bioimage Analysis in Python Course Dec 2019  
Cambridge

BBSRC bioscience for the future  
MRC Medical Research Council  
UK Research and Innovation  
UNIVERSITY OF OXFORD



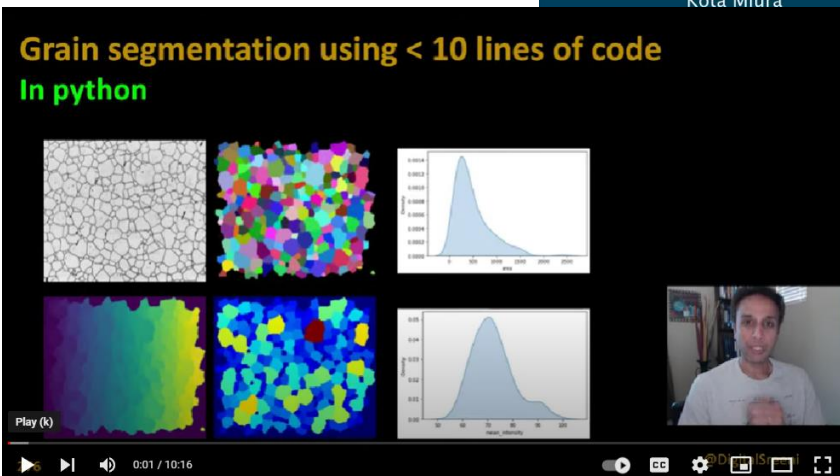
20200423

In defence of the scientific integrity of image data and analysis

Kota Miura

Analyst  
Center, Bioquant, University of Heidelberg  
European Bioimage Analysts (NEUBIAS)

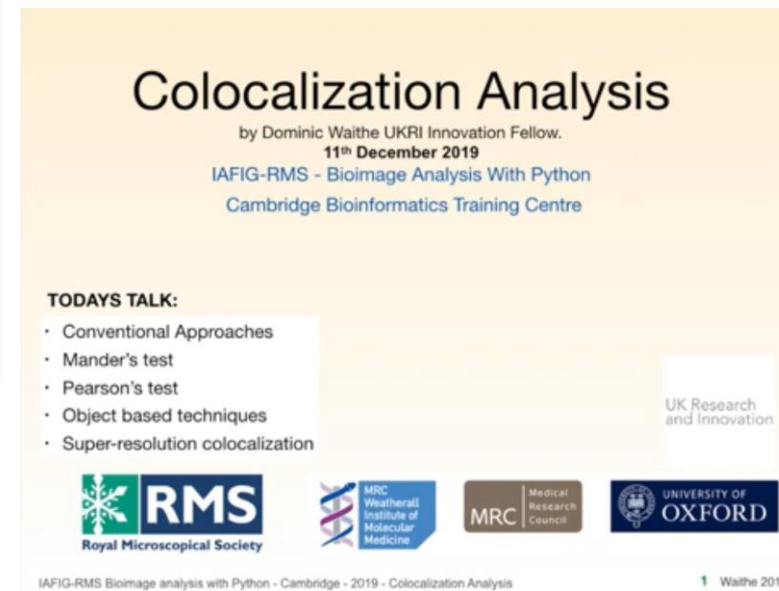
neubias  
network of european  
bioimage analysts



Grain segmentation using < 10 lines of code  
in python

Play (k)

0:01 / 10:16



Colocalization Analysis

by Dominic Waithe UKRI Innovation Fellow.  
11<sup>th</sup> December 2019  
IAFIG-RMS - Bioimage Analysis With Python  
Cambridge Bioinformatics Training Centre

TODAYS TALK:

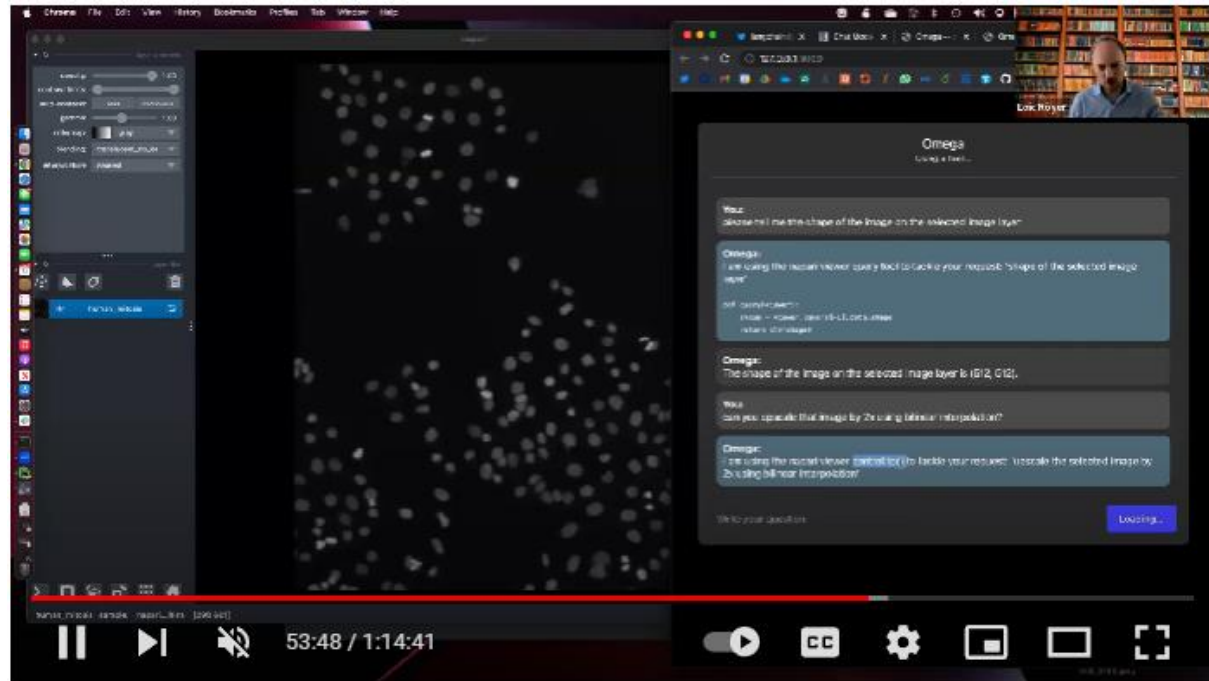
- Conventional Approaches
- Mander's test
- Pearson's test
- Object based techniques
- Super-resolution colocalization

UK Research and Innovation

RMS Royal Microscopical Society  
MRC Weatherall Institute of Molecular Medicine  
MRC Medical Research Council  
UNIVERSITY OF OXFORD

IAFIG-RMS Bioimage analysis with Python - Cambridge - 2019 - Colocalization Analysis

1 Waithe 2019



Code review of Napari-ChatGPT by Loic Royer (CZI Biohub)

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[https://www.youtube.com/watch?v=JMo6Sn-L\\_j4](https://www.youtube.com/watch?v=JMo6Sn-L_j4)



DigitalSreeni

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This channel walks you through the entire process of learning to code in P... >



308 - An introduction to language models with focu...



311 - Fine tuning GPT2 using custom documents



309 - Training your own Chatbot using GPT



<https://www.youtube.com/c/digitalsreeni>

Picture source: <https://twitter.com/digitalsreeni/status/1541578740584415233>