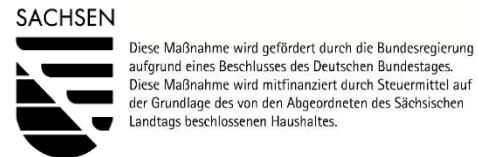


Supervised and Unsupervised Machine Learning for Bio-image Analysis

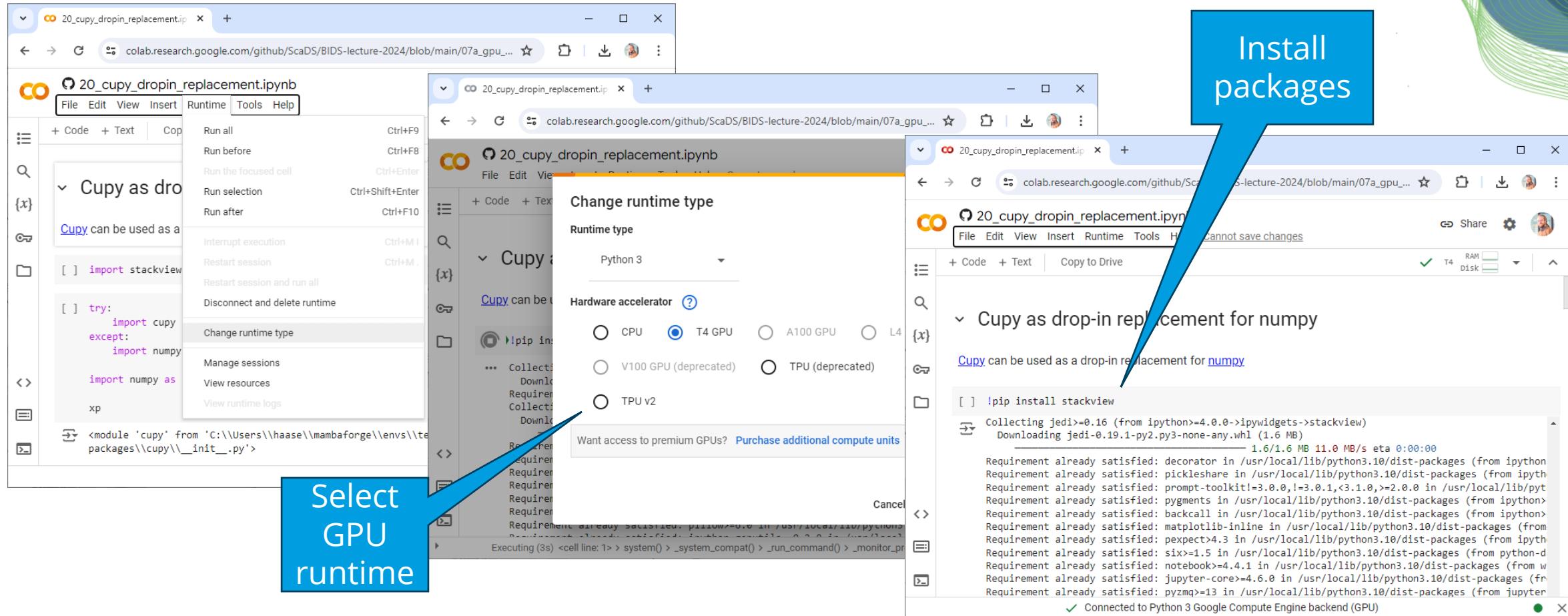
Robert Haase

Reusing materials from Johannes Soltwedel, Till Korten, Johannes Müller, Laura Žigutytė (TU Dresden), Ryan Savill (MPI-CBG), Matthias Täschner (ScaDS.AI/Uni Leipzig) and the Scikit-learn community.



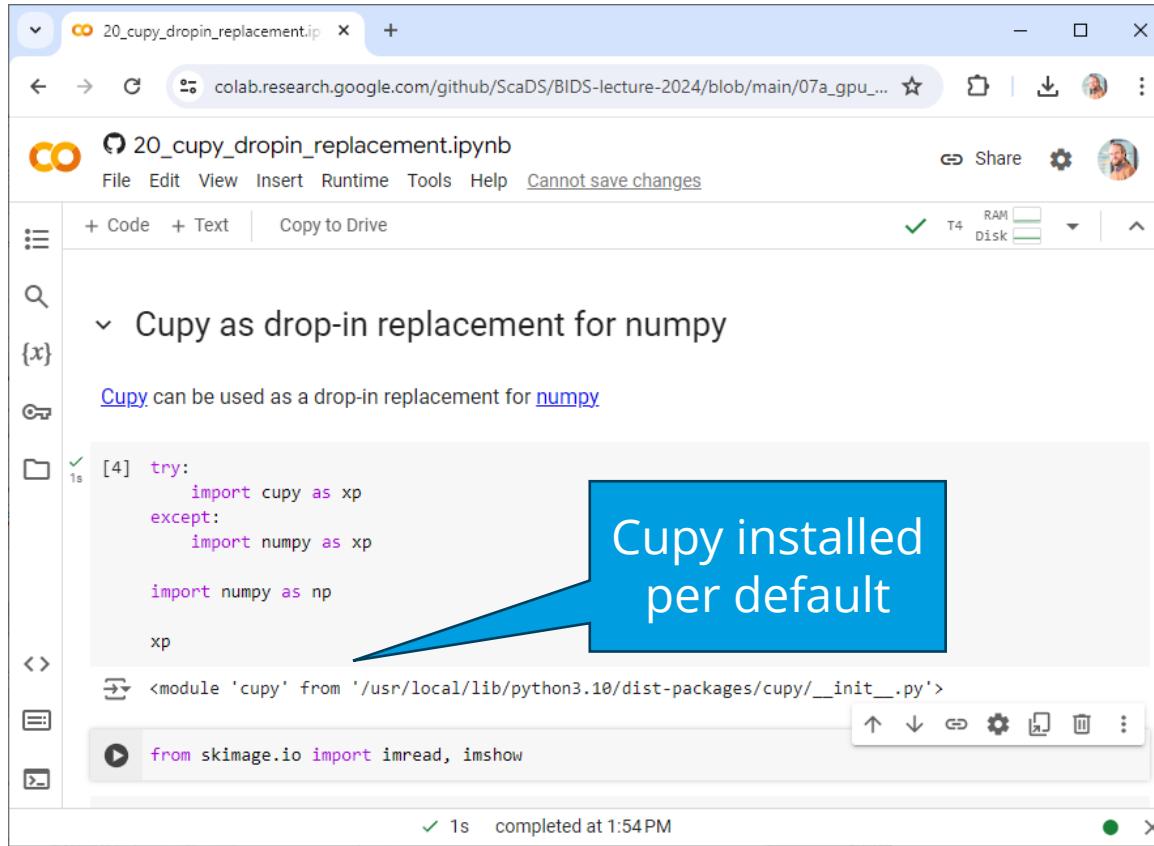
Follow-up: cupy

cupy was hard to make work. Consider using a GPU-runtime in Google Colab



Follow-up: cupy

cupy was hard to make work. Consider using a GPU-kernel in Google Colab

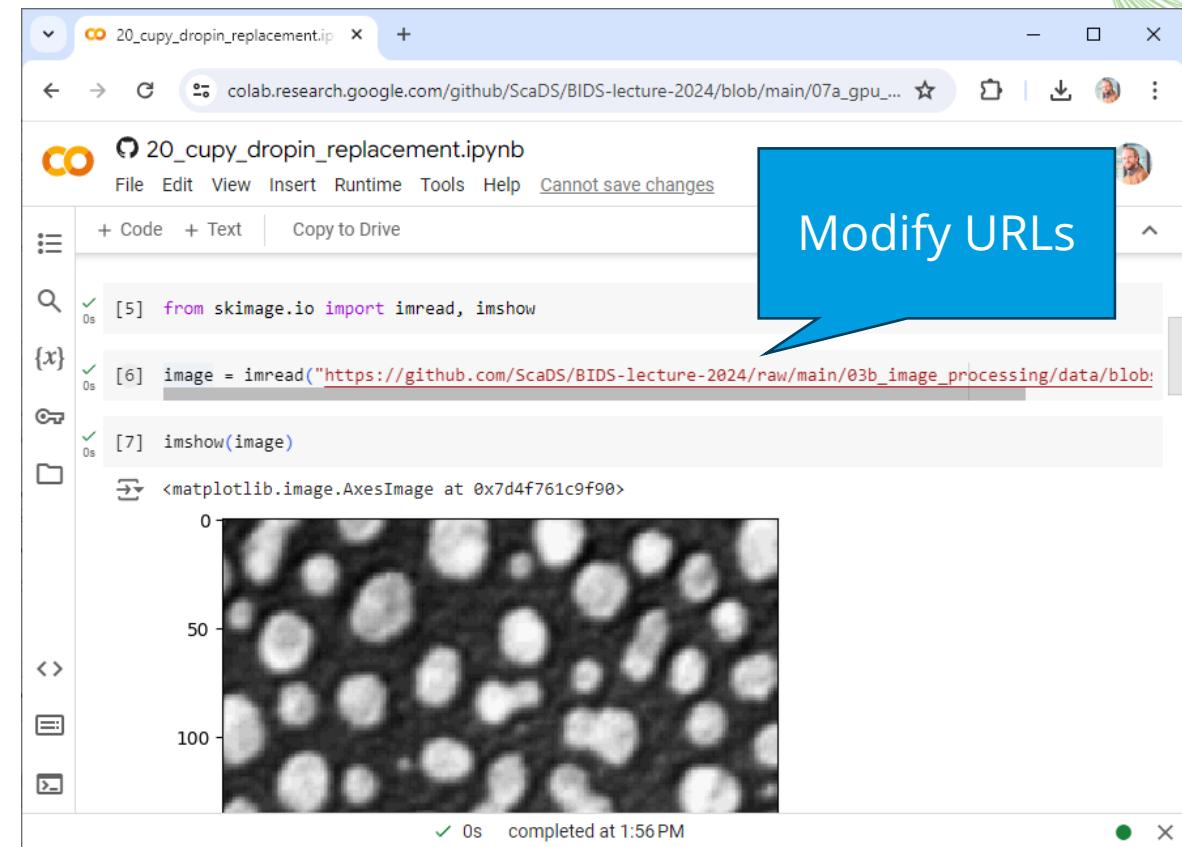


Cupy as drop-in replacement for numpy

Cupy can be used as a drop-in replacement for numpy

```
[4] try:  
    import cupy as xp  
except:  
    import numpy as xp  
  
import numpy as np  
  
xp
```

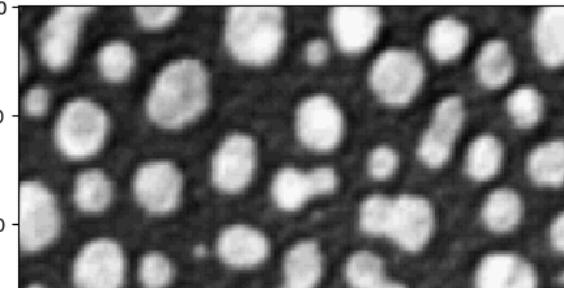
Cupy installed per default



Modify URLs

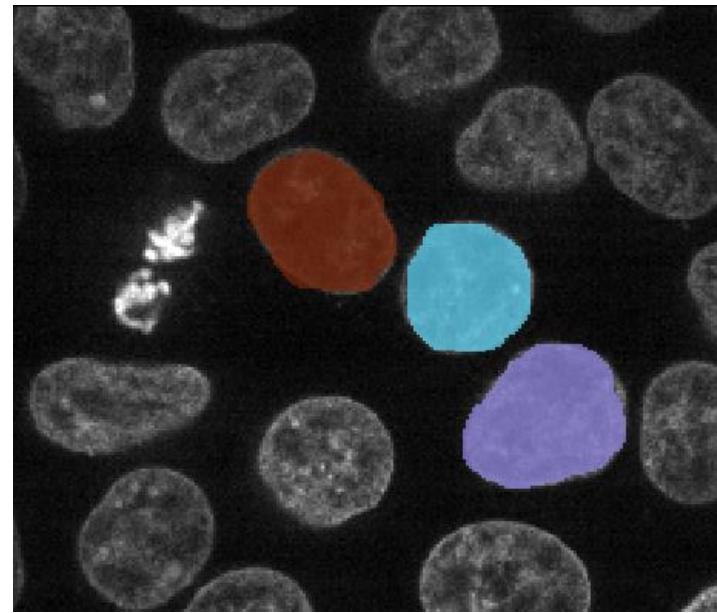
```
[5] from skimage.io import imread, imshow  
  
[6] image = imread("https://github.com/ScaDS/BIDS-lecture-2024/raw/main/03b_image_processing/data/blob:  
  
[7] imshow(image)
```

`<matplotlib.image.AxesImage at 0x7d4f761c9f90>`



Quiz: Recap

What kind of label image is this?



Instance
segmentation



Semantic
segmentation



Sparse
instance
segmentation



Sparse
semantic
segmentation



Terminology

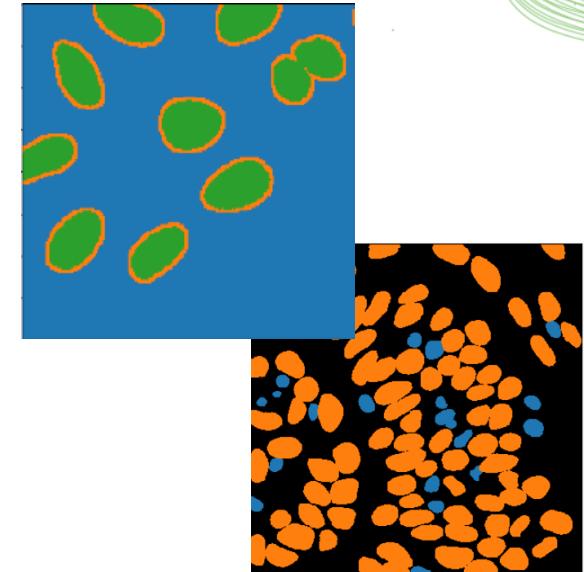
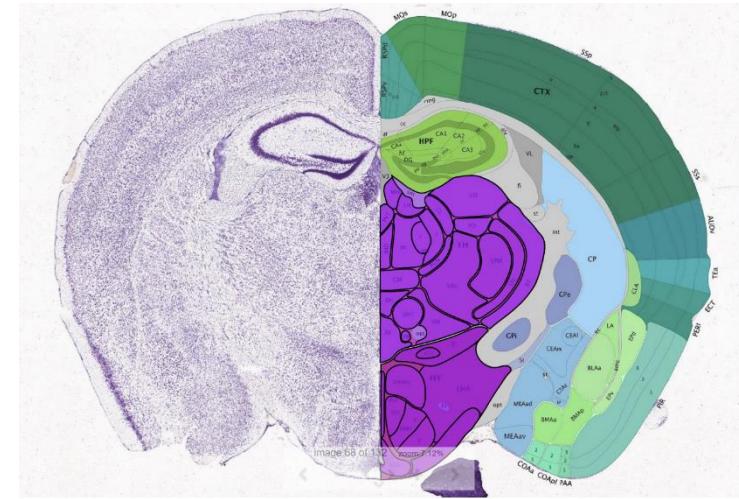
Instance segmentation



Instances:

- Cells, nuclei, cats, dogs, cars, trees

Semantic segmentation

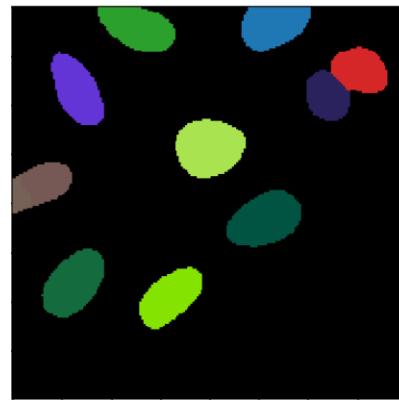


Regions:

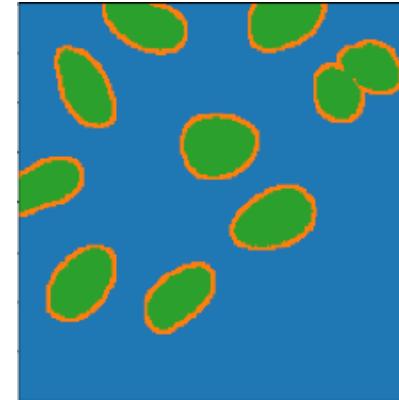
- Anatomical, geographical
- All pixels belonging to the same type of object have the same value

Terminology

Instance segmentation

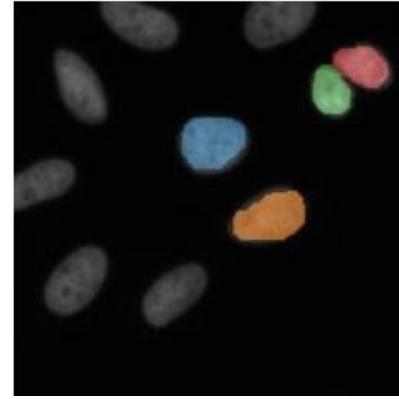


Semantic segmentation



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

Sparse instance annotation



Sparse semantic annotation

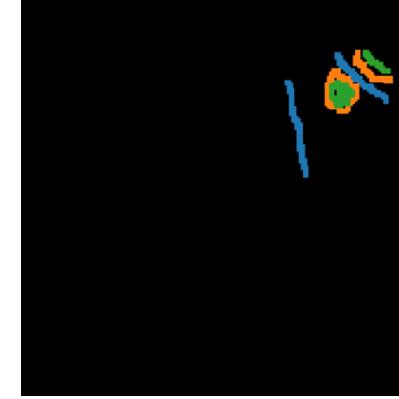


Image segmentation using thresholding

Recap: Finding the right workflow towards a good segmentation takes time

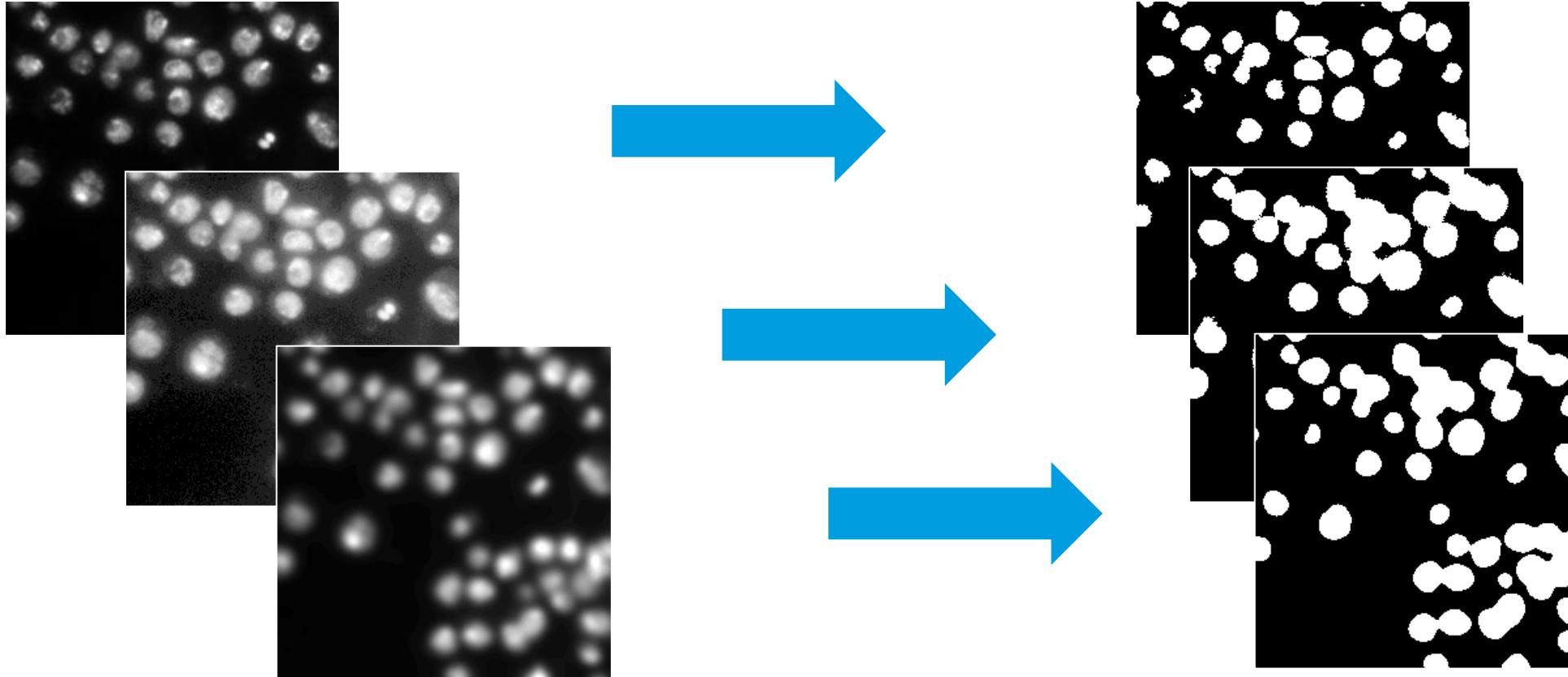


Image segmentation using thresholding

Recap: Combining images, e.g. using Difference of Gaussian (DoG)

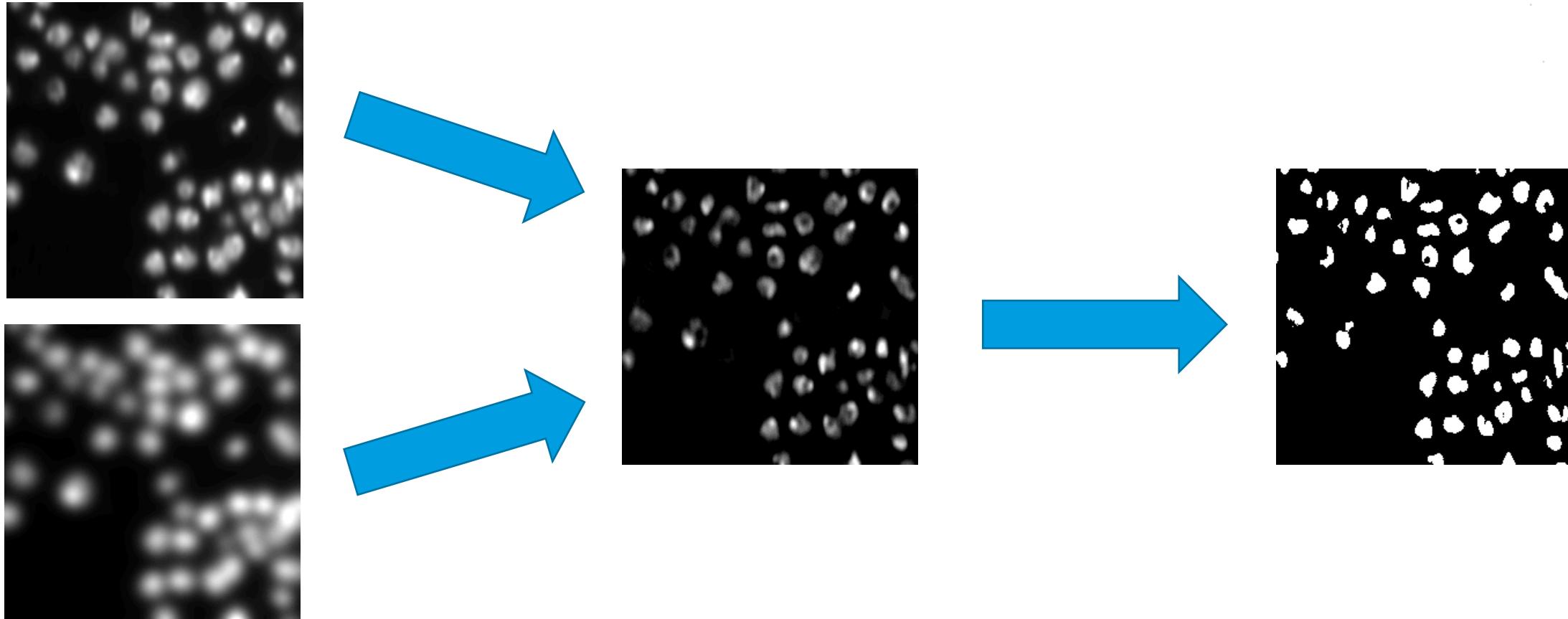
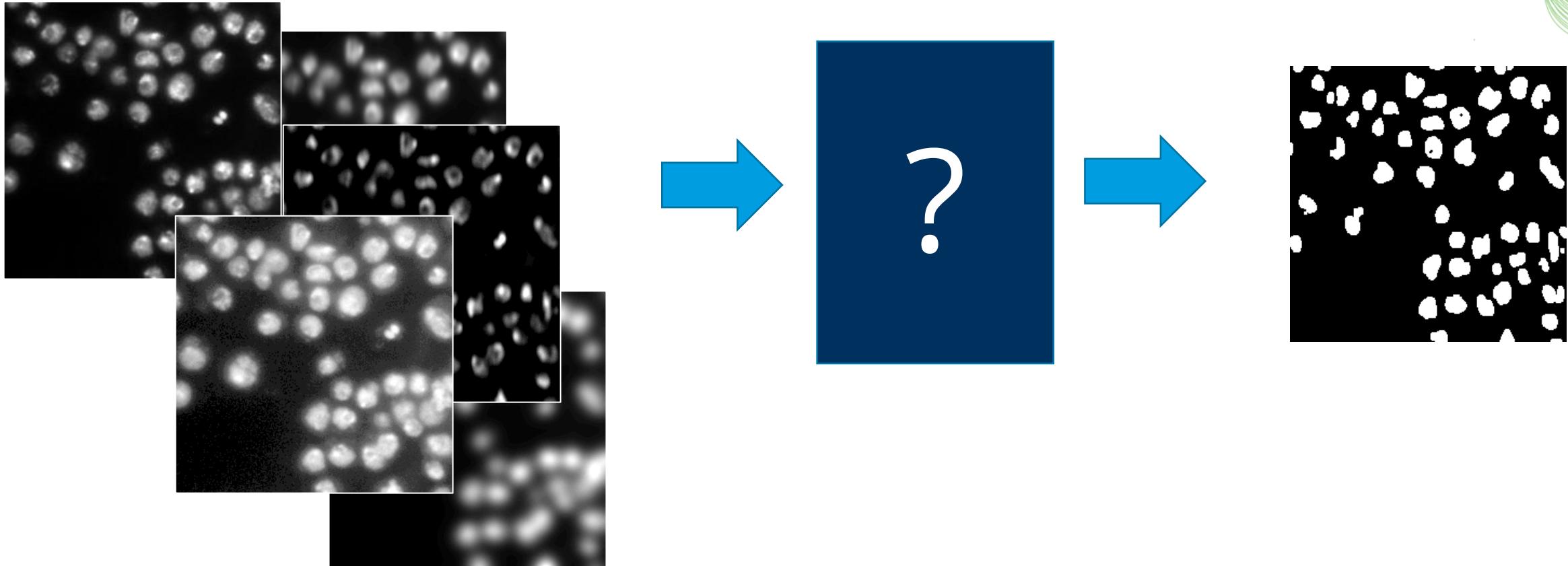


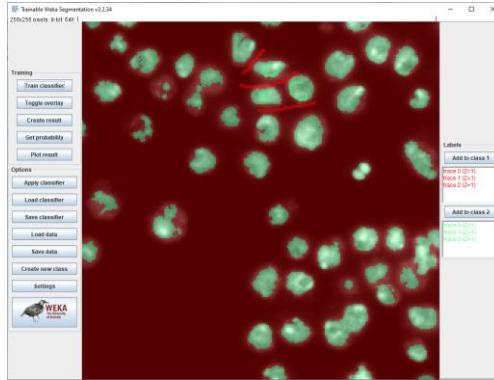
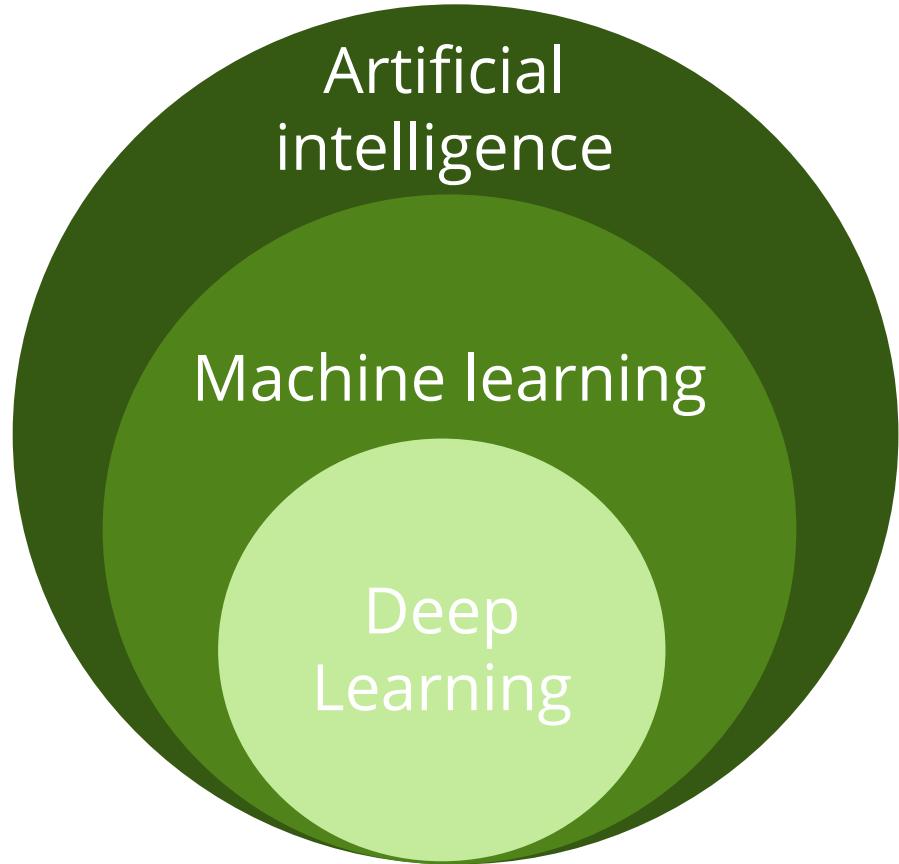
Image segmentation using thresholding

Might there be a technology for optimization which combination of images can be used to get the best segmentation result?

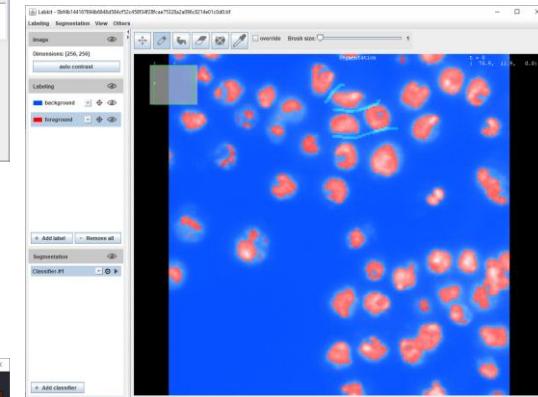


Machine learning

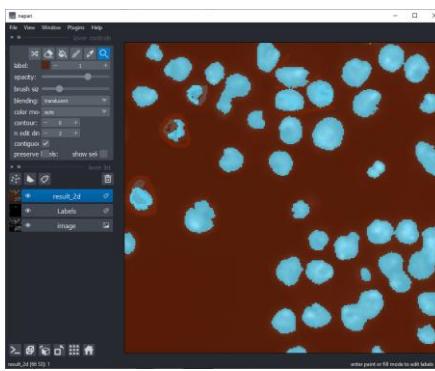
Finds more and more applications, also in life sciences.



Trainable Weka Segmentation.
<https://imagej.net/plugins/tws/>



LabKit
<https://imagej.net/plugins/labkit/>

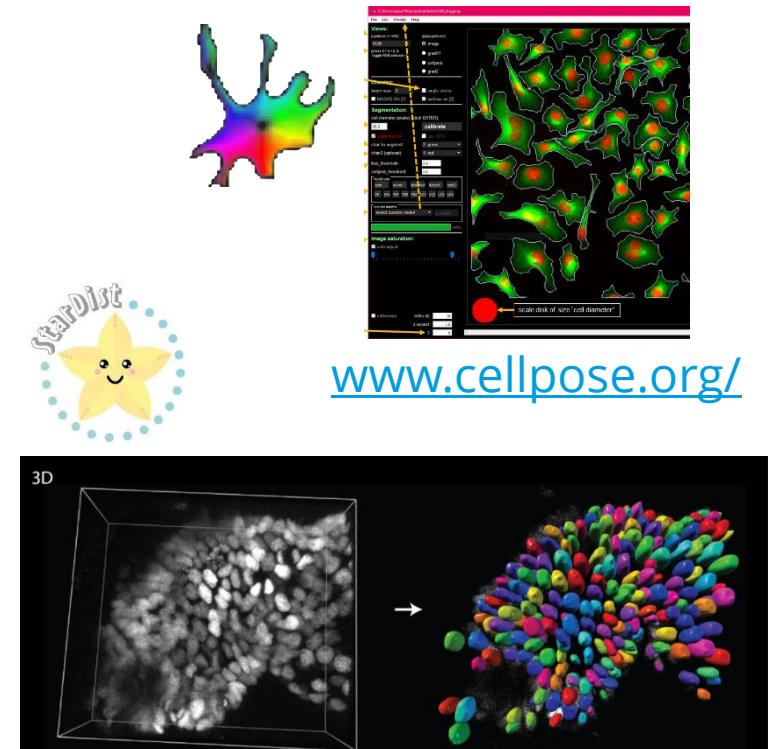
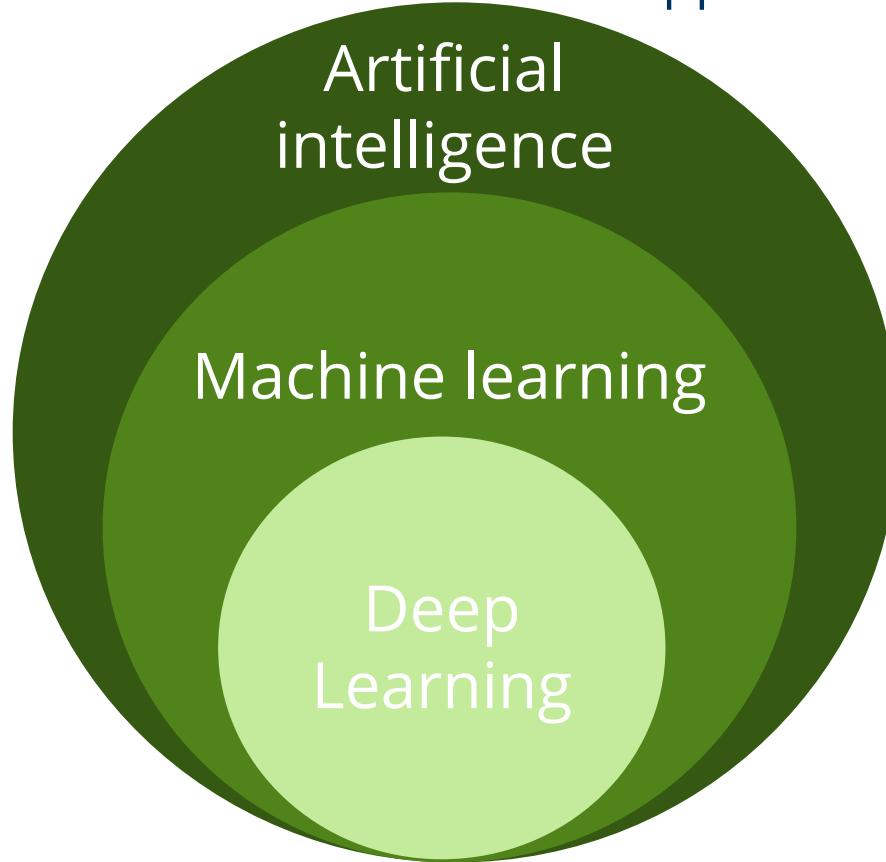


Python / scikit-learn /
napari / apoc

Machine learning

A research field in computer science

Finds more and more applications, also in life sciences.



<https://github.com/stardist/stardist>

www.cellpose.org/

BiolImage Model Zoo
Advanced AI models in one-click
Integrate with Fiji, Ilastik, ImJoy and more
Try model instantly with BioEngine
Contribute your models via Github
Link models to datasets and applications
[Explore the Zoo](#)

Community Partners:

All [models](#) [applications](#) [datasets](#)

Type a keyword and press enter

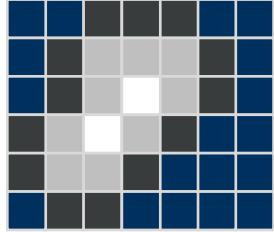
Mitochondria Segmentation for EM
Mitochondria segmentation for electron microscopy.
3d electron-microscopy mitochondria instance-segmentation ...

<https://bioimage.io/>

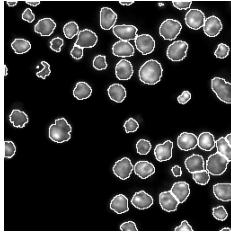
Machine learning

Automatic construction of predictive models from given data

Pixels,



Objects,

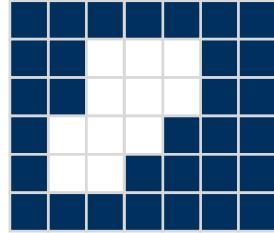


Images



Audio, Text, Measurements, ...

Dense
Segmentation
/ Binarization



Object
classification

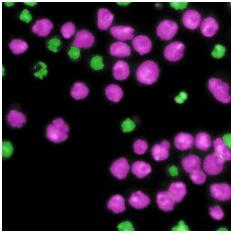
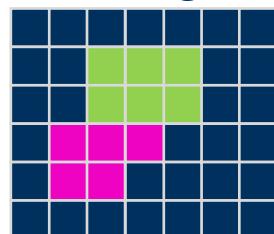


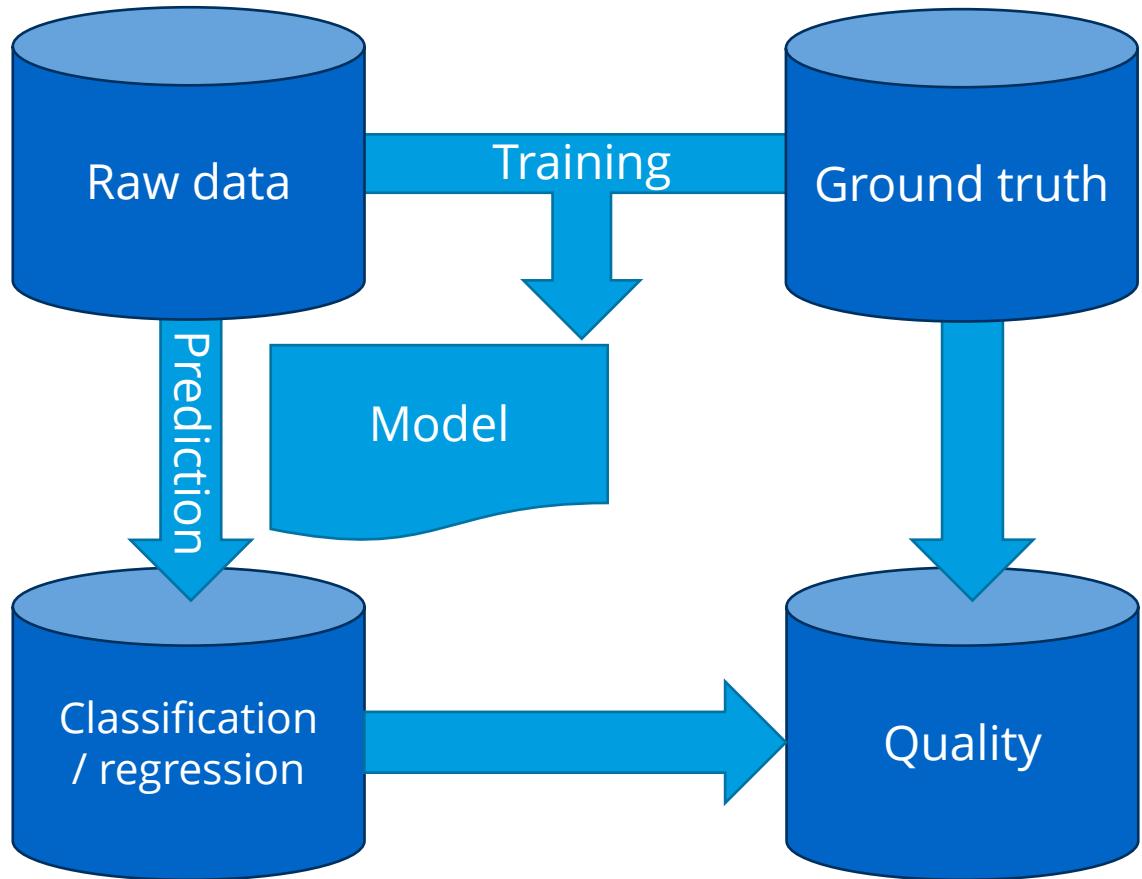
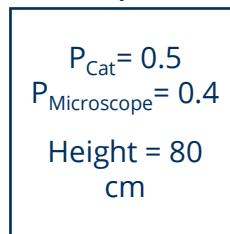
Image
classification



Instance segmentation



Cont. quantity

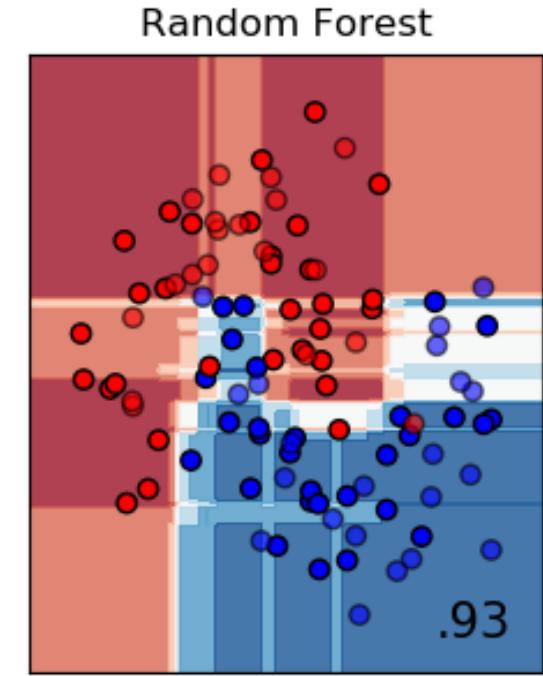
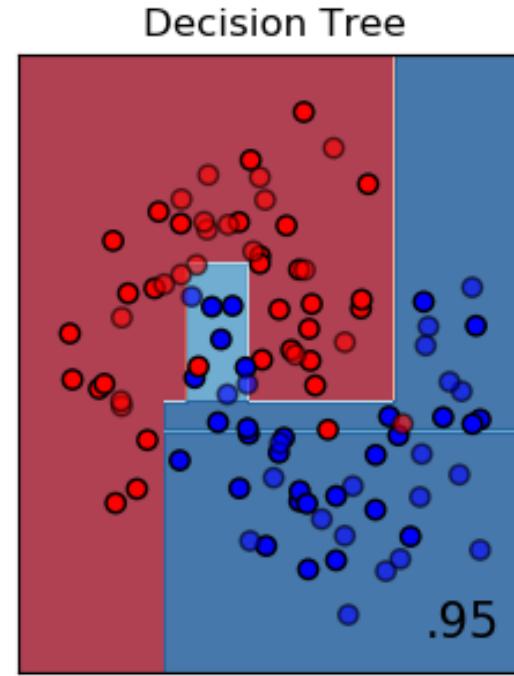
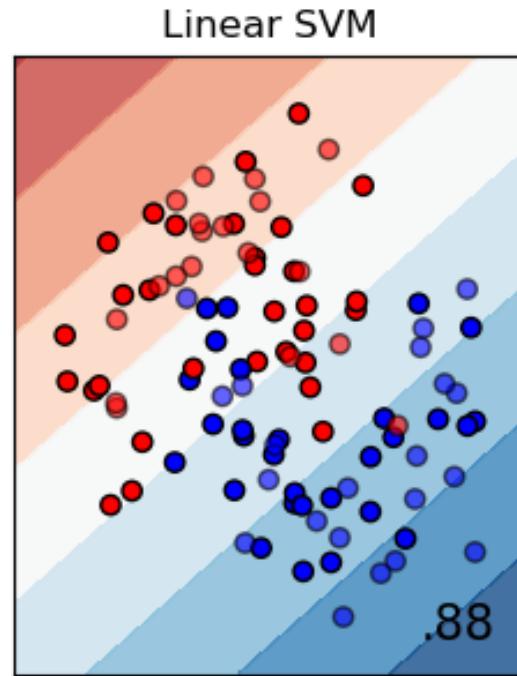
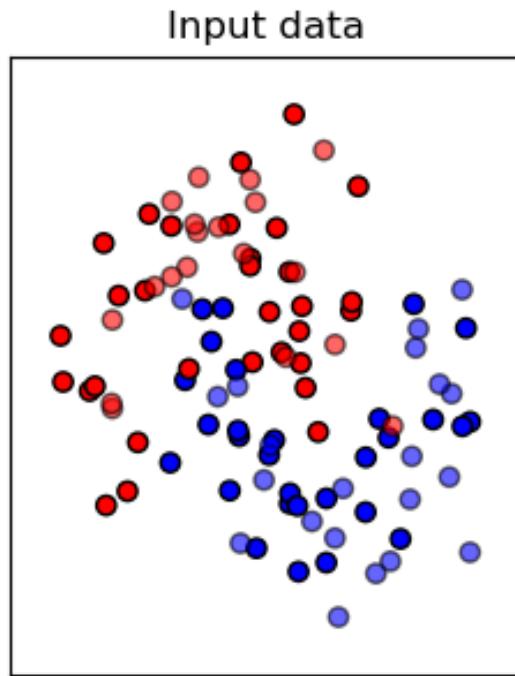


Annotated
raw data,
usually
generated by
humans

Precision,
Recall

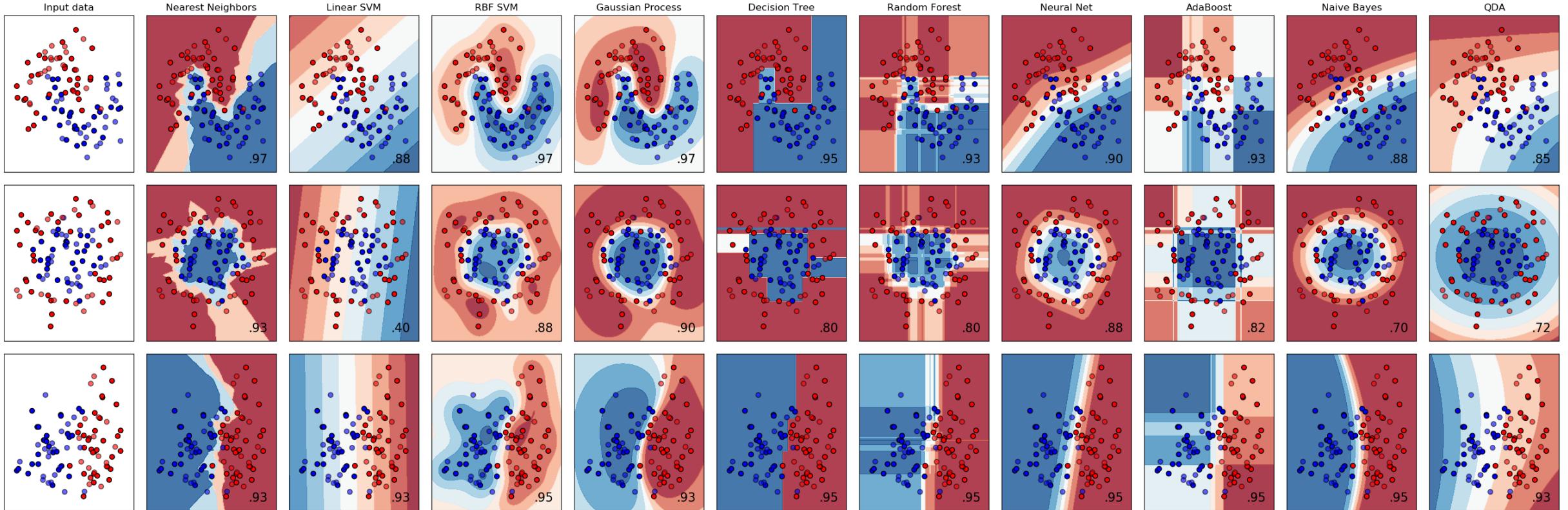
Goal

Guess classification (color) from position of a sample in parameter space.



Approaches

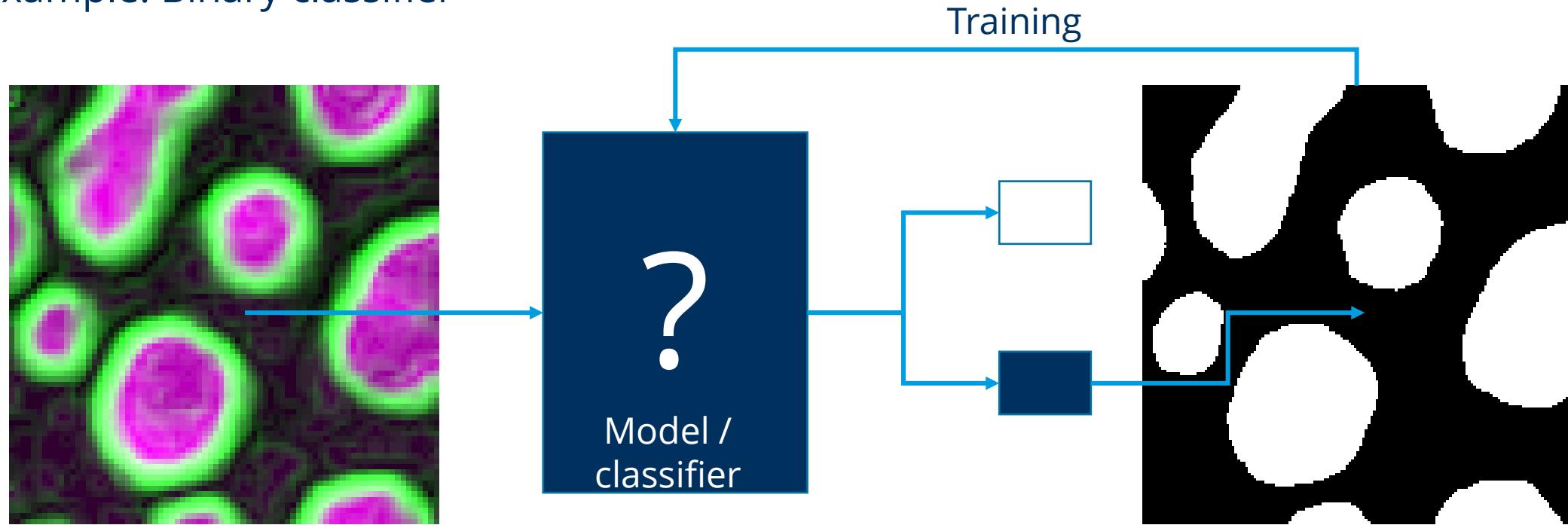
The right approach depends on data, computational resources and desired quality



Machine learning for image segmentation

Supervised machine learning: We give the computer some ground truth to learn from
The computer derives a *model* or a *classifier* which can judge if a pixel should be
foreground (white) or background (black)

Example: Binary classifier



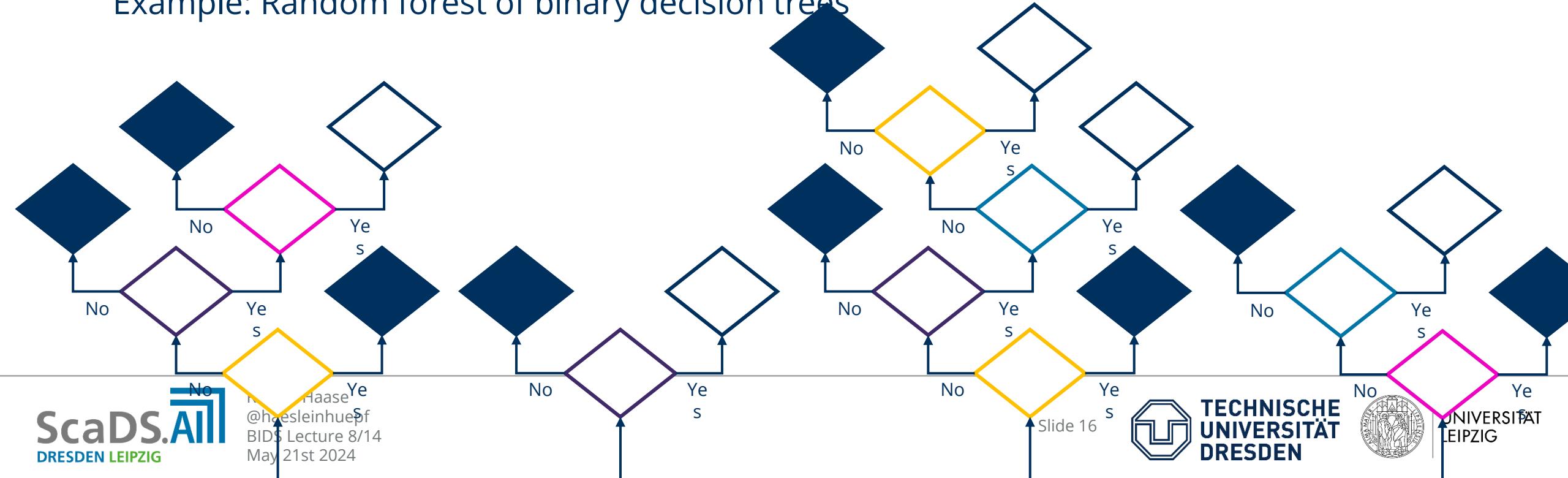
Random forest based image segmentation

Decision trees are classifiers, they decide if a pixel should be white or black

Random decision trees are randomly initialized, afterwards evaluated and selected

Random forests consist of many random decision trees

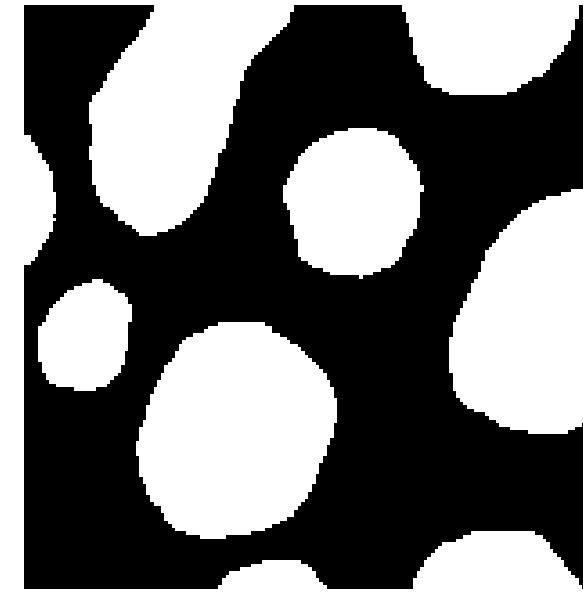
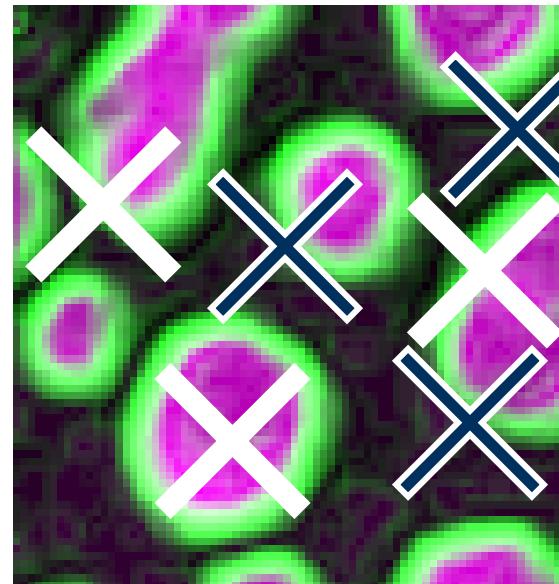
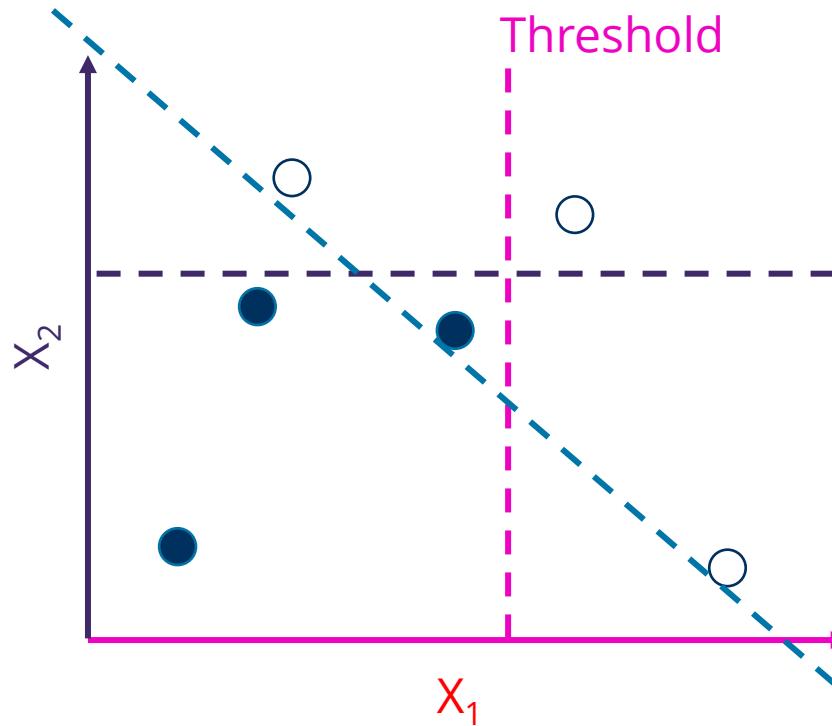
Example: Random forest of binary decision trees



Deriving random decision trees

For efficient processing, we randomly *sample* our data set

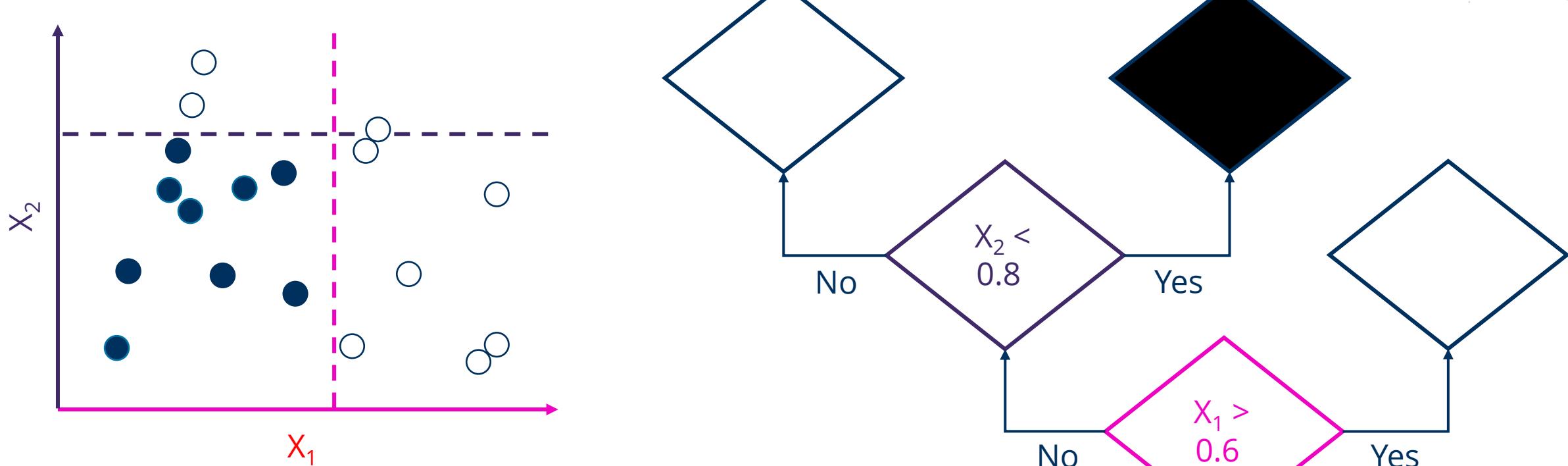
- Individual pixels, their intensity and their classification



Note: You cannot use a single threshold to make the decision

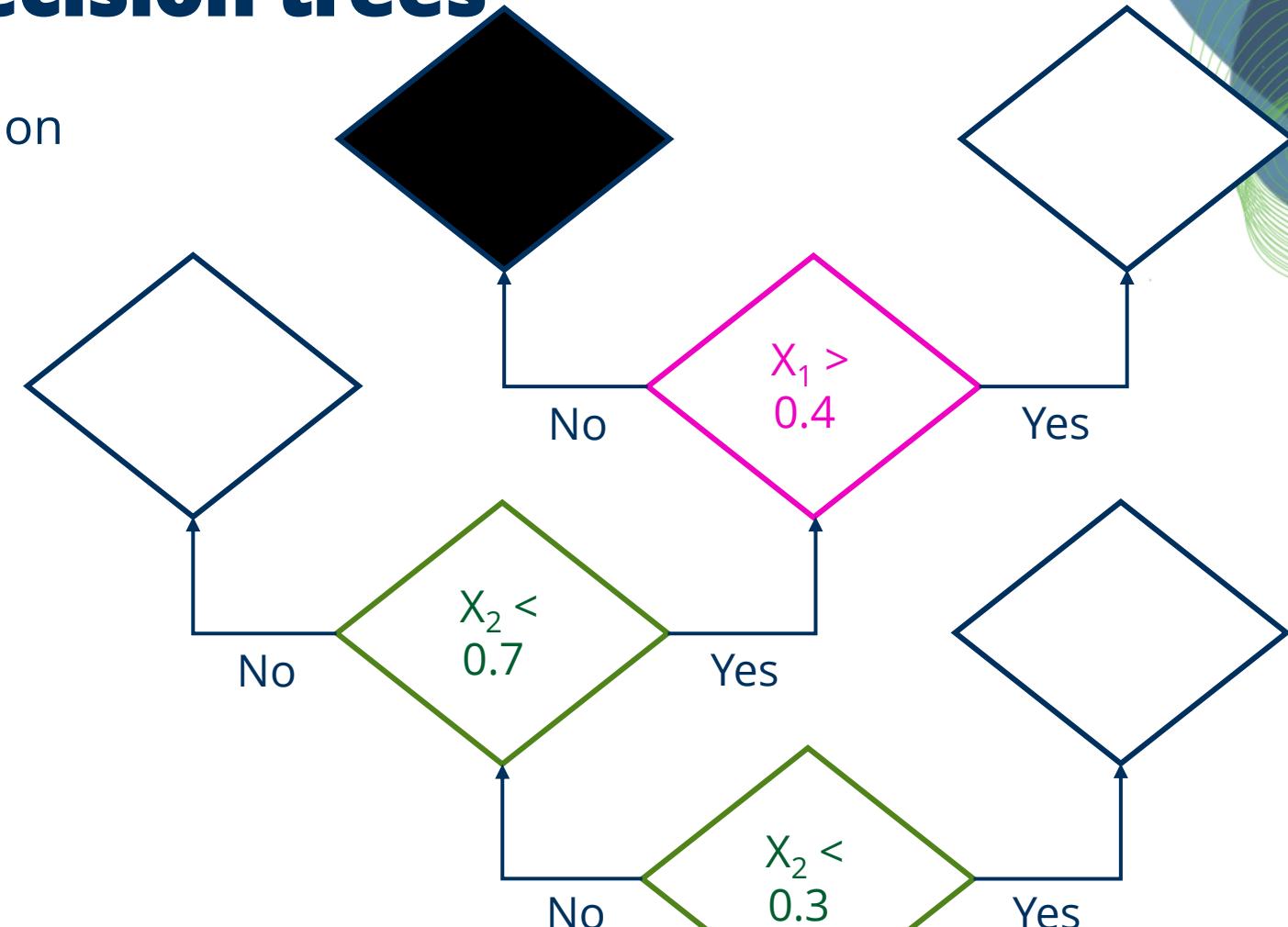
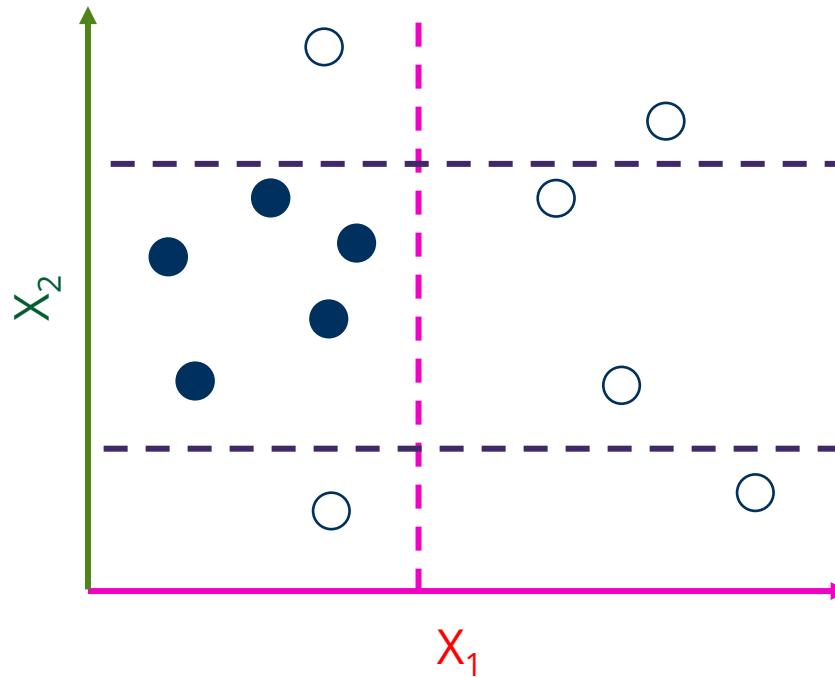
Deriving random decision trees

Decision trees combine several thresholds on several parameters



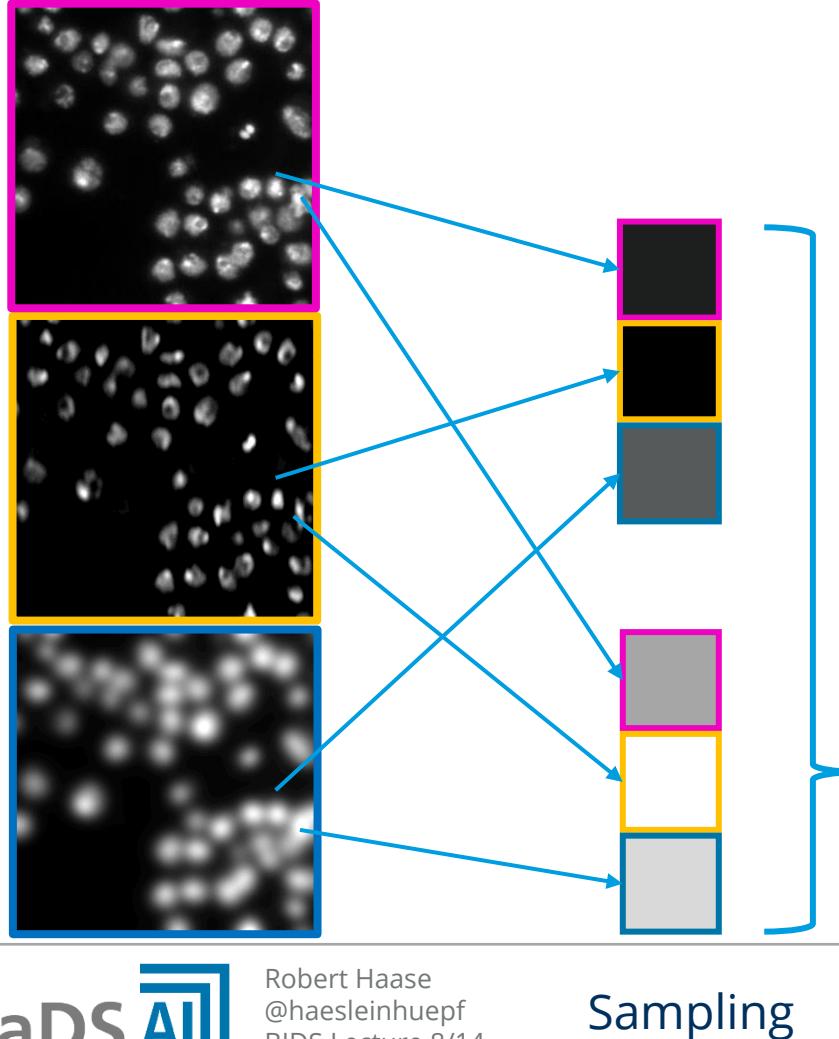
Deriving random decision trees

Depending on sampling, the decision trees are different



Random Forest Pixel Classifiers

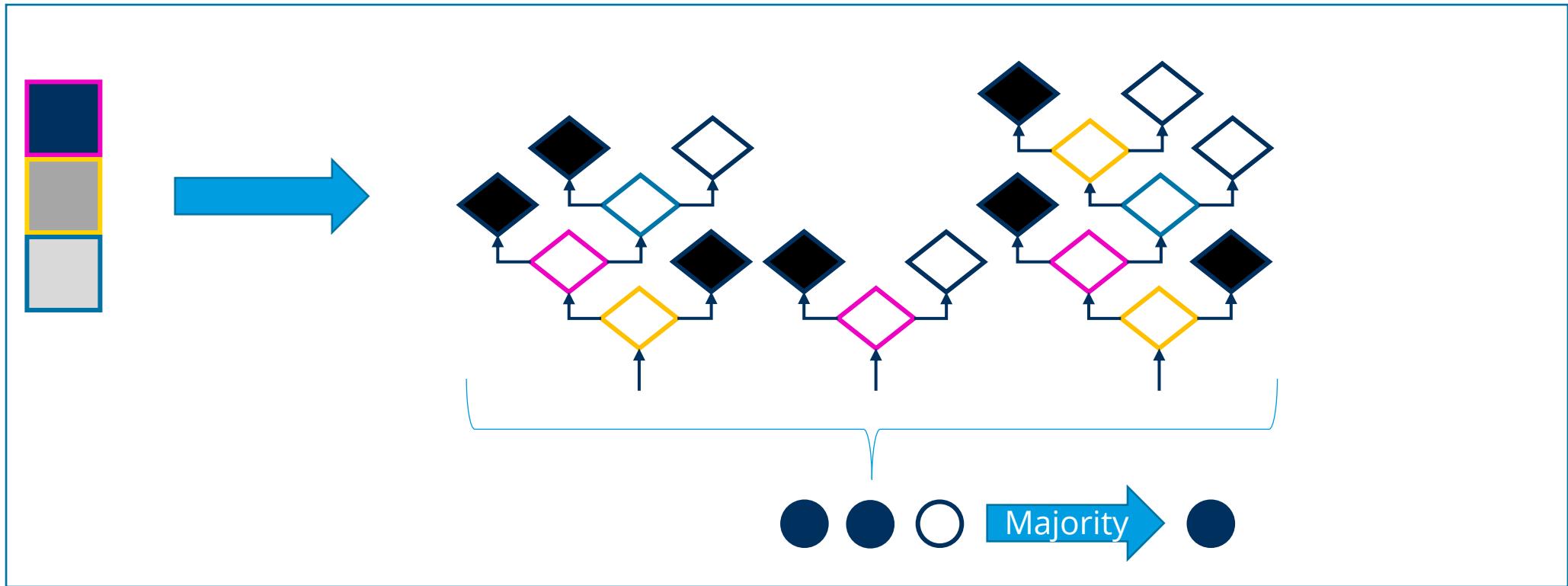
By training many decision trees, errors are equilibrated



Random Forest Pixel Classifiers

Combination of individual tree decisions by voting or max / mean

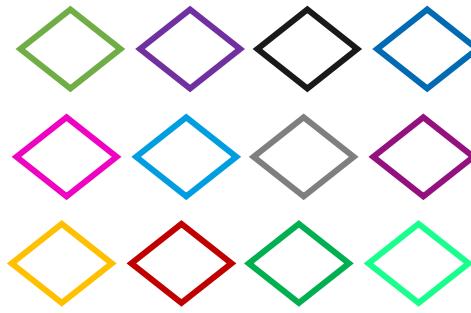
Prediction



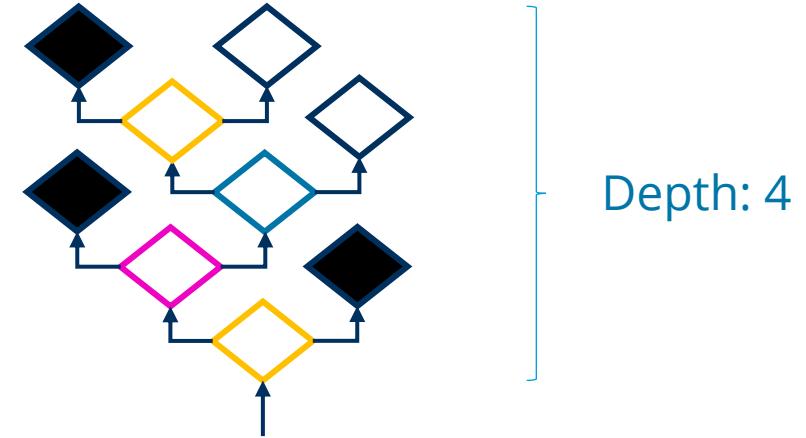
Random Forest Pixel Classifiers

Typical numbers for pixel classifiers in microscopy

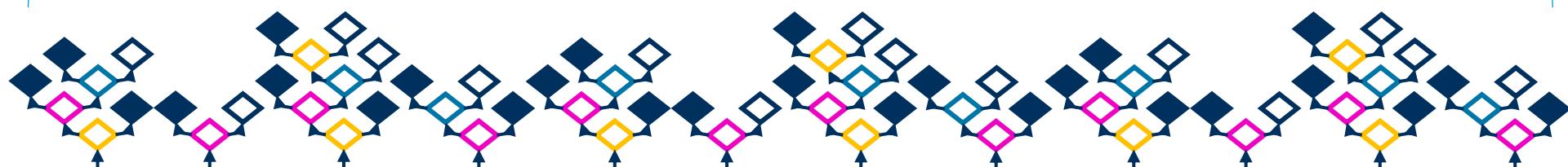
Available features:



- Gaussian blur image
- DoG image
- LoG image
- Hessian
-



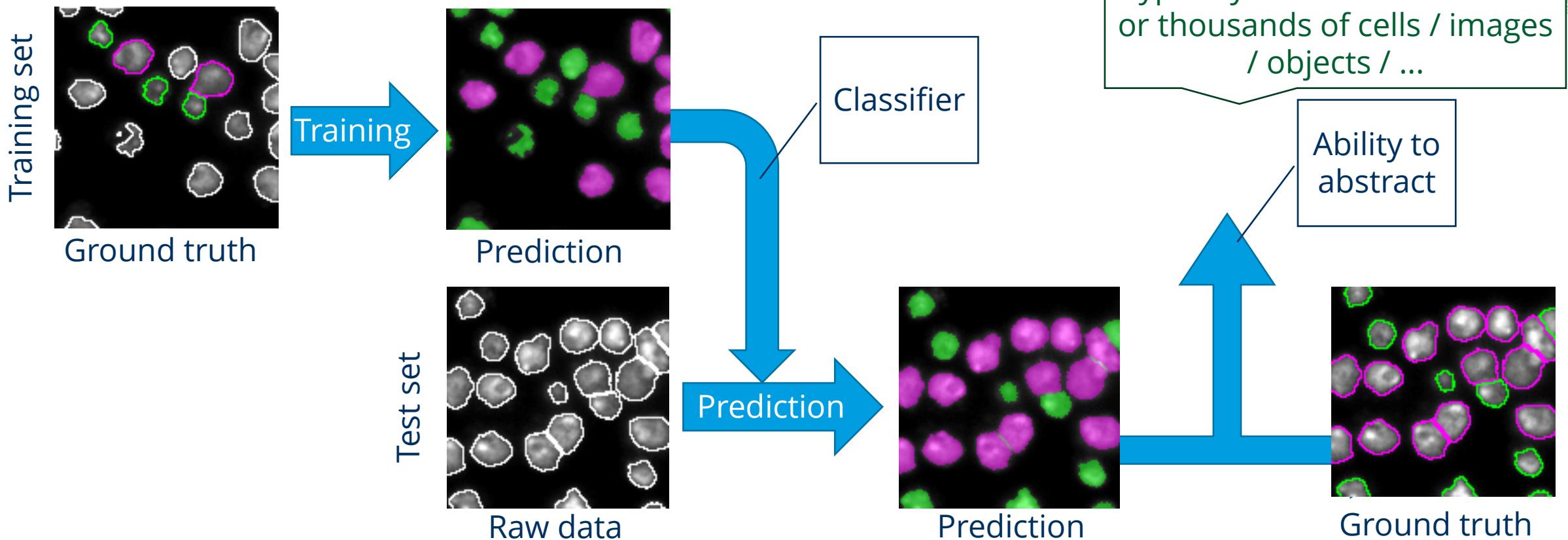
Number of trees: > 100



Model validation

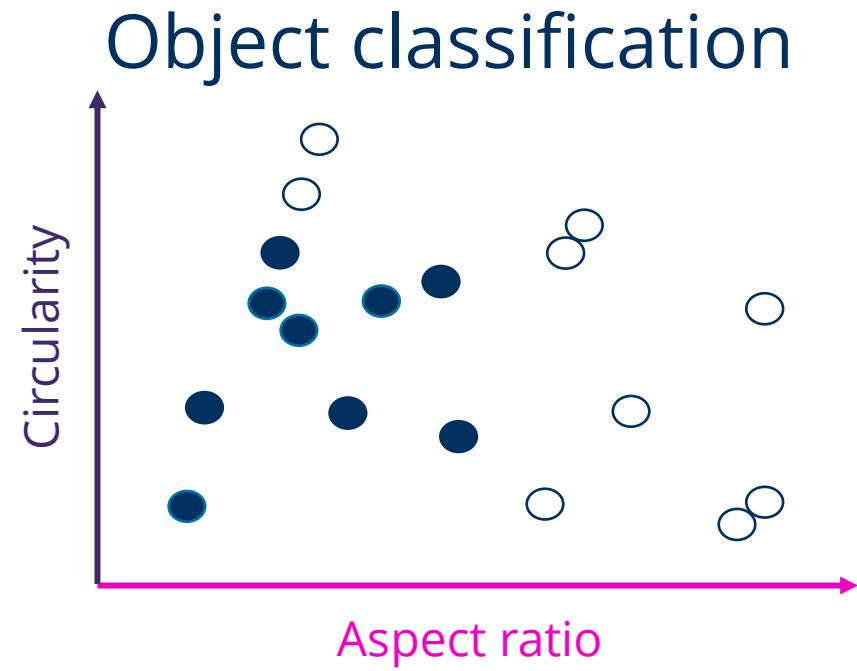
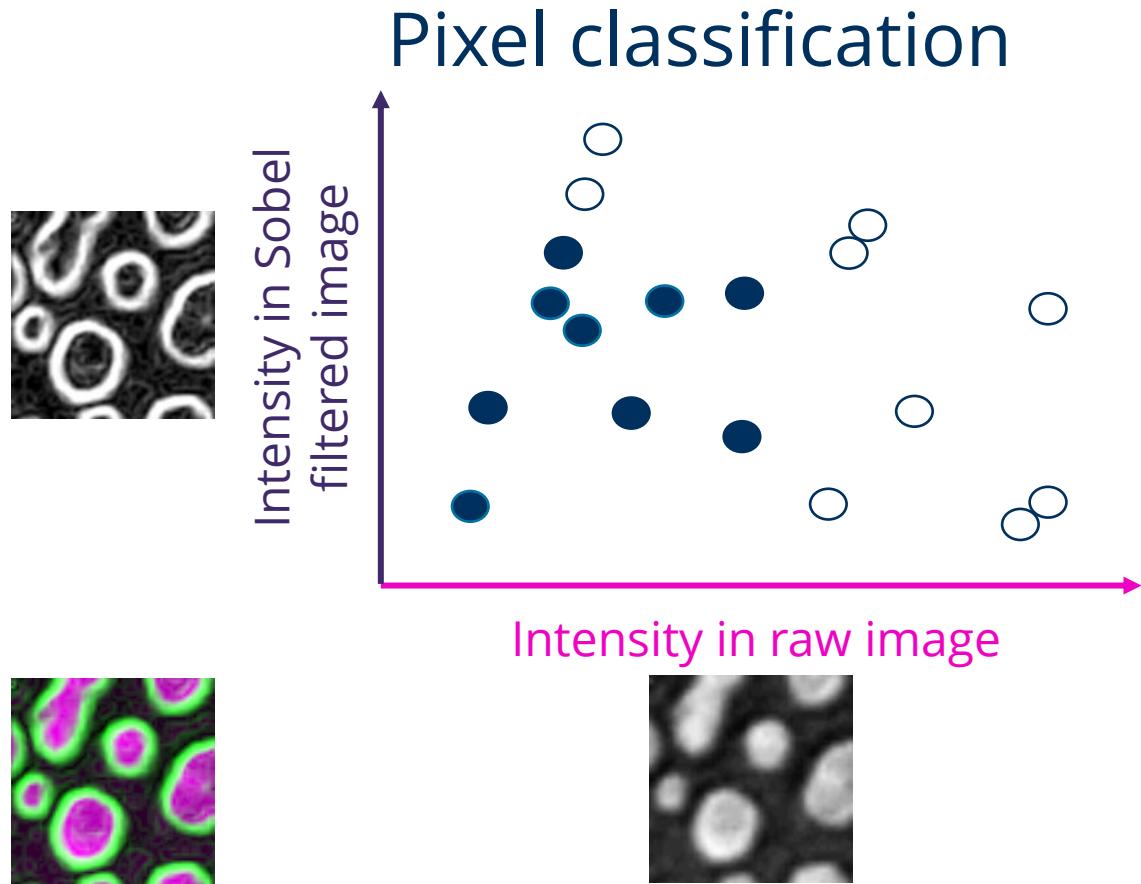
In order to assess model quality, we split the ground truth into two sets

- Training set (50%-90% of the available data)
- Test set (10%-50% of the available data)



Object classification

What if we exchange pixel features with object features?



- The algorithms work the same using with different features

Supervised and Unsupervised Machine Learning for Bio-image Analysis

Robert Haase

Reusing materials from Johannes Soltwedel, Till Korten, Johannes Müller, Laura Žigutytė (TU Dresden), Ryan Savill (MPI-CBG), Matthias Täschner (ScaDS.AI/Uni Leipzig) and the Scikit-learn community.

Using
Python

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und Forschung



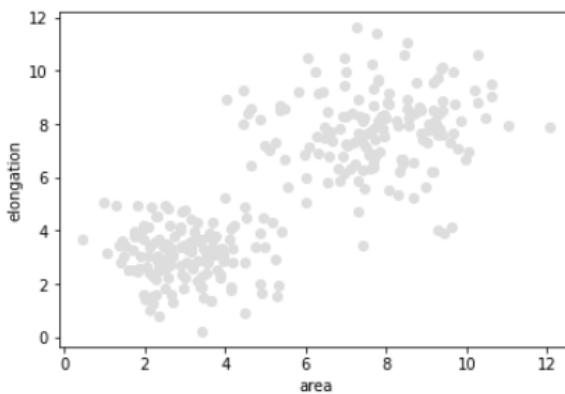
Diese Maßnahme wird gefördert durch die Bundesregierung aufgrund eines Beschlusses des Deutschen Bundestages.
Diese Maßnahme wird mitfinanziert durch Steuermittel auf der Grundlage des von den Abgeordneten des Sächsischen Landtags beschlossenen Haushaltes.

Tabular object classification

Classify objects starting from feature vectors (table columns)

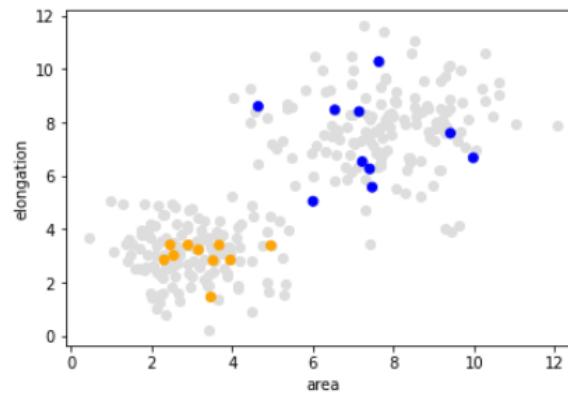
Raw data

| | area | elongation |
|----|----------|------------|
| 0 | 3.950088 | 2.848643 |
| 1 | 4.955912 | 3.390093 |
| 2 | 7.469852 | 5.575289 |
| 3 | 2.544467 | 3.017479 |
| 4 | 3.465662 | 1.463756 |
| 5 | 3.156507 | 3.232181 |
| 6 | 9.978705 | 6.676372 |
| 7 | 6.001683 | 5.047063 |
| 8 | 2.457139 | 3.416050 |
| 9 | 3.672295 | 3.407462 |
| 10 | 9.413702 | 7.598608 |



“Ground truth” annotation

annotation = [1, 1, 2, 1, 1, 1,

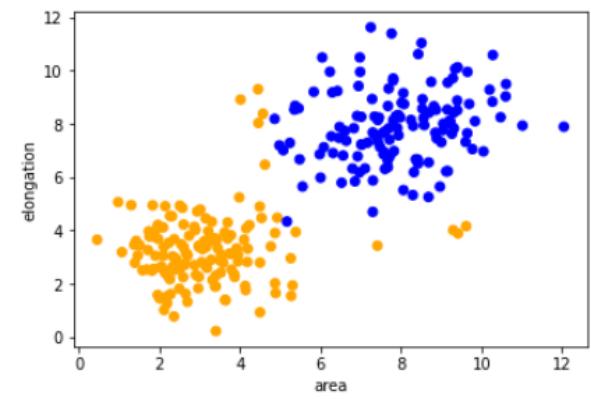


Classifier training

```
classifier = RandomForestClassifier()  
classifier.fit(train_data, train_annotation)
```

Classifier prediction

```
result = classifier.predict(validation_data)
```



Interactive pixel classification

Prepare an empty layer for annotations and keep a reference

```
labels = viewer.add_labels(  
    np.zeros(image.shape).astype(int))
```

Read annotations

```
manual_annotations = labels.data
```

```
from skimage.io import imshow  
  
imshow(manual_annotations,  
       vmin=0, vmax=2)
```

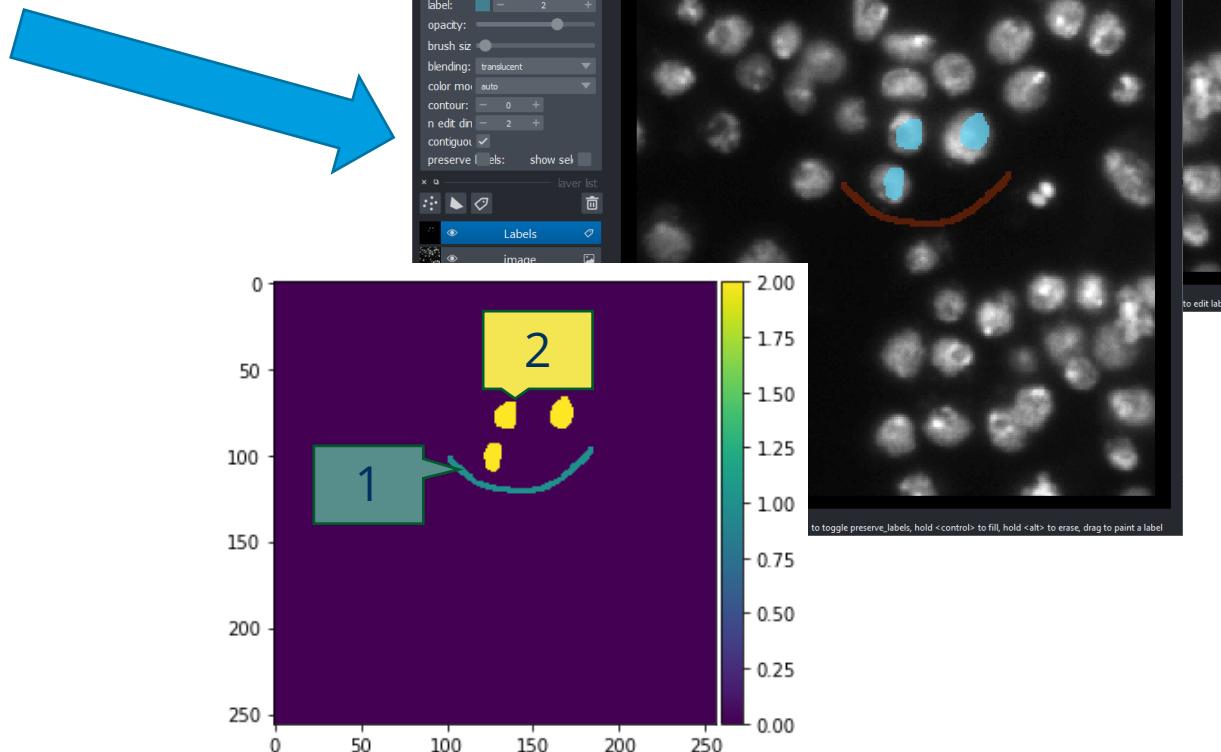
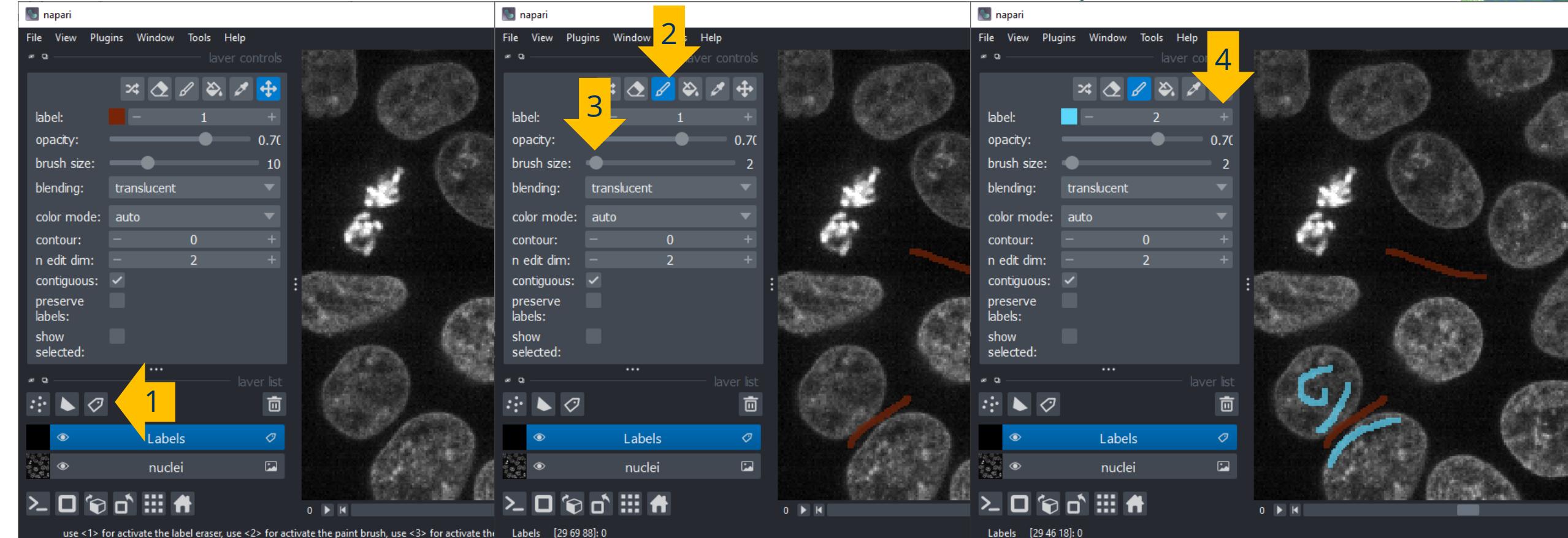


Image data source: BBB038v1, available from the Broad BioImage Benchmark Collection (Caicedo et al., Nature Methods, 2019).

Napari – common workflows

Pixel / object annotation drawing

- [1: Create empty labels layer]
- 2: Select paint brush tool
- 3: Decrease brush size
- 4: Increase label



Interactive pixel classification

Pixel classification using scikit-learn

- Expects one-dimensional arrays for features and ground truth

```
# train classifier  
  
from sklearn.ensemble import RandomForestClassifier  
  
classifier = RandomForestClassifier(max_depth=2, random_state=0)  
  
classifier.fit(X, y)
```

Image data

Ground truth /
annotation

y_* = classifier.predict(X)

prediction

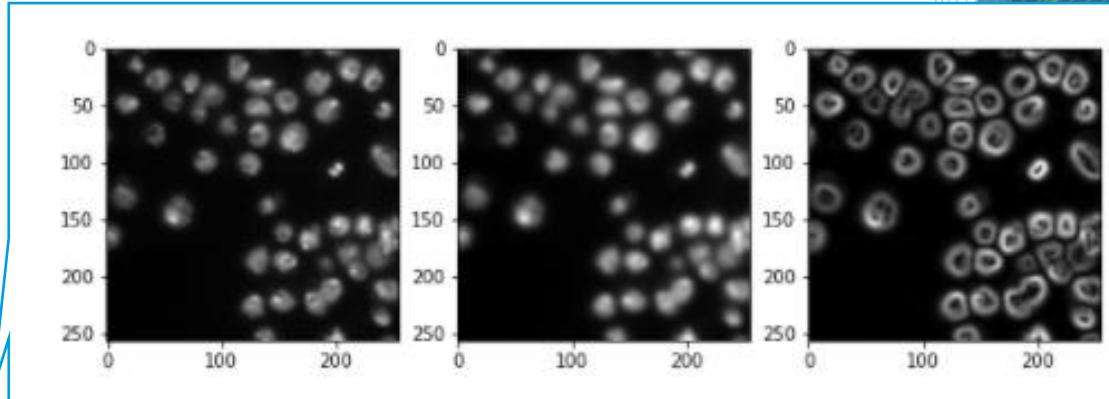
Interactive pixel classification

Pixel classification using scikit-learn

- Expects one-dimensional arrays for features and ground truth

```
# for training, we need to generate features
feature_stack = generate_feature_stack(image)
X, y = format_data(feature_stack, manual_annotations)

# train classifier
from sklearn.ensemble import RandomForestClassifier
classifier = RandomForestClassifier(max_depth=2, random_state=0)
classifier.fit(X, y)
```



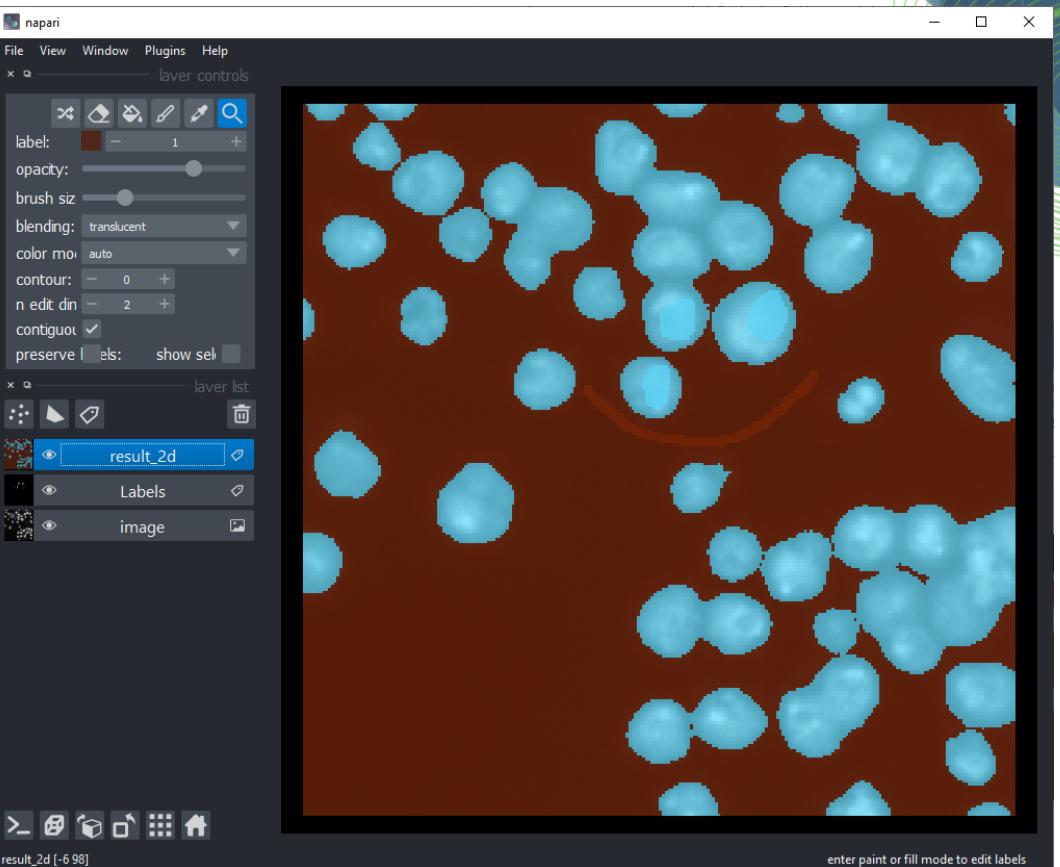
Interactive pixel classification

Pixel classification using scikit-learn

- Expects one-dimensional arrays for features and ground truth

```
# process the whole image and show result  
  
result_1d = classifier.predict(feature_stack.T)  
  
result_2d = result_1d.reshape(image.shape)  
  
viewer.add_labels(result_2d)
```

Convert 1D
result back to 2D



Interactive pixel classification

Jupyter notebooks and napari side-by-side

scikit_learn_random_forest_pixel_ x +
localhost:8889/notebooks/machine_learning/scikit_learn_random_forest_pixel_clas...
jupyter scikit_learn_random_forest_pixel_classifier (autosaved)
File Edit View Insert Cell Kernel Widgets Help Trusted Python
Interactive segmentation
We can also use napari to annotate some regions as negative (label = 1) and positive (label = 2).
In [30]: `import napari

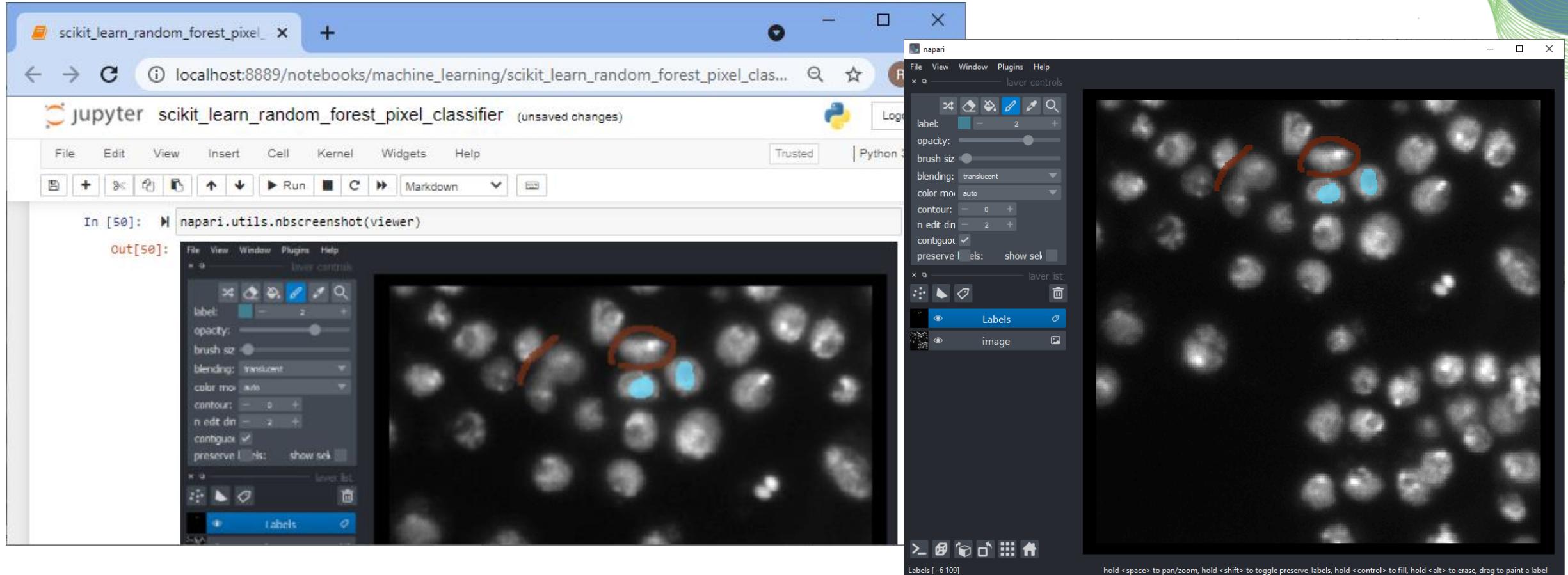
start napari
viewer = napari.Viewer()

add image
viewer.add_image(image)

add an empty Labels layer and keep it in a variable
labels = viewer.add_labels(np.zeros(image.shape).astype(int))`
Go ahead after annotating at least two regions with labels 1 and 2.
Take a screenshot of the annotation:
Labels [-6 -7] enter paint or fill mode to edit labels

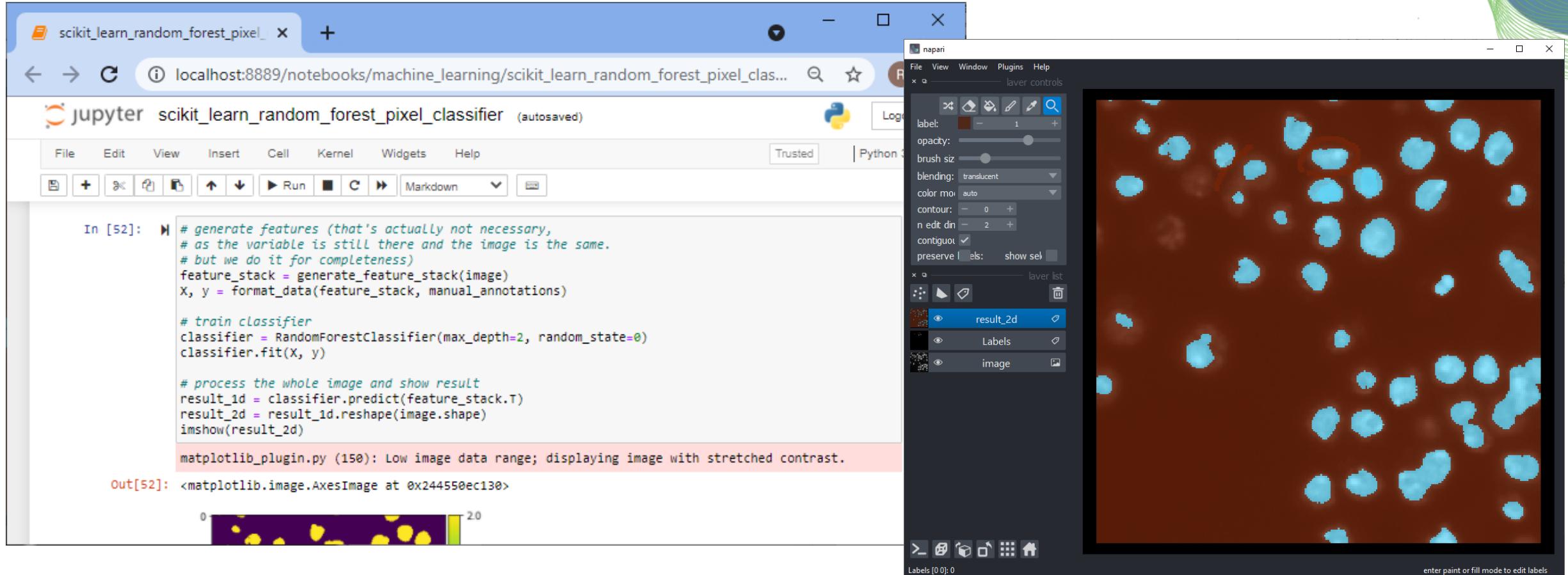
Interactive pixel classification

Jupyter notebooks and napari side-by-side



Interactive pixel classification

Jupyter notebooks and napari side-by-side



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GPU-accelerated

Using
Python

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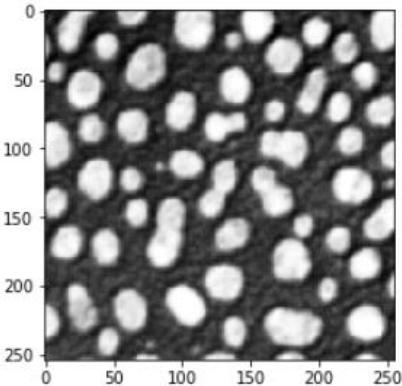
Bundesministerium
für Bildung
und Forschung



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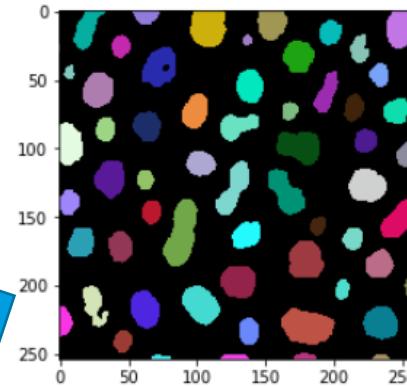
Accelerated pixel and object classification

APOC is a python library that makes use of OpenCL-compatible Graphics Cards to accelerate pixel and object classification

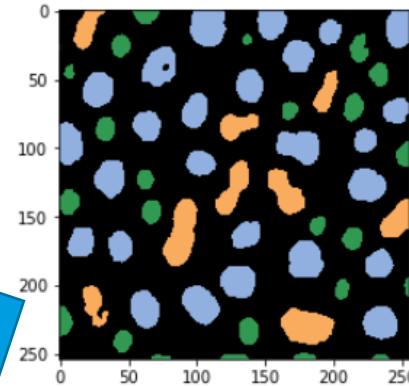


Raw image

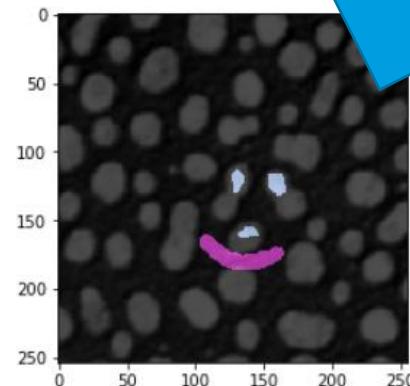
Object segmentation



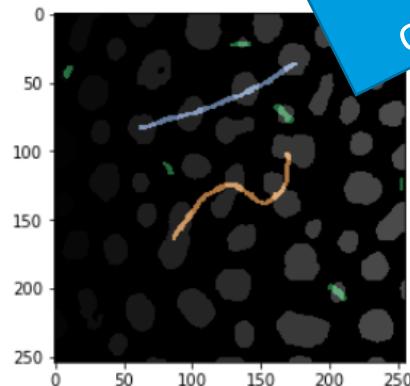
Object label image



Class label image



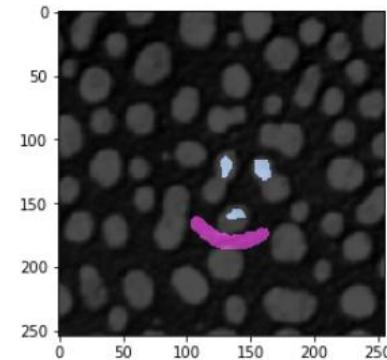
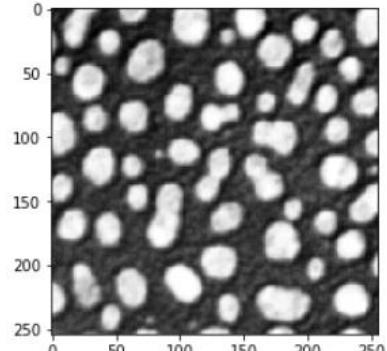
Pixel annotation



Object annotation

Object segmentation

Pixel classification + connected component labeling



```
# define features
features = "gaussian_blur=1 gaussian_blur=5 sobel_of_gaussian_blur=1"

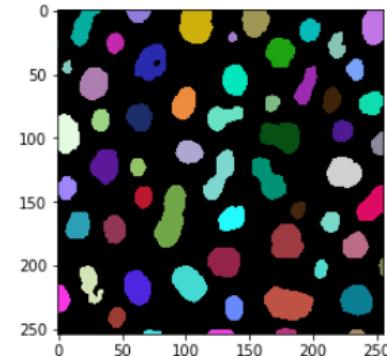
# this is where the model will be saved
cl_filename = 'my_object_segmenter.cl'

# delete classifier in case the file exists already
apoc.erase_classifier(cl_filename)

# train classifier
clf = apoc.ObjectSegmenter(opencl_filename=cl_filename, positive_class_identifier=2)
clf.train(features, manual_annotations, image)

segmentation_result = clf.predict(features=features, image=image)
cle.imshow(segmentation_result, labels=True)
```

Object segmentation



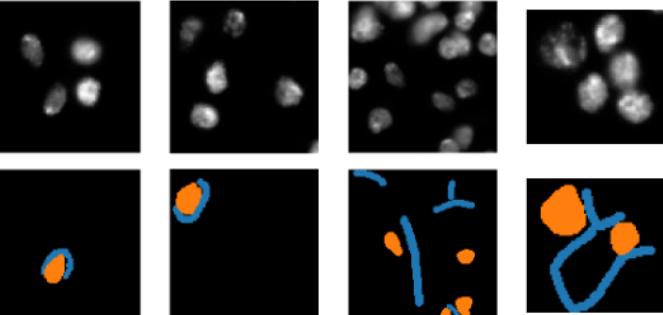
Training on folders of annotated images

```
[2]: image_folder = "data/BBBC007/images/"
masks_folder = "data/BBBC007/masks/"

[3]: file_list = os.listdir(image_folder)

# show all images
fig, axs = plt.subplots(1, 4, figsize=(15,15))
for i, filename in enumerate(file_list):
    image = imread(image_folder + filename)
    stackview.imshow(image, plot=axs[i])
plt.show()

# show corresponding label images
fig, axs = plt.subplots(1, 4, figsize=(15,15))
for i, filename in enumerate(file_list):
    masks = imread(masks_folder + filename)
    stackview.imshow(masks, plot=axs[i], labels=True)
plt.show()
```



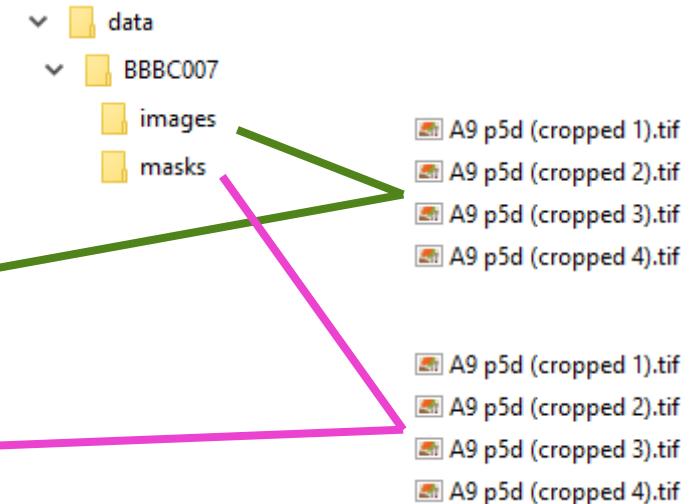
Training

If the folders are setup properly, we can pass the folders to the training.

```
[4]: # setup classifier and where it should be saved
segmenter = apoc.ObjectSegmenter(opencl_filename="data/object_segmenter_trained_on_f

# setup feature set used for training
features = apoc.PredefinedFeatureSet.object_size_1_to_5_px.value

# train classifier on folders
apoc.train_classifier_from_image_folders(
    segmenter,
    features,
    image = image_folder,
    ground_truth = masks_folder)
```



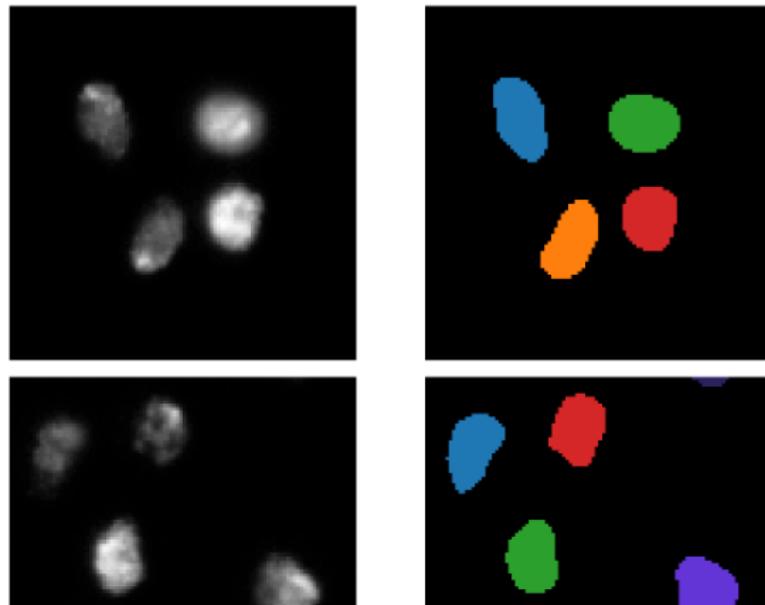
Prediction

After the training, we can apply the classifier to all images in the image folder. The following line reloads the classifier from disk. In that way we can ensure that it was stored correctly.

```
[5]: segmenter = apoc.ObjectSegmenter(opencl_filename="data/object_segmenter_trained_on_f
[6]: # show all images
for i, filename in enumerate(file_list):
    fig, axs = plt.subplots(1, 2, figsize=(15,15))

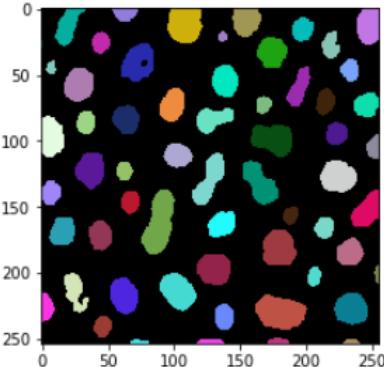
    image = imread(image_folder + filename)
    stackview.imshow(image, plot=axs[0])

    labels = segmenter.predict(image)
    stackview.imshow(labels, plot=axs[1], labels=True)
```

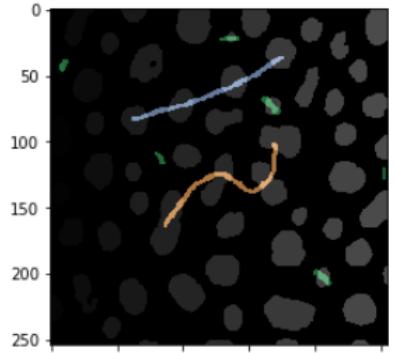


Object classification

Feature extraction + tabular classification



Object label image



Object annotation

```
# for the classification we define size and shape as criteria
features = 'area mean_max_distance_to_centroid_ratio'

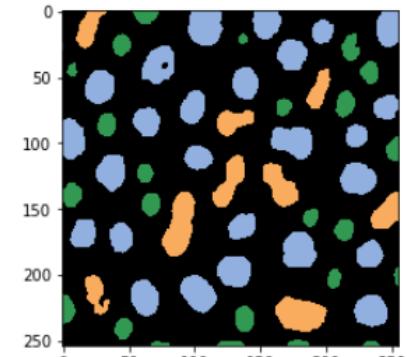
# This is where the model will be saved
cl_filename_object_classifier = "my_object_classifier.cl"

# delete classifier in case the file exists already
apoc.erase_classifier(cl_filename_object_classifier)

# train the classifier
classifier = apoc.ObjectClassifier(cl_filename_object_classifier)
classifier.train(features, segmentation_result, annotation, image)

# determine object classification
classification_result = classifier.predict(segmentation_result, image)
cle.imshow(classification_result, labels=True)
```

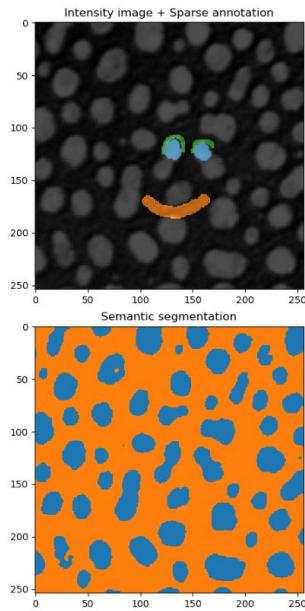
Object classification



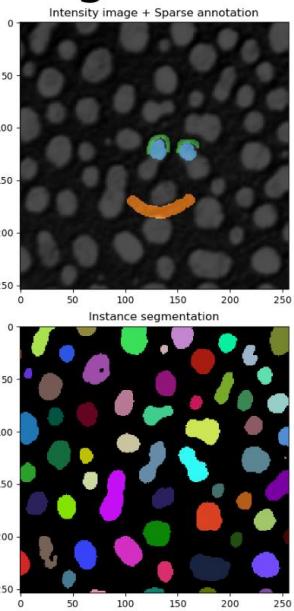
Class label image

Other classification / regression tasks

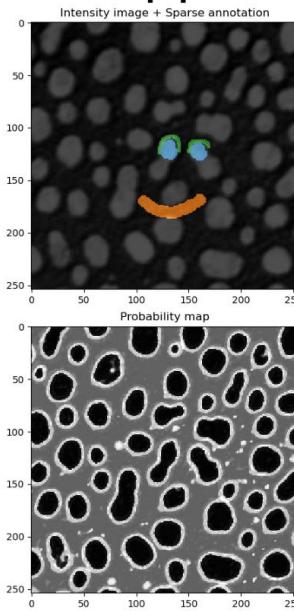
Pixel-
Classifier



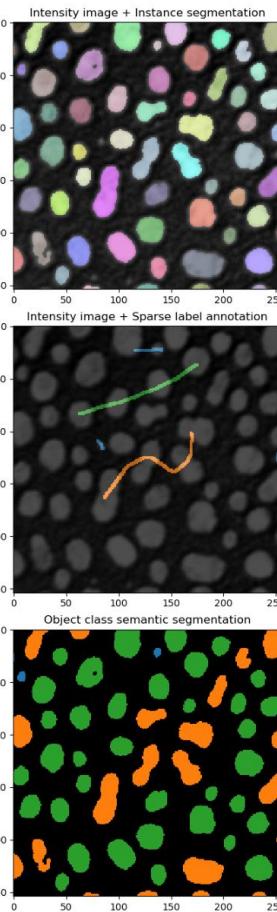
Object-
Segmenter



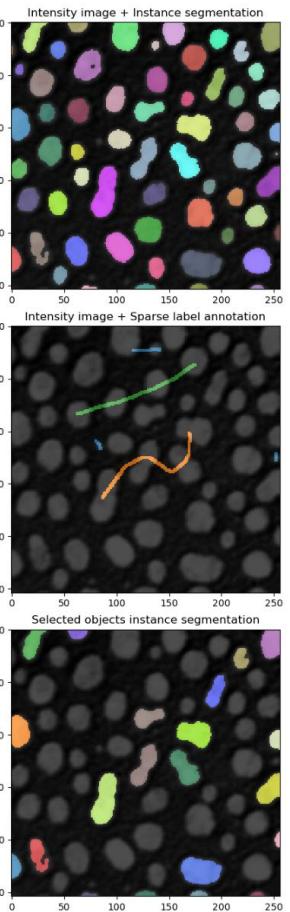
Probability-
Mapper



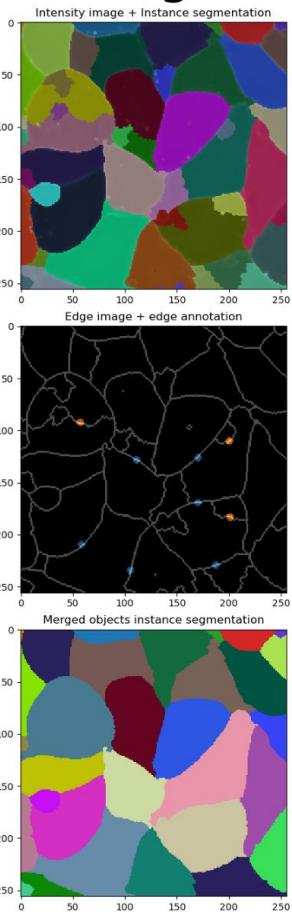
Object-
Mapper



Object-
Classifier

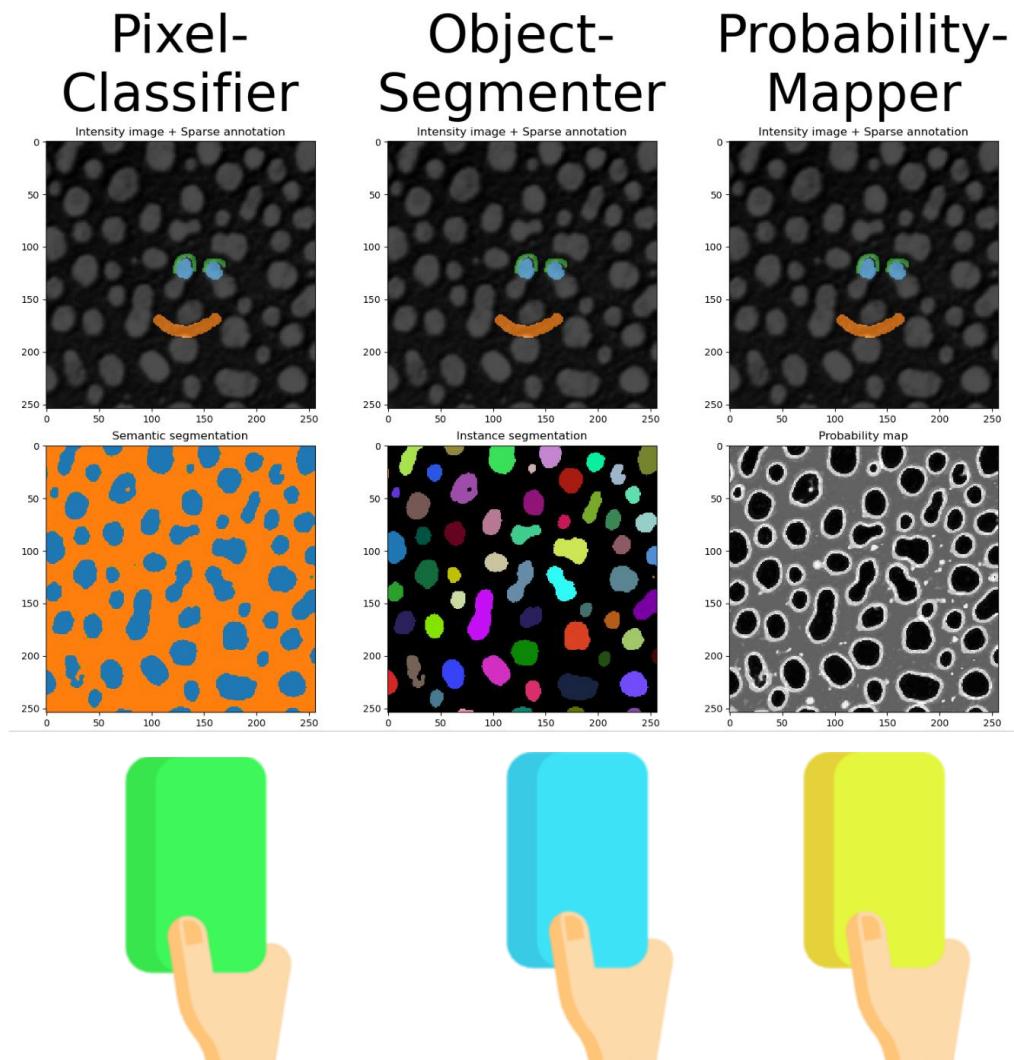


Object-
Selector



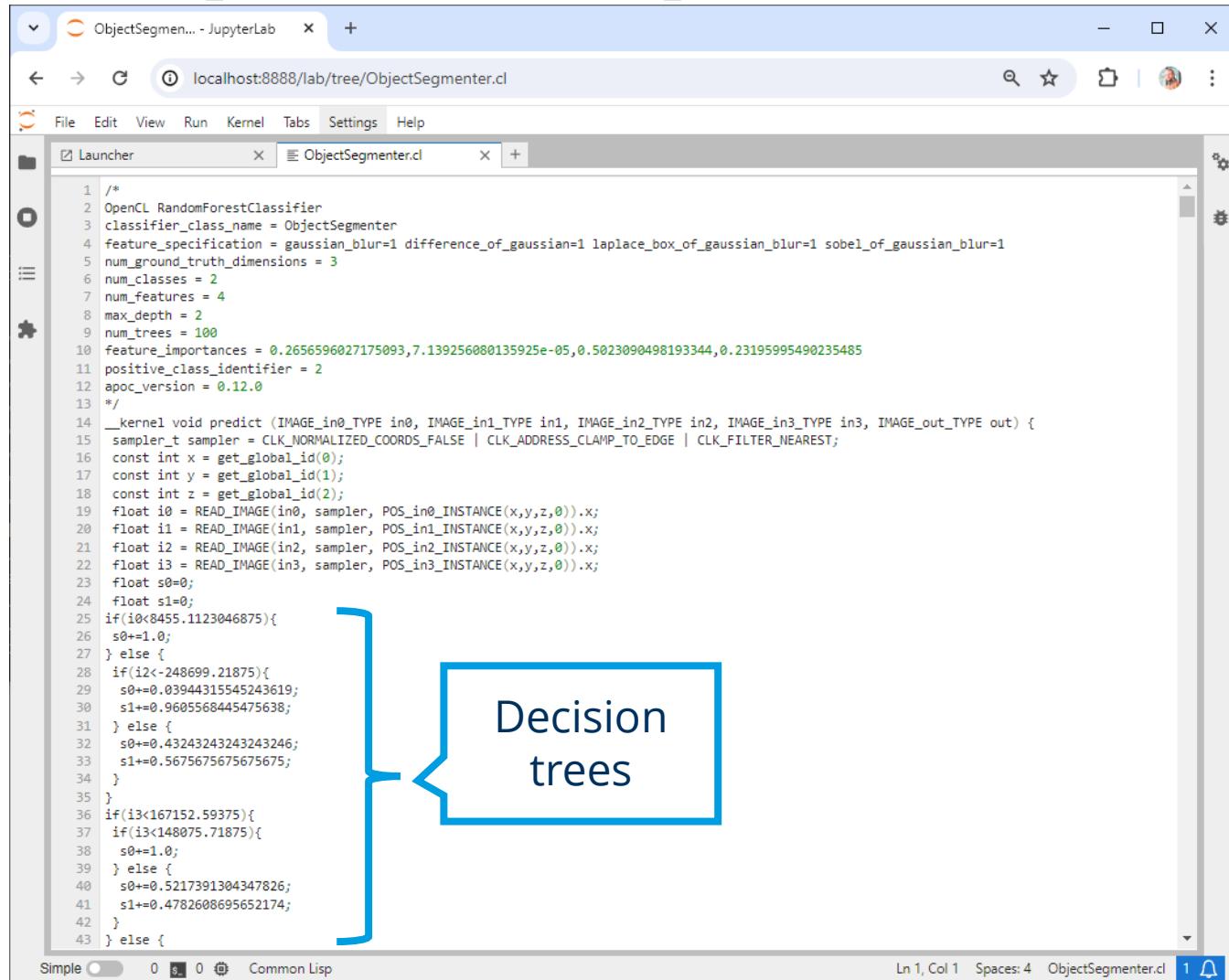
Quiz: Classification versus Regression

Which of these three solves a regression task?



Under the hood: clesperanto / OpenCL

classifier.cl files
can be *read*



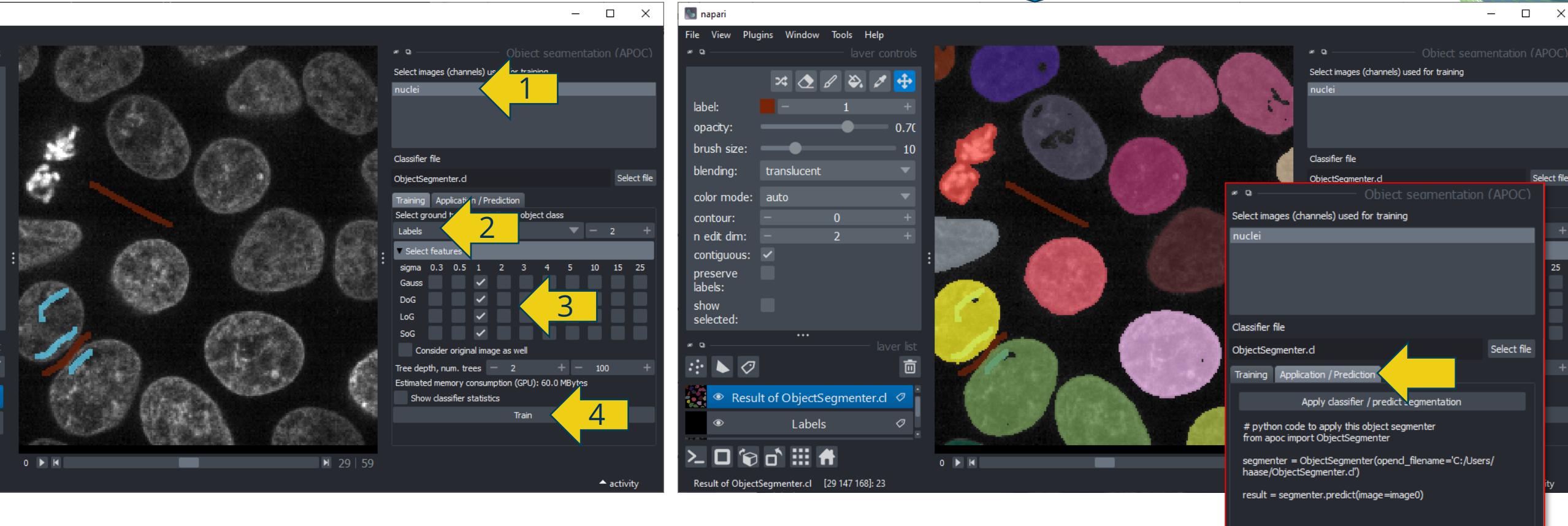
```
/*
OpenCL RandomForestClassifier
classifier_class_name = ObjectSegmenter
feature_specification = gaussian_blur=1 difference_of_gaussian=1 laplace_box_of_gaussian_blur=1 sobel_of_gaussian_blur=1
num_ground_truth_dimensions = 3
num_classes = 2
num_features = 4
max_depth = 2
num_trees = 100
feature_importances = 0.2656596027175093, 7.139256080135925e-05, 0.5023090498193344, 0.23195995490235485
positive_class_identifier = 2
apoc_version = 0.12.0
*/
__kernel void predict (IMAGE_in0_TYPE in0, IMAGE_in1_TYPE in1, IMAGE_in2_TYPE in2, IMAGE_in3_TYPE in3, IMAGE_out_TYPE out) {
    sampler_t sampler = CLK_NORMALIZED_COORDS_FALSE | CLK_ADDRESS_CLAMP_TO_EDGE | CLK_FILTER_NEAREST;
    const int x = get_global_id(0);
    const int y = get_global_id(1);
    const int z = get_global_id(2);
    float i0 = READ_IMAGE(in0, sampler, POS_in0_INSTANCE(x,y,z,0)).x;
    float i1 = READ_IMAGE(in1, sampler, POS_in1_INSTANCE(x,y,z,0)).x;
    float i2 = READ_IMAGE(in2, sampler, POS_in2_INSTANCE(x,y,z,0)).x;
    float i3 = READ_IMAGE(in3, sampler, POS_in3_INSTANCE(x,y,z,0)).x;
    float s0=0;
    float s1=0;
    if(i0>8455.1123046875){
        s0+=1.0;
    } else {
        if(i2<-248699.21875){
            s0+=0.03944315545243619;
            s1+=0.9605568445475638;
        } else {
            s0+=0.43243243243243246;
            s1+=0.5675675675675675;
        }
    }
    if(i3<167152.59375){
        if(i3<148075.71875){
            s0+=1.0;
        } else {
            s0+=0.5217391304347826;
            s1+=0.4782608695652174;
        }
    } else {

```

Decision trees

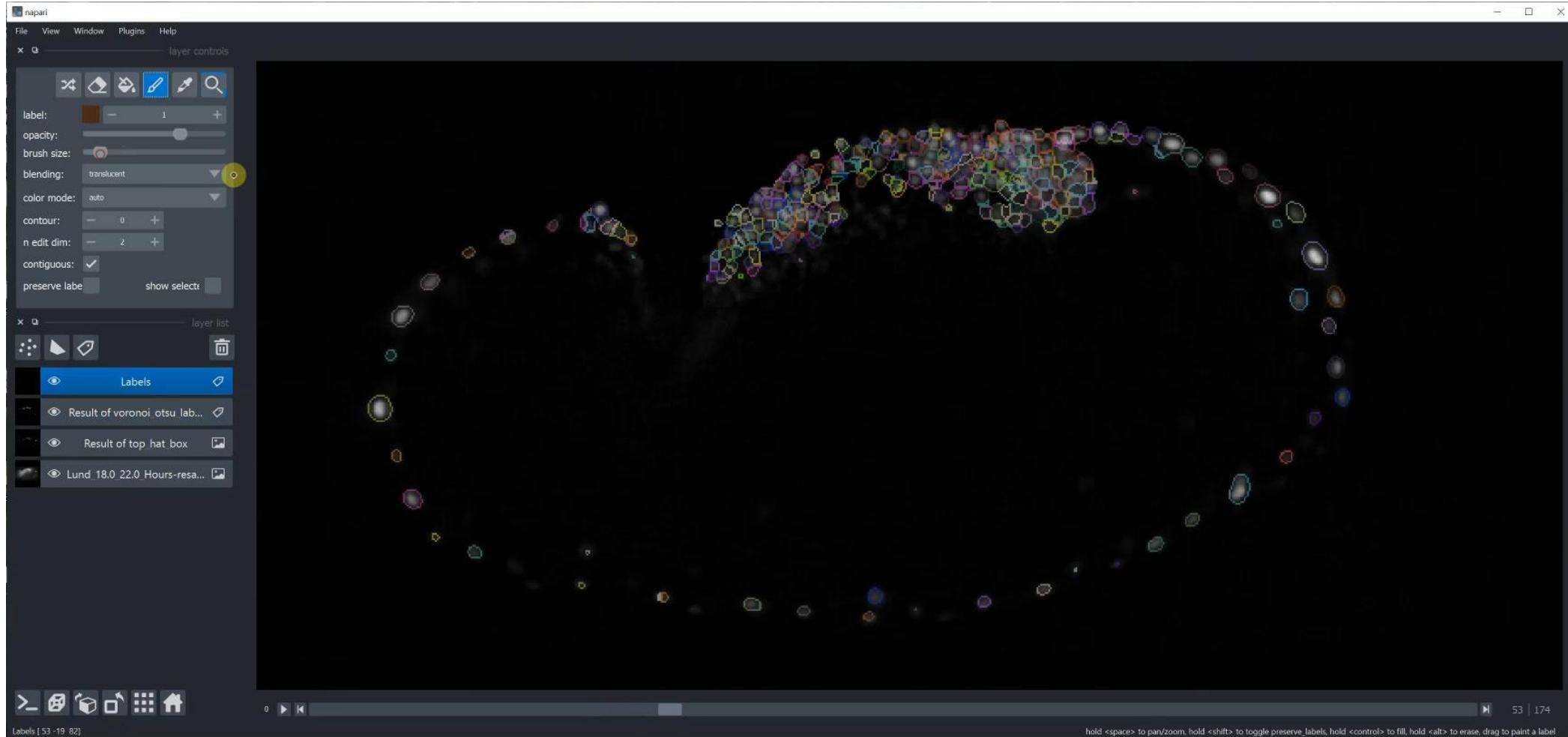
Graphical user interface: Object segmentation

- 1: Select image[s]
- 2: Select ground truth annotation
- [3: Select features]
- 4: Train / predict



Graphical user interface: Object classification

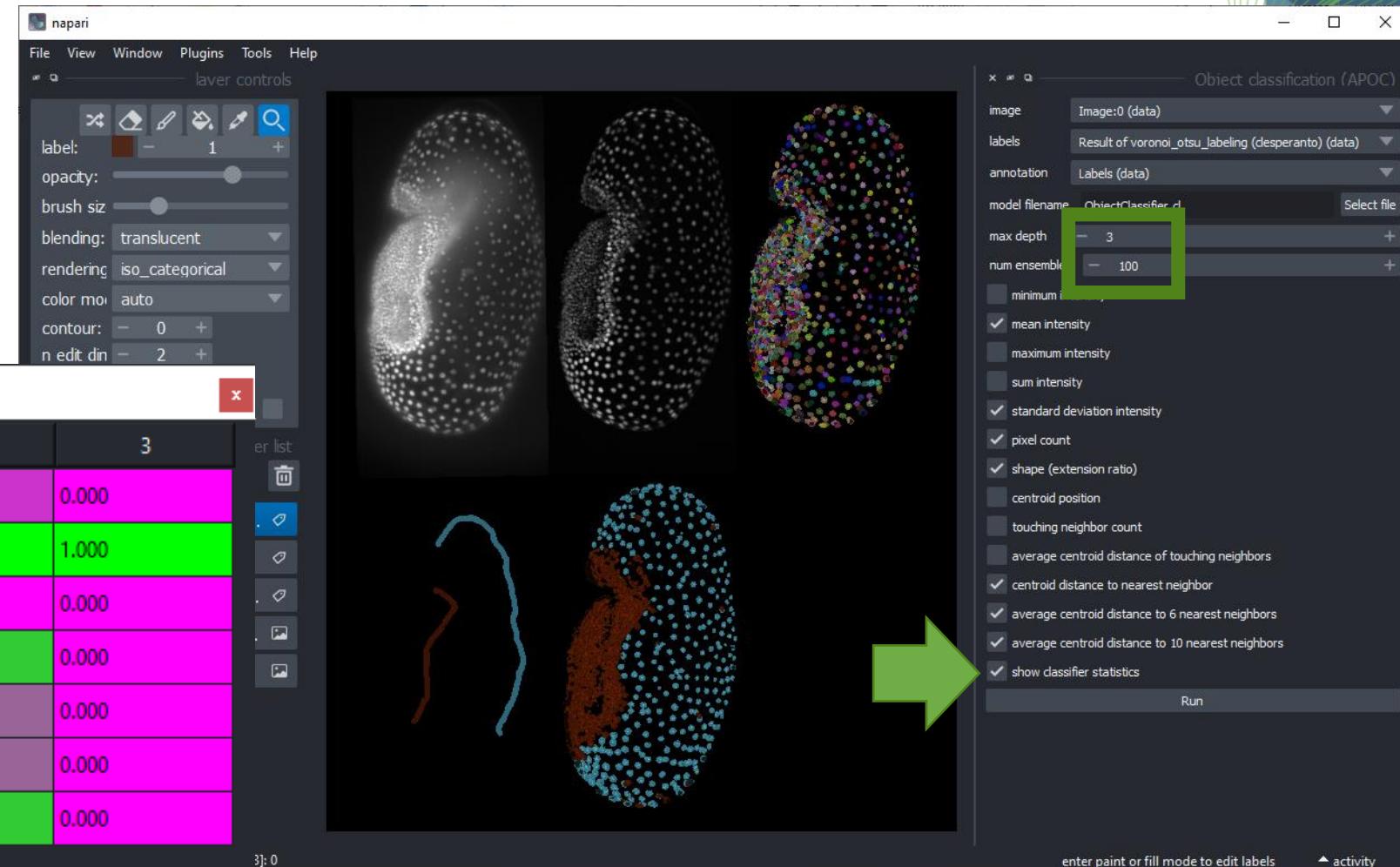
Annotation / classification of segmented objects



Graphical user interface: Object classification

Inspect how the random forest classifier makes decisions

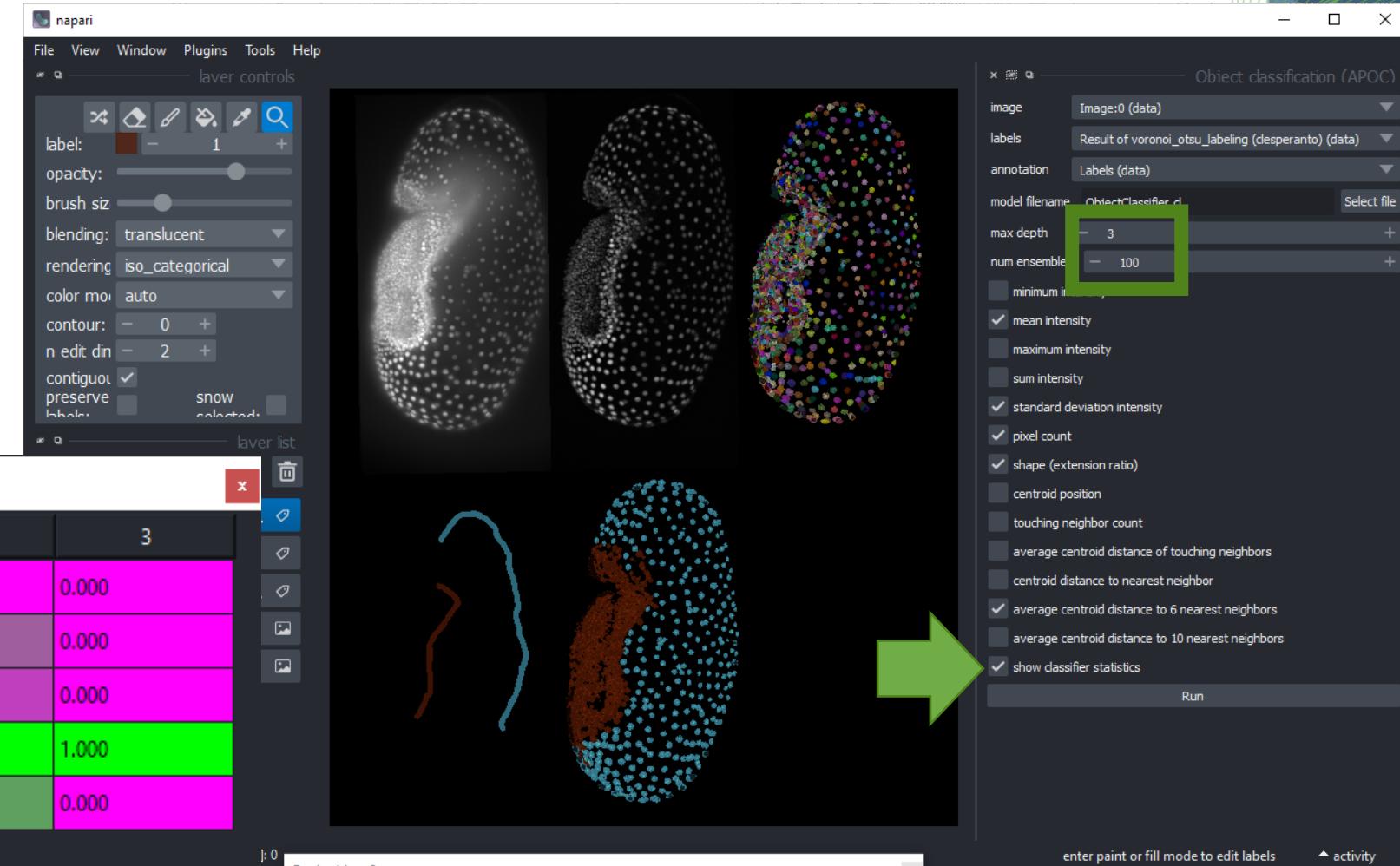
Note: Beware of correlated parameters!



Graphical user interface: Object classification

Inspect how the random forest classifier makes decisions

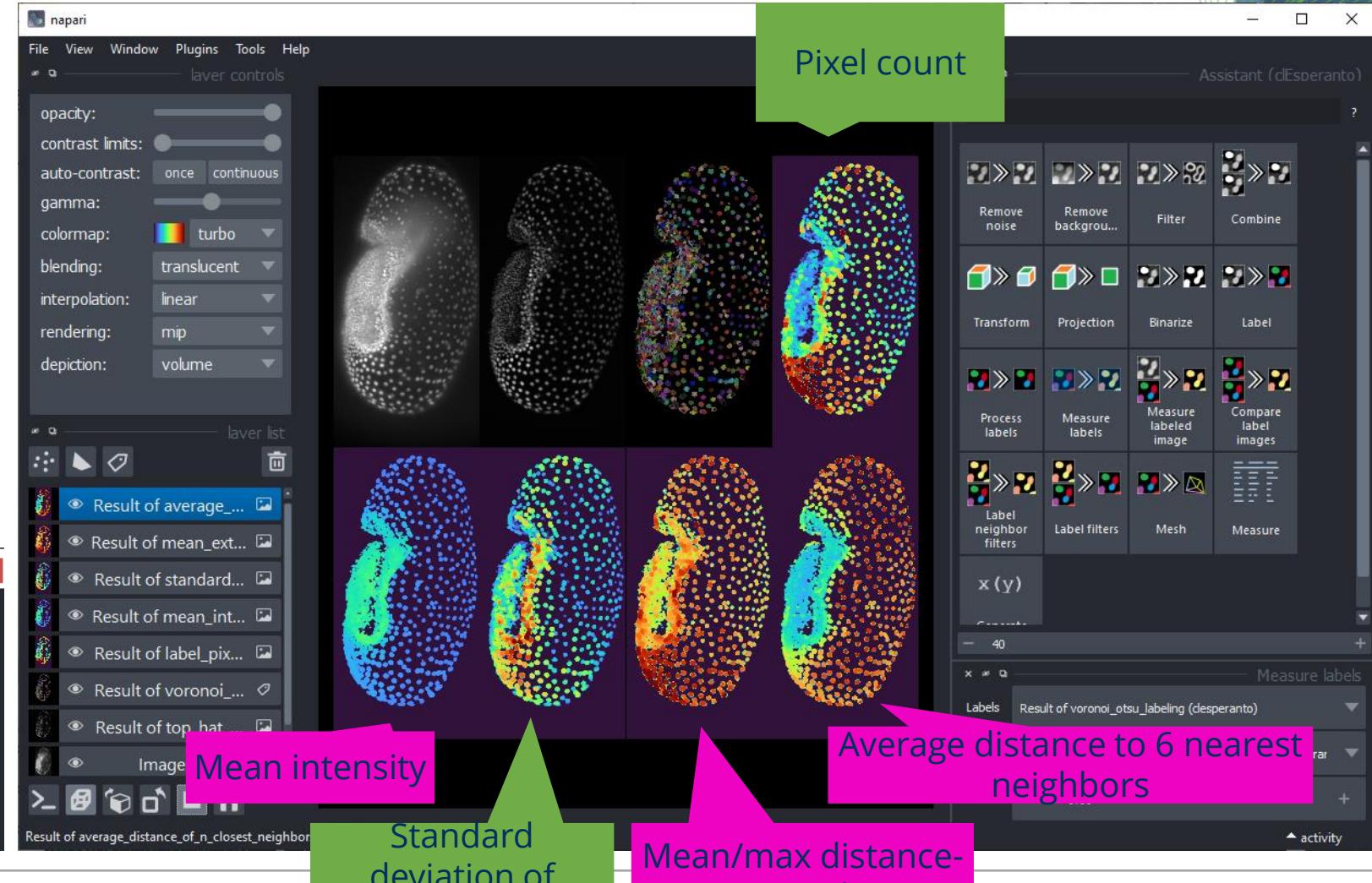
Note: Beware of correlated parameters!



Graphical user interface: Object classification

Inspect how the random forest classifier makes decisions

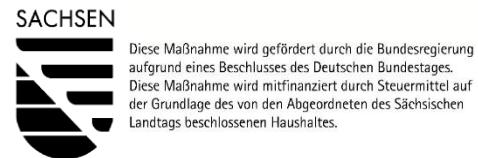
Note: Beware of correlated parameters!



Supervised and Unsupervised Machine Learning for Bio-image Analysis

Robert Haase

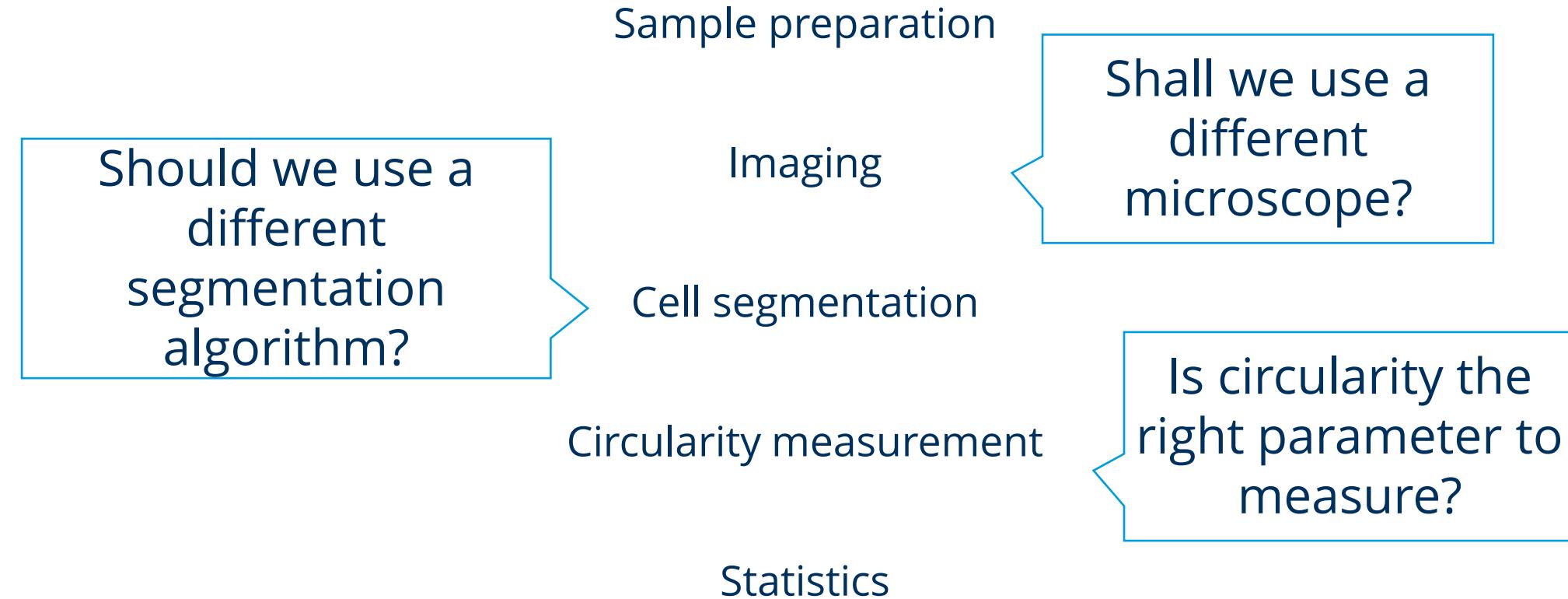
Reusing materials from Johannes Soltwedel, Till Korten, Johannes Müller, Laura Žigutytė (TU Dresden), Ryan Savill (MPI-CBG), Matthias Täschner (ScaDS.AI/Uni Leipzig) and the Scikit-learn community.



Hypothesis-driven quantitative biology

Hypothesis: Cell shape can be influenced by modifying X.

Null-Hypothesis: Circularity of modified cells is similar to cells in the control group.



Hypothesis generating quantitative biology

Hypothesis: Cell shape can be influenced by modifying X.

Question: Which image-derived parameter is influenced when modifying X?

Sample preparation

Imaging

Cell segmentation algorithm A, algorithm B, algorithm C

Measurement of circularity, solidity, elongation, extend, texture, intensity, topology ...

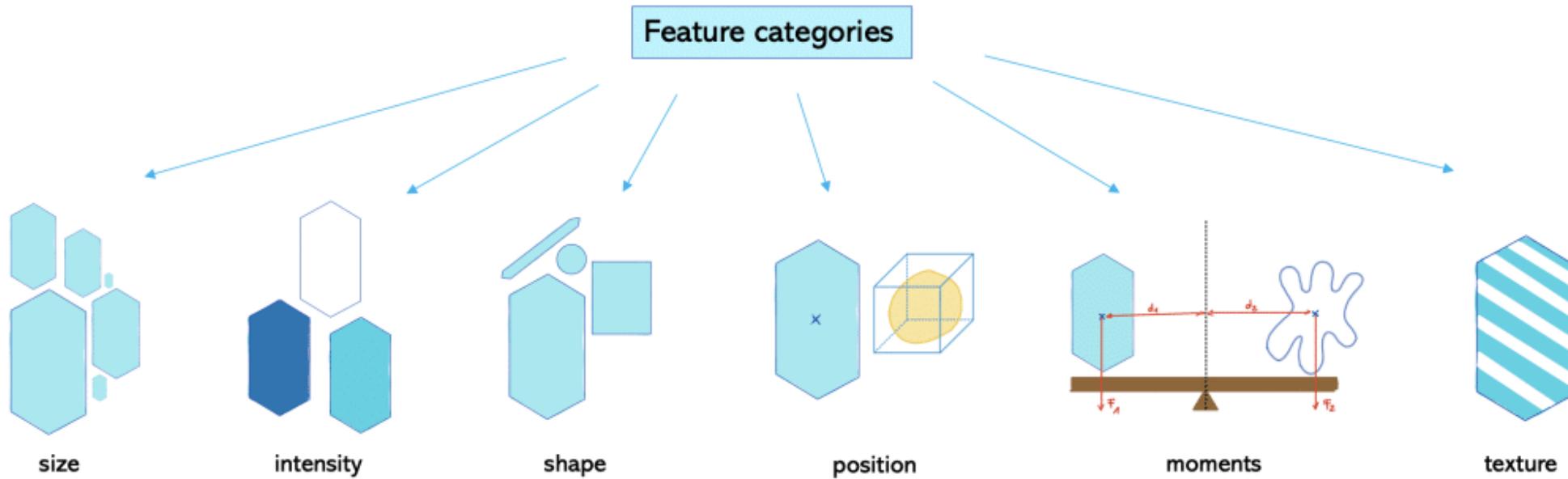
Statistics

Which segmentation algorithms allow measurements that show a relationship with X?

Why?

Which parameter shows any relationship with X?

Feature selection



Which of these features reflect the phenotype we are perceiving?

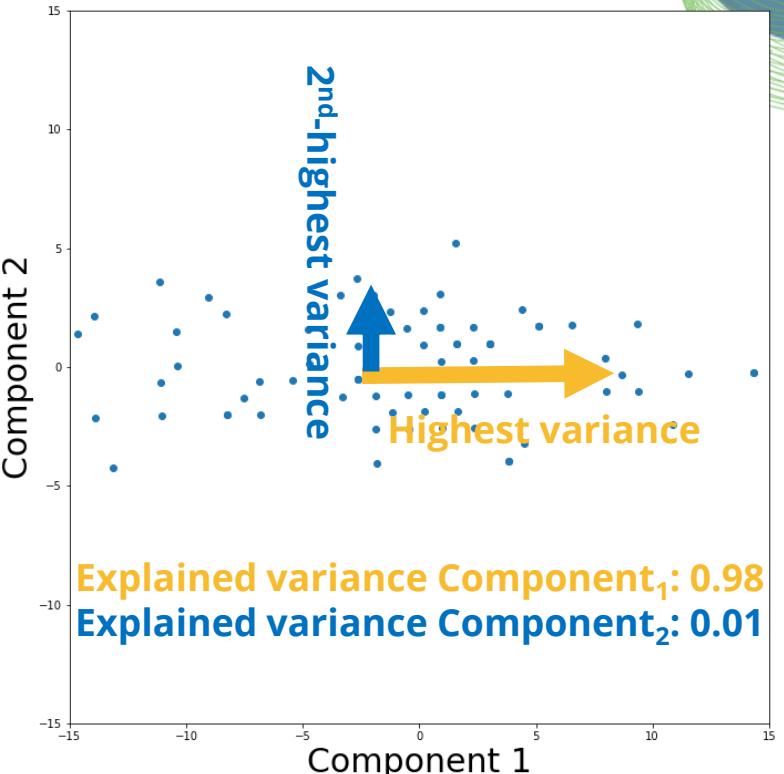
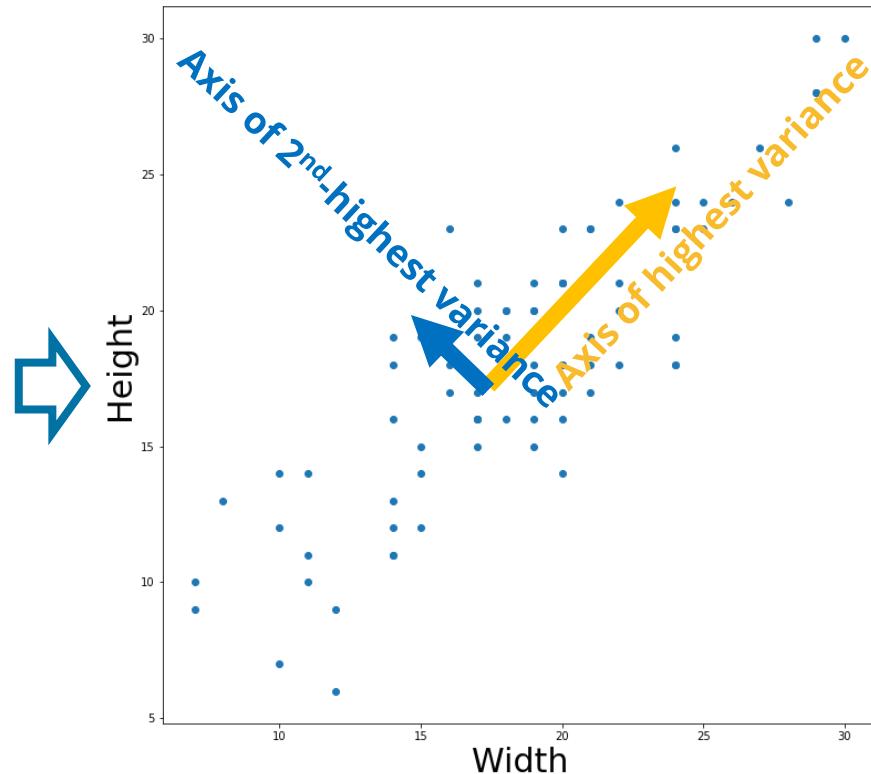
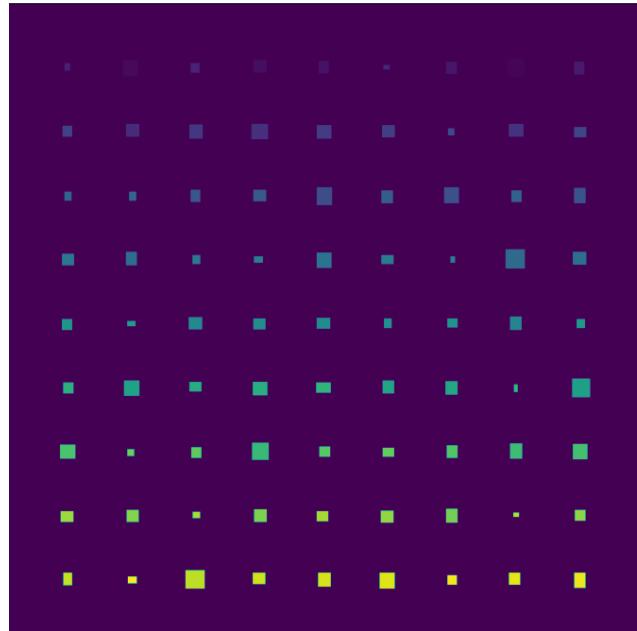
Feature selection: challenges

- Features are not independent
 - Area and diameter
 - Roundness, circularity, solidity, extent, aspect ratio, elongation, Feret's diameter, ...
- Best classification most likely involves multiple features
- Vast amount of features can hardly be visualized
- Need for dimensionality reduction
 - Principal component analysis (PCA)
 - t-Distributed Stochastic Neighbour Embedding (t-SNE)
 - Uniform Manifold Approximation and Projection (UMAP)
- Grouping objects (clustering)

PCA: Principal Component Analysis

Decomposes data into linear combinations of features that explain the highest variance

Example: Squares of different size

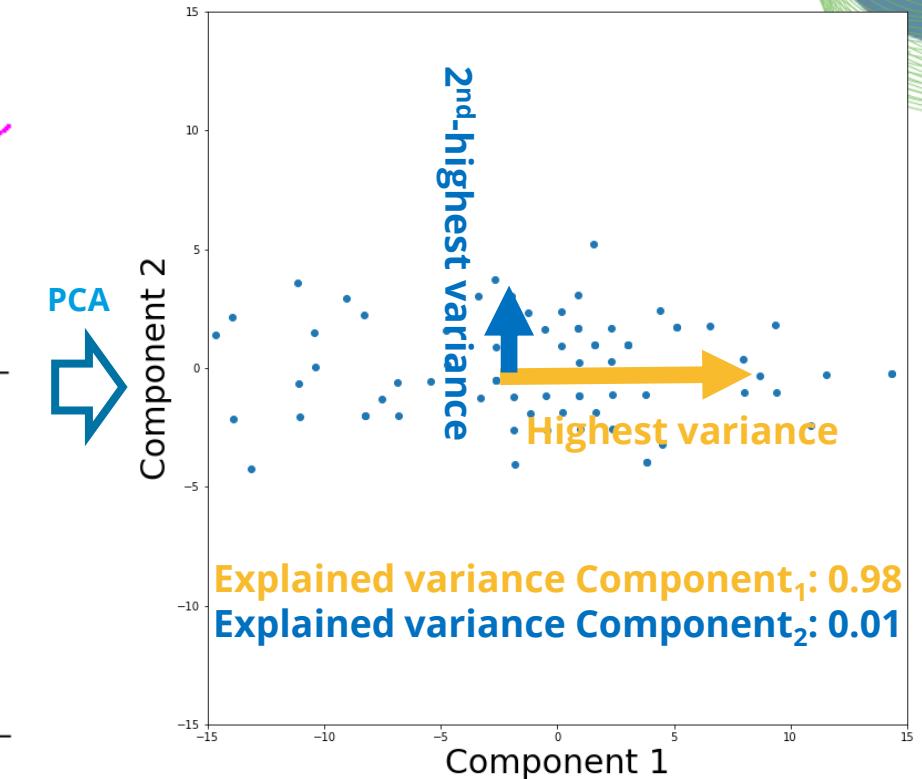
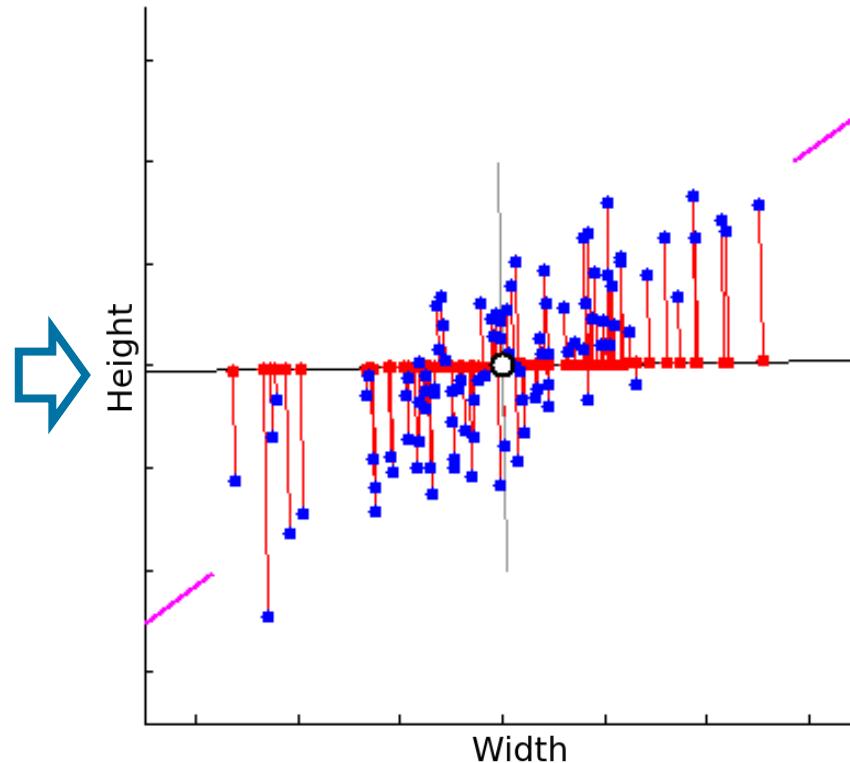
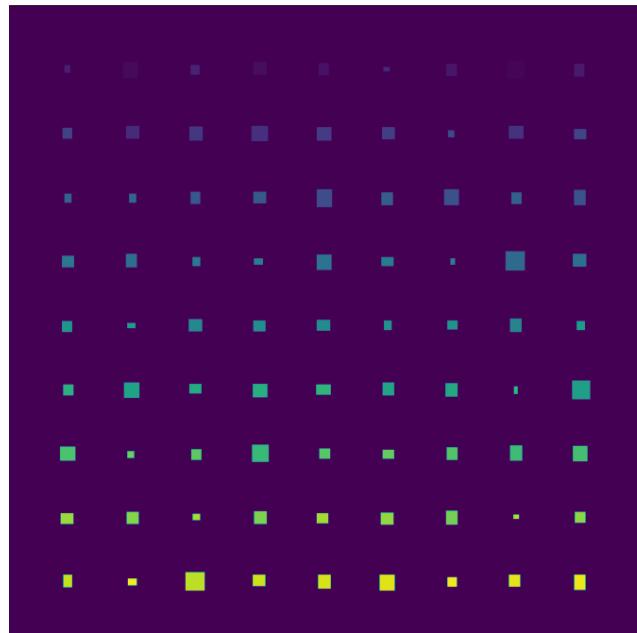


→ PCA transforms width/height measurements into a coordinate system that explains existing variance better

PCA: Principal Component Analysis

Decomposes data into linear combinations of features that explain the highest variance

Example: Squares of different size



→ PCA transforms width/height measurements into a coordinate system that explains existing variance better

PCA in Python: sklearn.decomposition.PCA

- Import package

```
from sklearn.decomposition import PCA
```

- Apply PCA

```
pca = PCA(n_components=2)  
pca.fit(standardized_data)
```

- Transform data into new coordinate system

```
transformed_data = pca.transform(data)
```

Important!

Always check the explained variance along the PCA component axes!

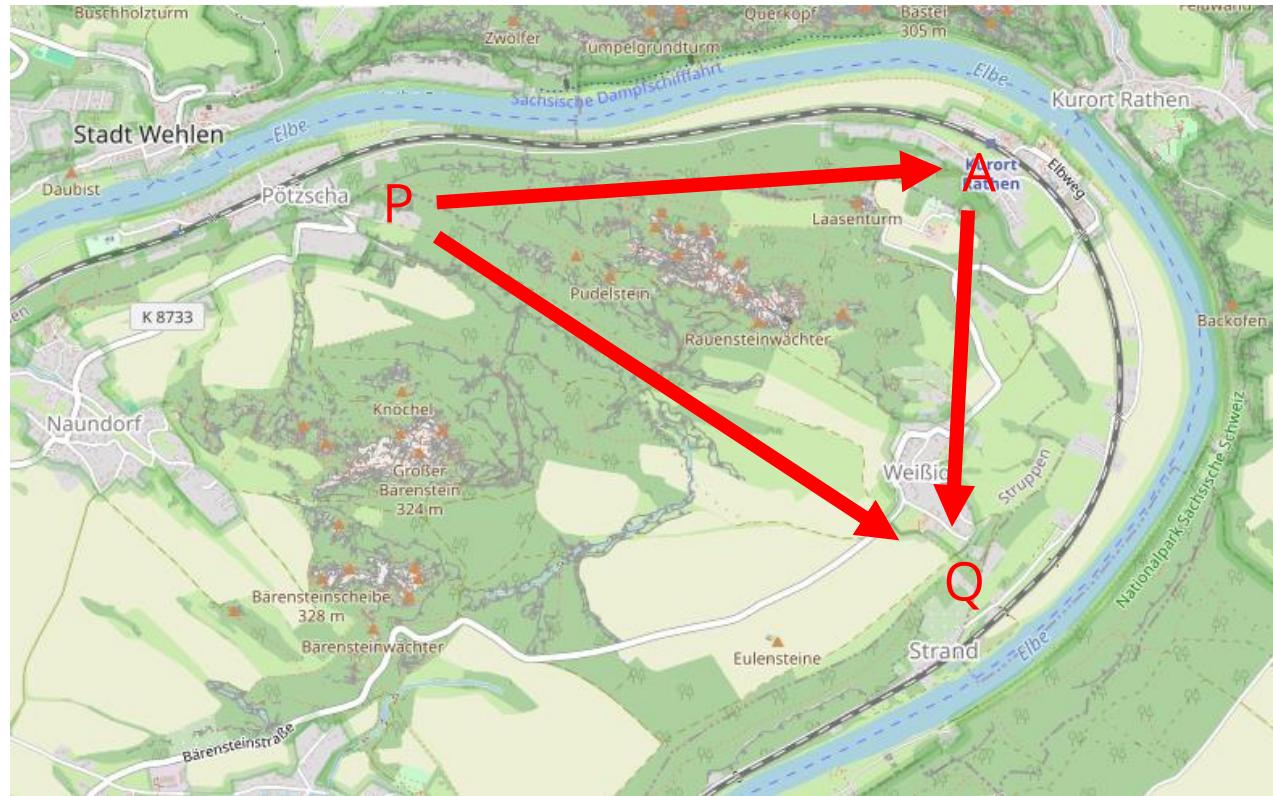
```
pca.explained_variance_ratio_
```

```
array([0.98773142, 0.01226858])
```

The screenshot shows the official scikit-learn website. At the top, there's a navigation bar with links for Install, User Guide, API, Examples, Community, and More. Below the header, the title "scikit-learn" and subtitle "Machine Learning in Python" are displayed. There are three main sections: "Classification" (with a grid of small images), "Regression" (with a line plot), and "Clustering" (with a scatter plot). Each section has a brief description, applications, and algorithms listed, followed by an "Examples" button.

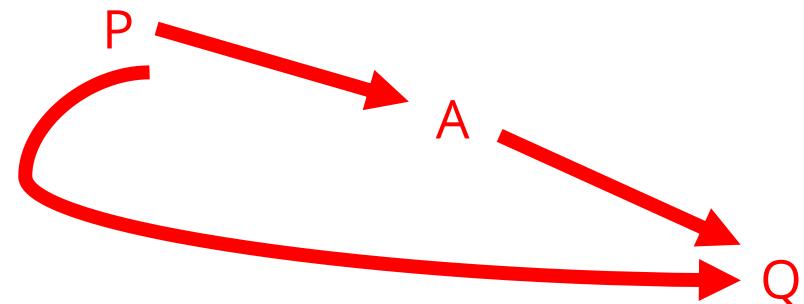
Non-Euclidian spaces

Not all dimensions (features) might be distances



Use travel time between P and Q as metric for distance

→ Travelling from Stadt Wehlen to Strand by bike is probably faster if you make a detour through Rathen



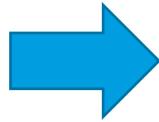
Dimensionality reduction: UMAP

Uniform Manifold Approximation Projection

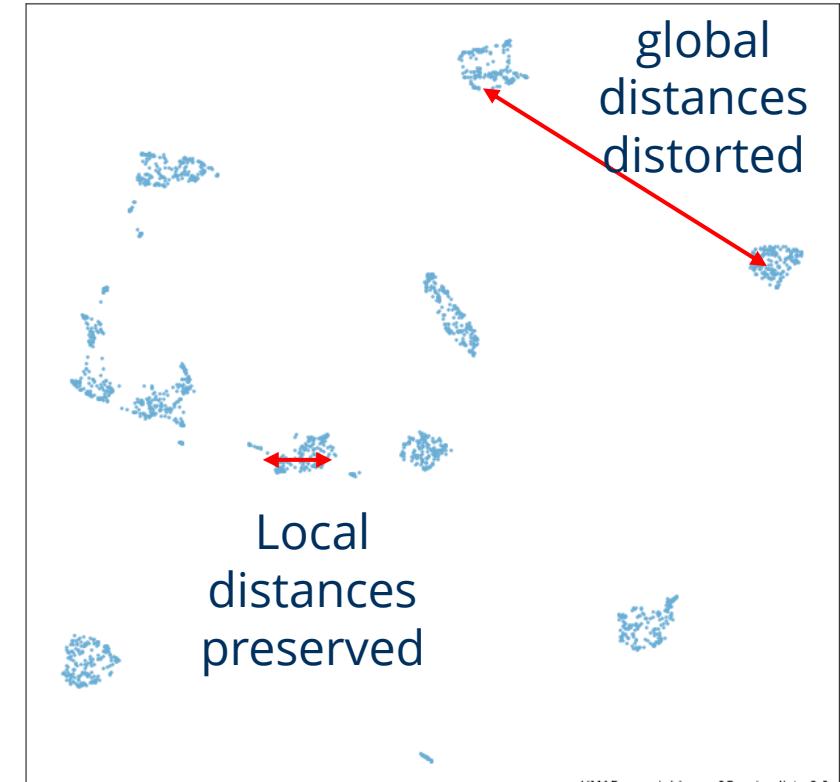
Preserve local distances at the expense of global distortions

Many dimensions

| | count | mean | std |
|------------------------------|-------|------------|------------|
| label | 44.0 | 22.500000 | 12.845233 |
| area | 44.0 | 401.863636 | 202.852288 |
| bbox_area | 44.0 | 542.750000 | 295.106376 |
| equivalent_diameter | 44.0 | 21.781085 | 6.174086 |
| convex_area | 44.0 | 423.295455 | 216.613747 |
| max_intensity | 44.0 | 234.909091 | 17.517856 |
| mean_intensity | 44.0 | 190.116971 | 15.034153 |
| min_intensity | 44.0 | 128.000000 | 0.000000 |
| extent | 44.0 | 0.758804 | 0.063276 |
| local_centroid-0 | 44.0 | 11.439824 | 4.126230 |
| local_centroid-1 | 44.0 | 10.138666 | 3.491815 |
| solidity | 44.0 | 0.953153 | 0.024749 |
| feret_diameter_max | 44.0 | 26.382434 | 8.915046 |
| major_axis_length | 44.0 | 25.876797 | 9.591558 |
| minor_axis_length | 44.0 | 18.872898 | 5.158791 |
| orientation | 44.0 | 0.053057 | 0.691430 |
| eccentricity | 44.0 | 0.600434 | 0.165688 |
| standard_deviation_intensity | 44.0 | 29.556705 | 5.507399 |
| aspect_ratio | 44.0 | 1.374342 | 0.397611 |
| roundness | 44.0 | 0.762889 | 0.156695 |
| circularity | 44.0 | 0.918858 | 0.133288 |



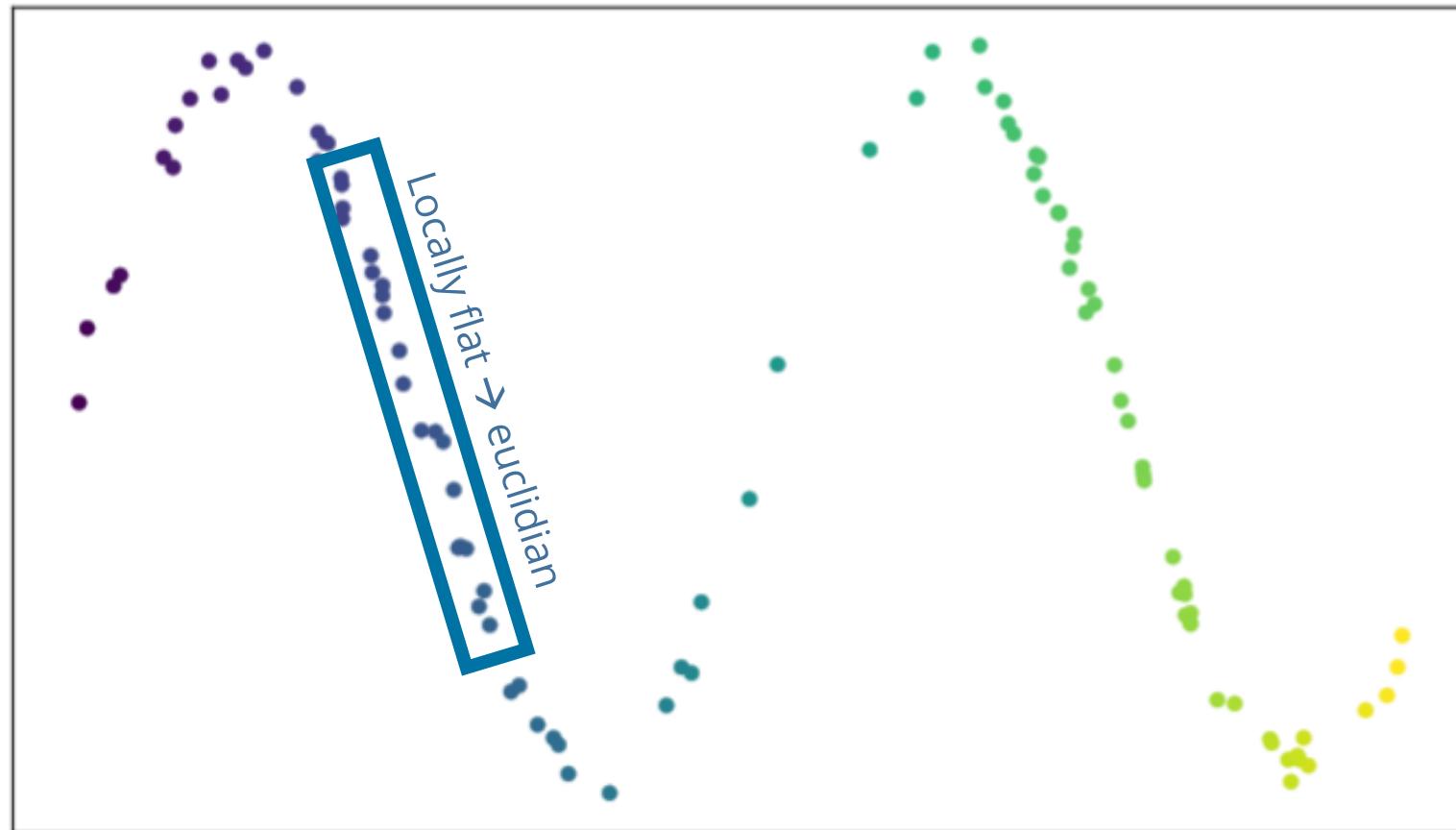
UMAP 2



UMAP 1

Dimensionality reduction: UMAP

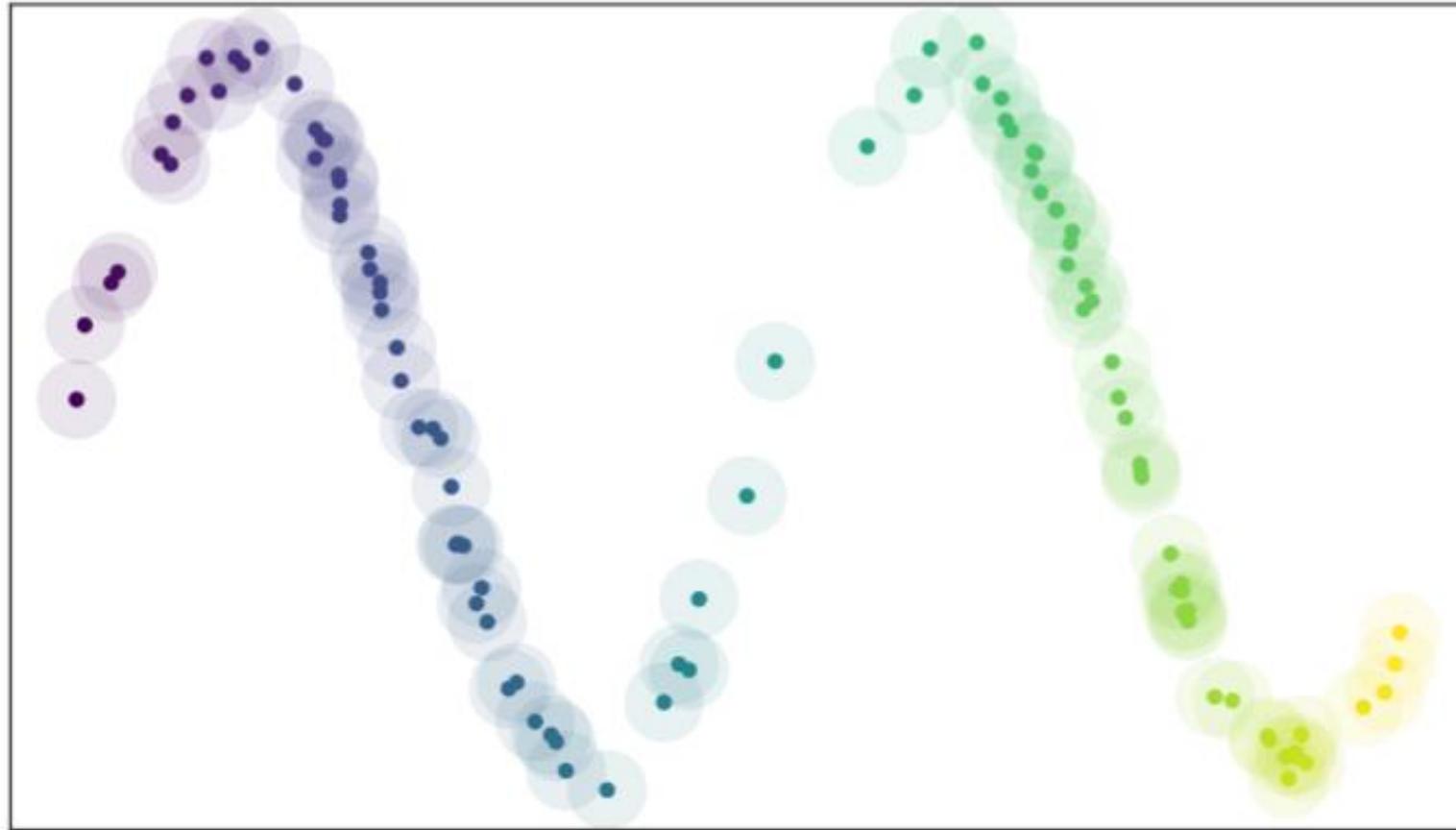
Initial situation: Our data suggests an underlying structure ("topology")



Goal:

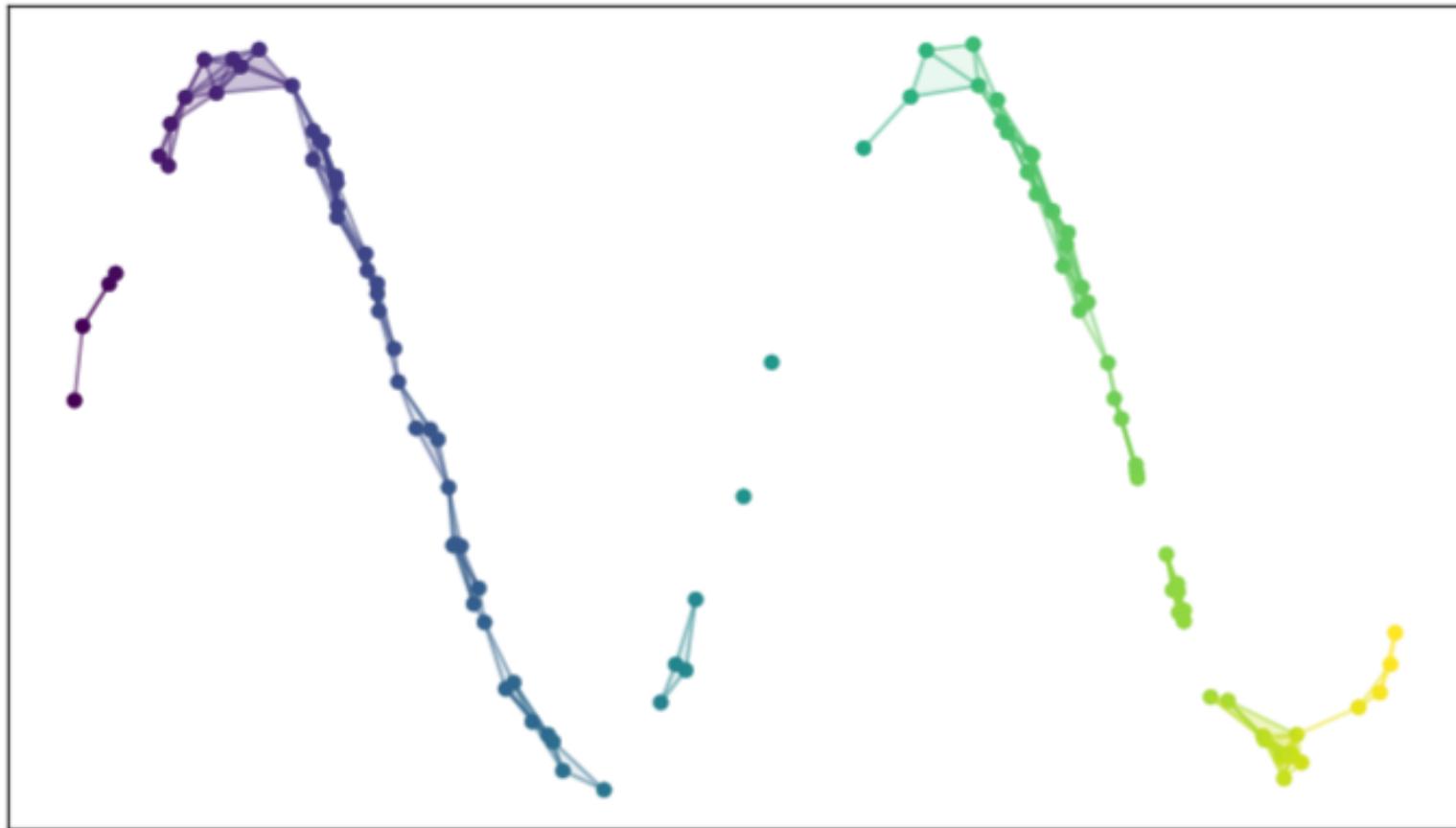
Reconstruct underlying topology to identify a space that best explains differences in our data

Dimensionality reduction



Naïve approach:
Points within a
defined radius are
considered
neighbors

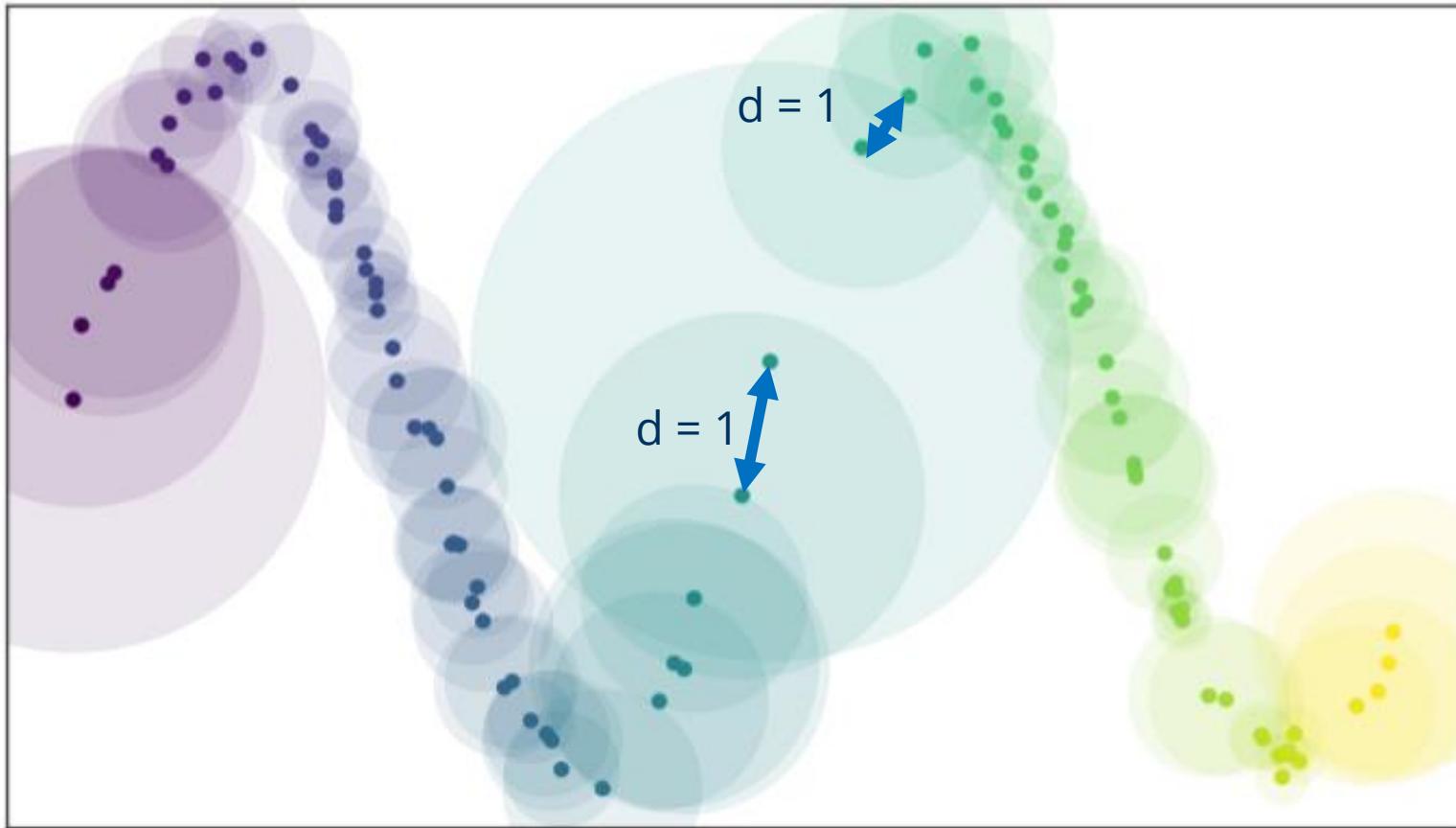
Dimensionality reduction



Naïve approach:
Points within a
defined radius are
considered
neighbors

Result:
Neighborhood
graph with
interruptions

Dimensionality reduction: UMAP

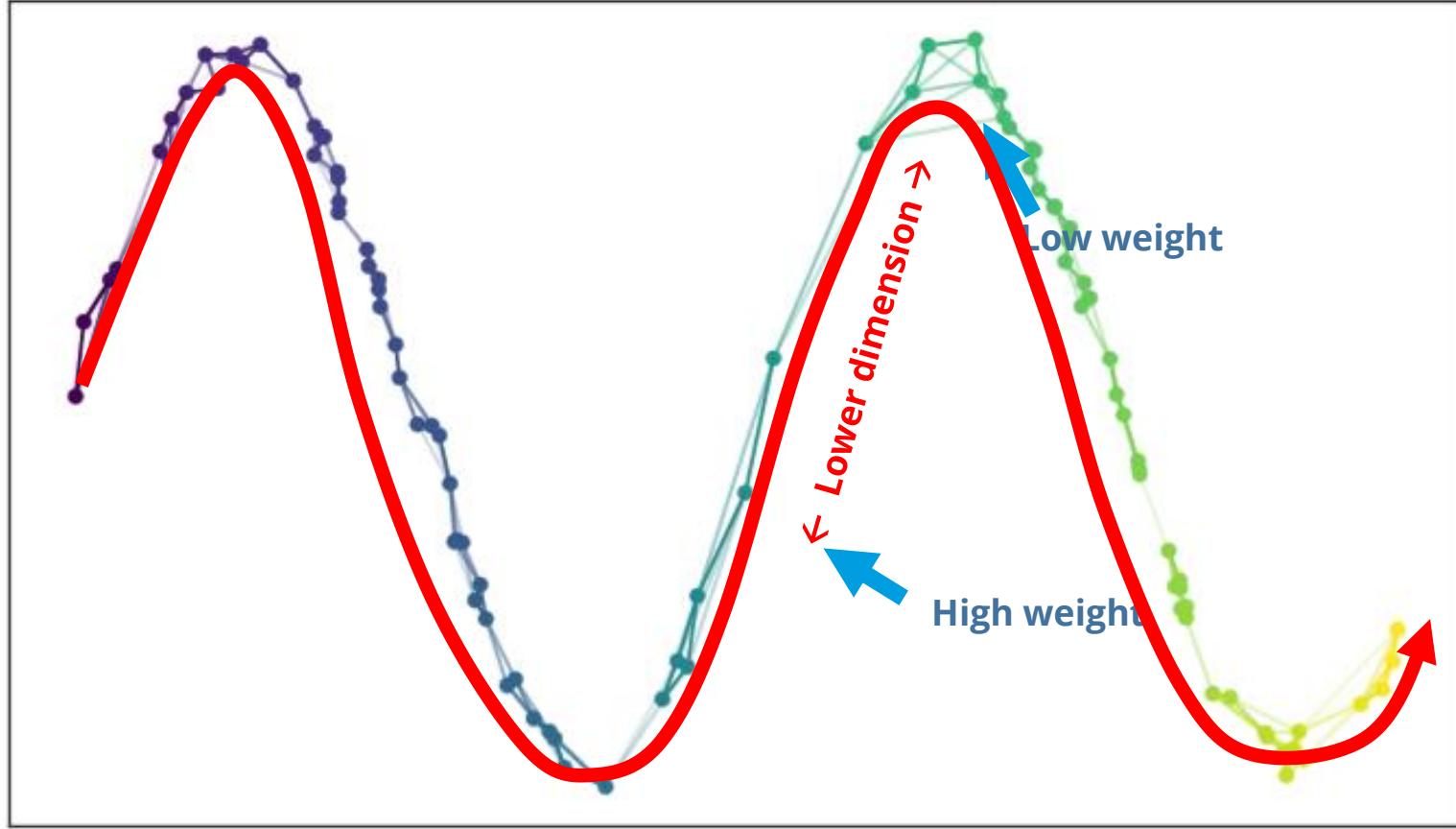


Approach:

Normalize distances
by dividing by the
average distance to
 n nearest neighbors

(Example: $n=1$)

Reduce dimensionality preserving fuzzy topology



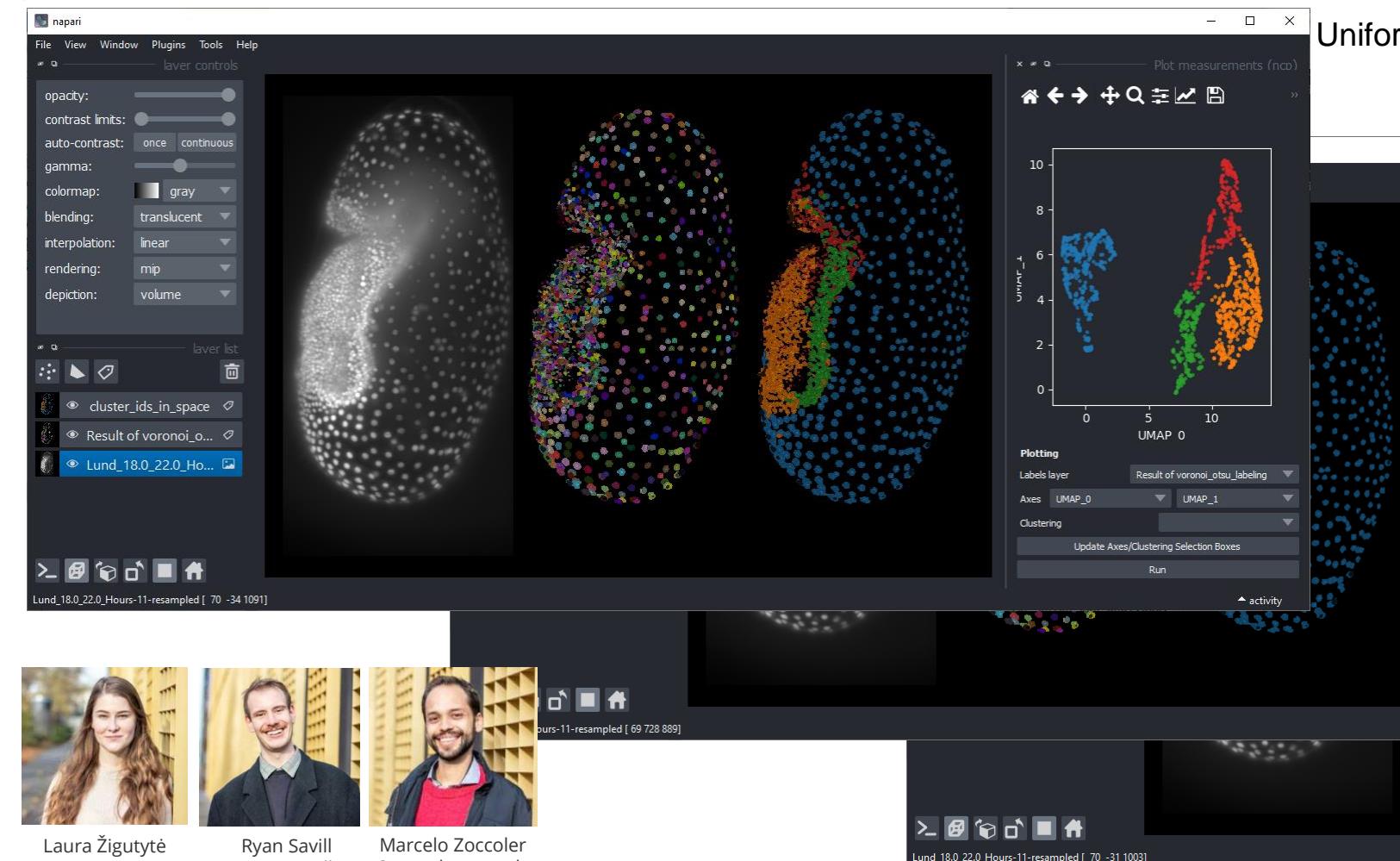
Approach:

Normalize distances
by dividing by the
average distance to
n nearest neighbors

Build a graph
considering
normalized
distances

Project data into
lower dimensional
space

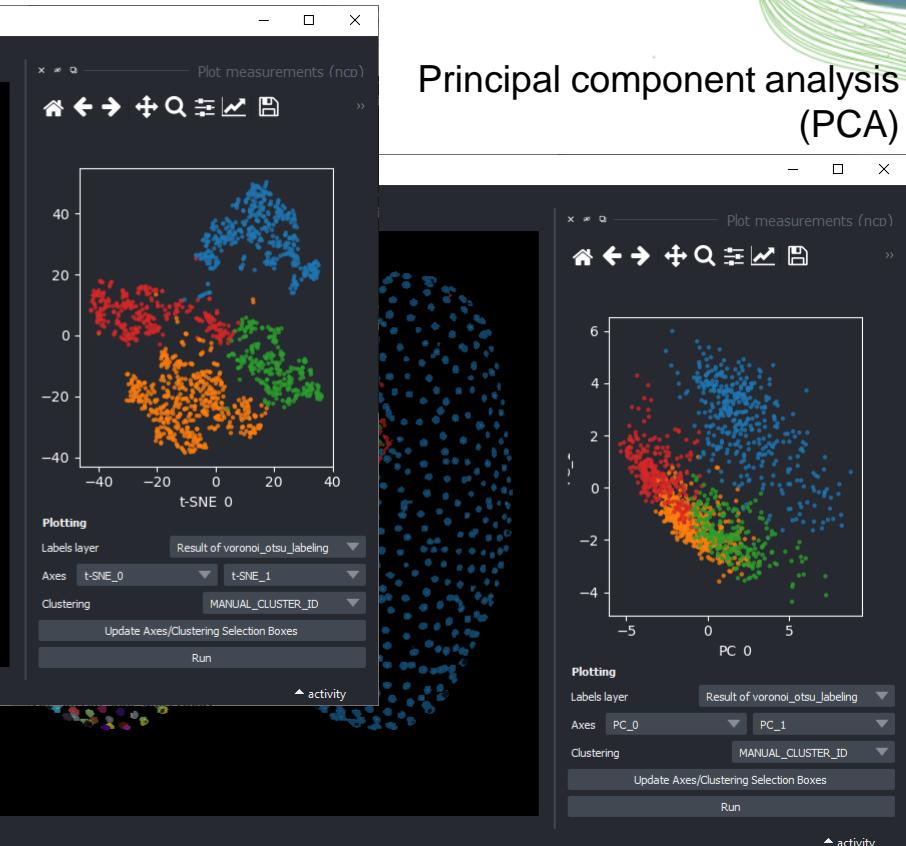
Dimensionality reduction



Uniform manifold approximation and projection (UMAP)

t-distributed stochastic neighbor embedding (t-SNE)

Principal component analysis (PCA)



UMAP in Python

Selecting columns from a pandas DataFrame

```
[8]: measurements.describe().T
```

| | count | mean | std |
|------------------------------|-------|------------|------------|
| label | 44.0 | 22.500000 | 12.845233 |
| area | 44.0 | 401.863636 | 202.852288 |
| bbox_area | 44.0 | 542.750000 | 295.106376 |
| equivalent_diameter | 44.0 | 21.781085 | 6.174086 |
| convex_area | 44.0 | 423.295455 | 216.613747 |
| max_intensity | 44.0 | 234.909091 | 17.517856 |
| mean_intensity | 44.0 | 190.116971 | 15.034153 |
| min_intensity | 44.0 | 128.000000 | 0.000000 |
| extent | 44.0 | 0.758804 | 0.063276 |
| local_centroid-0 | 44.0 | 11.439824 | 4.126230 |
| local_centroid-1 | 44.0 | 10.138666 | 3.491815 |
| solidity | 44.0 | 0.953153 | 0.024749 |
| feret_diameter_max | 44.0 | 26.382434 | 8.915046 |
| major_axis_length | 44.0 | 25.876797 | 9.591558 |
| minor_axis_length | 44.0 | 18.872898 | 5.158791 |
| orientation | 44.0 | 0.053057 | 0.691430 |
| eccentricity | 44.0 | 0.600434 | 0.165688 |
| standard_deviation_intensity | 44.0 | 29.556705 | 5.507399 |
| aspect_ratio | 44.0 | 1.374342 | 0.397611 |
| roundness | 44.0 | 0.762889 | 0.156695 |
| circularity | 44.0 | 0.918858 | 0.133288 |

```
[9]: selected_measurements = measurements[[
    'area',
    'equivalent_diameter',
    'convex_area',
    'max_intensity',
    'mean_intensity',
    'min_intensity',
    'extent',
    'solidity',
    'feret_diameter_max',
    'major_axis_length',
    'minor_axis_length',
    'eccentricity',
    'standard_deviation_intensity',
    'aspect_ratio',
    'roundness',
    'circularity']]
```

```
selected_measurements.describe().T
```

```
[9]:
```

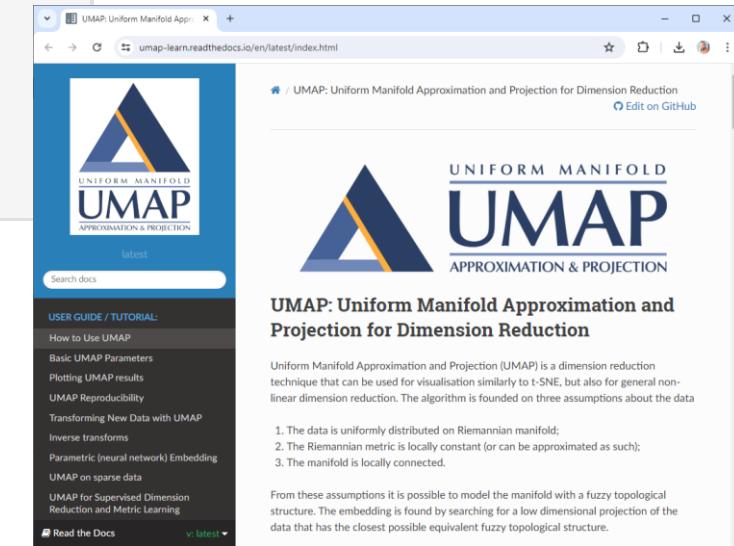
| | count | mean | std |
|------------------------------|-------|------------|------------|
| area | 44.0 | 401.863636 | 202.852288 |
| equivalent_diameter | 44.0 | 21.781085 | 6.174086 |
| convex_area | 44.0 | 423.295455 | 216.613747 |
| max_intensity | 44.0 | 234.909091 | 17.517856 |
| mean_intensity | 44.0 | 190.116971 | 15.034153 |
| min_intensity | 44.0 | 128.000000 | 0.000000 |
| extent | 44.0 | 0.758804 | 0.063276 |
| solidity | 44.0 | 0.953153 | 0.024749 |
| feret_diameter_max | 44.0 | 26.382434 | 8.915046 |
| major_axis_length | 44.0 | 25.876797 | 9.591558 |
| minor_axis_length | 44.0 | 18.872898 | 5.158791 |
| eccentricity | 44.0 | 0.600434 | 0.165688 |
| standard_deviation_intensity | 44.0 | 29.556705 | 5.507399 |
| aspect_ratio | 44.0 | 1.374342 | 0.397611 |
| roundness | 44.0 | 0.762889 | 0.156695 |
| circularity | 44.0 | 0.918858 | 0.133288 |

Select reasonable features

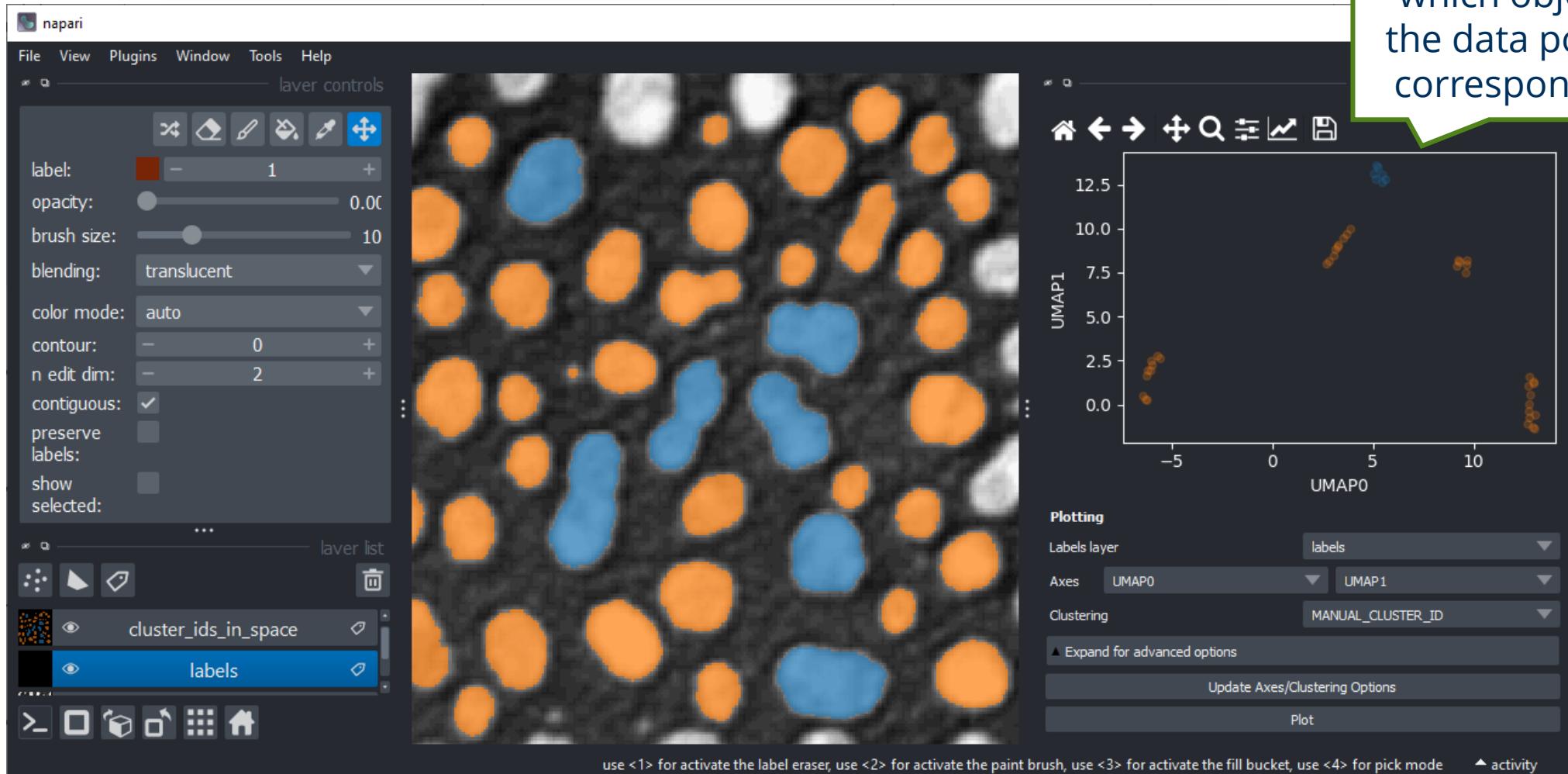
UMAP in Python

```
[10]: # configure UMAP algorithm  
umap = UMAP(n_neighbors=5, n_components=2)  
  
# apply algorithm  
transformed_data = umap.fit_transform(selected_measurements.values.tolist())  
  
# store results back in table  
measurements['UMAP0'] = transformed_data[:,0]  
measurements['UMAP1'] = transformed_data[:,1]
```

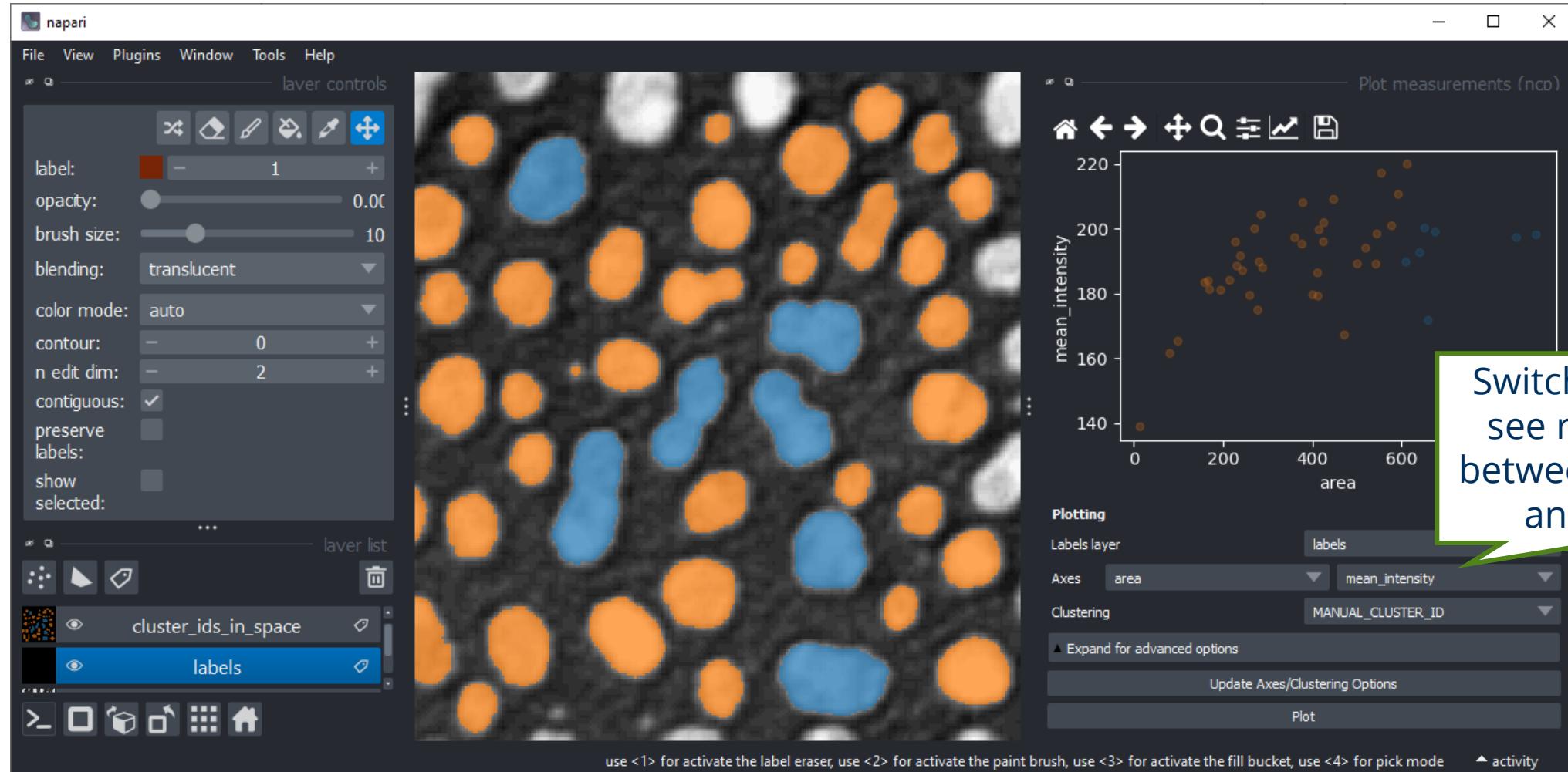
Data conversion



Annotating UMAPs in Napari



Interpreting annotations in Napari



Correlation statistics

```
[16]: def colorize(styler):
    styler.background_gradient(axis=None, cmap="PiYG")
    return styler

df = measurements.corr().T
df.style.pipe(colorize)
```

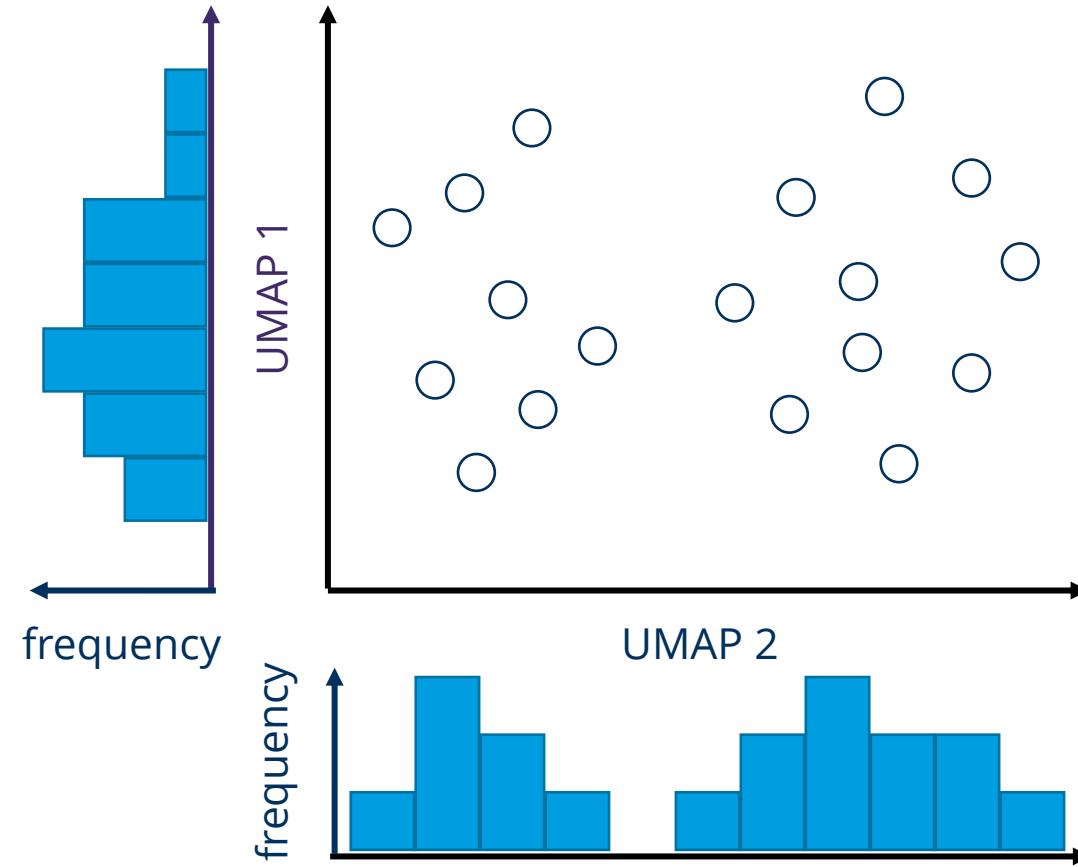
| | label | area | bbox_area | equivalent_diameter | convex_area | max_intensity | mean_intensity | min_intensity | extent | local_centroid-0 | local_centroid-1 | solidity | feret_diameter_max | major_axis_length | minor_axis_length | orientation | eccentricity |
|------------------------------|-----------|-----------|-----------|---------------------|-------------|---------------|----------------|---------------|-----------|------------------|------------------|-----------|--------------------|-------------------|-------------------|-------------|--------------|
| label | 1.000000 | 0.261682 | 0.223070 | 0.249249 | 0.250594 | 0.110791 | 0.235692 | nan | 0.031673 | 0.177363 | 0.227746 | 0.090163 | 0.208067 | 0.198908 | 0.237521 | 0.319053 | 0.059804 |
| area | 0.261682 | 1.000000 | 0.973718 | 0.978723 | 0.997560 | 0.511730 | 0.530250 | nan | -0.362472 | 0.847281 | 0.935689 | -0.243908 | 0.930981 | 0.911069 | 0.859240 | 0.280673 | 0.348585 |
| bbox_area | 0.223070 | 0.973718 | 1.000000 | 0.948328 | 0.985584 | 0.481524 | 0.476951 | nan | -0.546728 | 0.902854 | 0.904551 | -0.416707 | 0.973189 | 0.967337 | 0.752580 | 0.213080 | 0.479196 |
| equivalent_diameter | 0.249249 | 0.978723 | 0.948328 | 1.000000 | 0.974614 | 0.633984 | 0.618553 | nan | -0.395696 | 0.858779 | 0.947036 | -0.266587 | 0.931696 | 0.904412 | 0.904698 | 0.197456 | 0.363799 |
| convex_area | 0.250594 | 0.997560 | 0.985584 | 0.974614 | 1.000000 | 0.506730 | 0.517356 | nan | -0.413323 | 0.862417 | 0.934090 | -0.305706 | 0.948048 | 0.932682 | 0.832264 | 0.263176 | 0.389269 |
| max_intensity | 0.110791 | 0.511730 | 0.481524 | 0.633984 | 0.506730 | 1.000000 | 0.825115 | nan | -0.324093 | 0.504879 | 0.603305 | -0.253635 | 0.536089 | 0.502524 | 0.645600 | -0.139025 | 0.246172 |
| mean_intensity | 0.235692 | 0.530250 | 0.476951 | 0.618553 | 0.517356 | 0.825115 | 1.000000 | nan | -0.160940 | 0.412859 | 0.609264 | -0.077797 | 0.458515 | 0.422638 | 0.707711 | 0.132754 | 0.017030 |
| min_intensity | nan | nan | nan | nan | nan | nan | nan | nan | nan | nan | nan | nan | nan | nan | nan | nan | nan |
| extent | 0.031673 | -0.362472 | -0.546728 | -0.395696 | -0.413323 | -0.324093 | -0.160940 | nan | 1.000000 | -0.631158 | -0.375580 | 0.853431 | -0.631776 | -0.664733 | -0.062873 | 0.252915 | -0.756019 |
| local_centroid-0 | 0.177363 | 0.847281 | 0.902854 | 0.858779 | 0.862417 | 0.504879 | 0.412859 | nan | -0.631158 | 1.000000 | 0.706437 | -0.439244 | 0.937673 | 0.932889 | 0.623186 | 0.003490 | 0.560853 |
| local_centroid-1 | 0.227746 | 0.935689 | 0.904551 | 0.947036 | 0.934090 | 0.603305 | 0.609264 | nan | -0.375580 | 0.706437 | 1.000000 | -0.290177 | 0.863585 | 0.840724 | 0.875044 | 0.271191 | 0.318154 |
| solidity | 0.090163 | -0.243908 | -0.416707 | -0.266587 | -0.305706 | -0.253635 | -0.077797 | nan | 0.853431 | -0.439244 | -0.290177 | 1.000000 | -0.512903 | -0.556555 | 0.049965 | 0.279509 | -0.723572 |
| feret_diameter_max | 0.208067 | 0.930981 | 0.973189 | 0.931696 | 0.948048 | 0.536089 | 0.458515 | nan | -0.631776 | 0.937673 | 0.863585 | -0.512903 | 1.000000 | 0.996744 | 0.690639 | 0.077145 | 0.614849 |
| major_axis_length | 0.198908 | 0.911069 | 0.967337 | 0.904412 | 0.932682 | 0.502524 | 0.422638 | nan | -0.664733 | 0.932889 | 0.840724 | -0.556555 | 0.996744 | 1.000000 | 0.639308 | 0.076773 | 0.647021 |
| minor_axis_length | 0.237521 | 0.859240 | 0.752580 | 0.904698 | 0.832264 | 0.645600 | 0.707711 | nan | -0.062873 | 0.623186 | 0.875044 | 0.049965 | 0.690639 | 0.639308 | 1.000000 | 0.278107 | -0.012148 |
| orientation | 0.319053 | 0.280673 | 0.213080 | 0.197456 | 0.263176 | -0.139025 | 0.132754 | nan | 0.252915 | 0.003490 | 0.271191 | 0.279509 | 0.077145 | 0.076773 | 0.278107 | 1.000000 | -0.305652 |
| eccentricity | 0.059804 | 0.348585 | 0.479196 | 0.363799 | 0.389269 | 0.246172 | 0.017030 | nan | -0.756019 | 0.560853 | 0.318154 | -0.723572 | 0.614849 | 0.647021 | -0.012148 | -0.305652 | 1.000000 |
| standard_deviation_intensity | 0.189165 | 0.288670 | 0.267528 | 0.402328 | 0.285105 | 0.867057 | 0.902001 | nan | -0.216260 | 0.284331 | 0.379400 | -0.169801 | 0.306228 | 0.280378 | 0.455324 | -0.089349 | 0.107307 |
| aspect_ratio | 0.036433 | 0.411794 | 0.581132 | 0.386884 | 0.462720 | 0.121313 | -0.044872 | nan | -0.848271 | 0.678234 | 0.321805 | -0.787587 | 0.690082 | 0.736200 | -0.030443 | -0.181927 | 0.853302 |
| roundness | -0.055815 | -0.415592 | -0.569335 | -0.406856 | -0.464090 | -0.191680 | 0.009002 | nan | 0.834550 | -0.638667 | -0.359961 | 0.801971 | -0.690444 | -0.732103 | 0.003699 | 0.224205 | -0.955978 |
| circularity | -0.054152 | -0.626241 | -0.718764 | -0.701230 | -0.659125 | -0.636372 | -0.411166 | nan | 0.808533 | -0.785693 | -0.644979 | 0.773934 | -0.832660 | -0.839196 | -0.435236 | 0.242901 | -0.779895 |
| UMAP0 | -0.065835 | -0.442711 | -0.413779 | -0.509190 | -0.435101 | -0.324496 | -0.387465 | nan | 0.168523 | -0.391875 | -0.488311 | 0.068021 | -0.457079 | -0.437340 | -0.479807 | 0.025473 | -0.204662 |
| UMAP1 | 0.139702 | 0.819263 | 0.813951 | 0.793707 | 0.821940 | 0.391350 | 0.365621 | nan | -0.375632 | 0.720004 | 0.753502 | -0.260000 | 0.753713 | 0.736954 | 0.702828 | 0.277117 | 0.251959 |
| MANUAL_CLUSTER_ID | 0.080739 | 0.677335 | 0.719434 | 0.590973 | 0.700457 | 0.156570 | 0.074372 | nan | -0.371454 | 0.582543 | 0.616873 | -0.418390 | 0.671673 | 0.686248 | 0.387847 | 0.163152 | 0.424045 |

My annotation
seems related to
area

My annotation
seems not related
to intensity

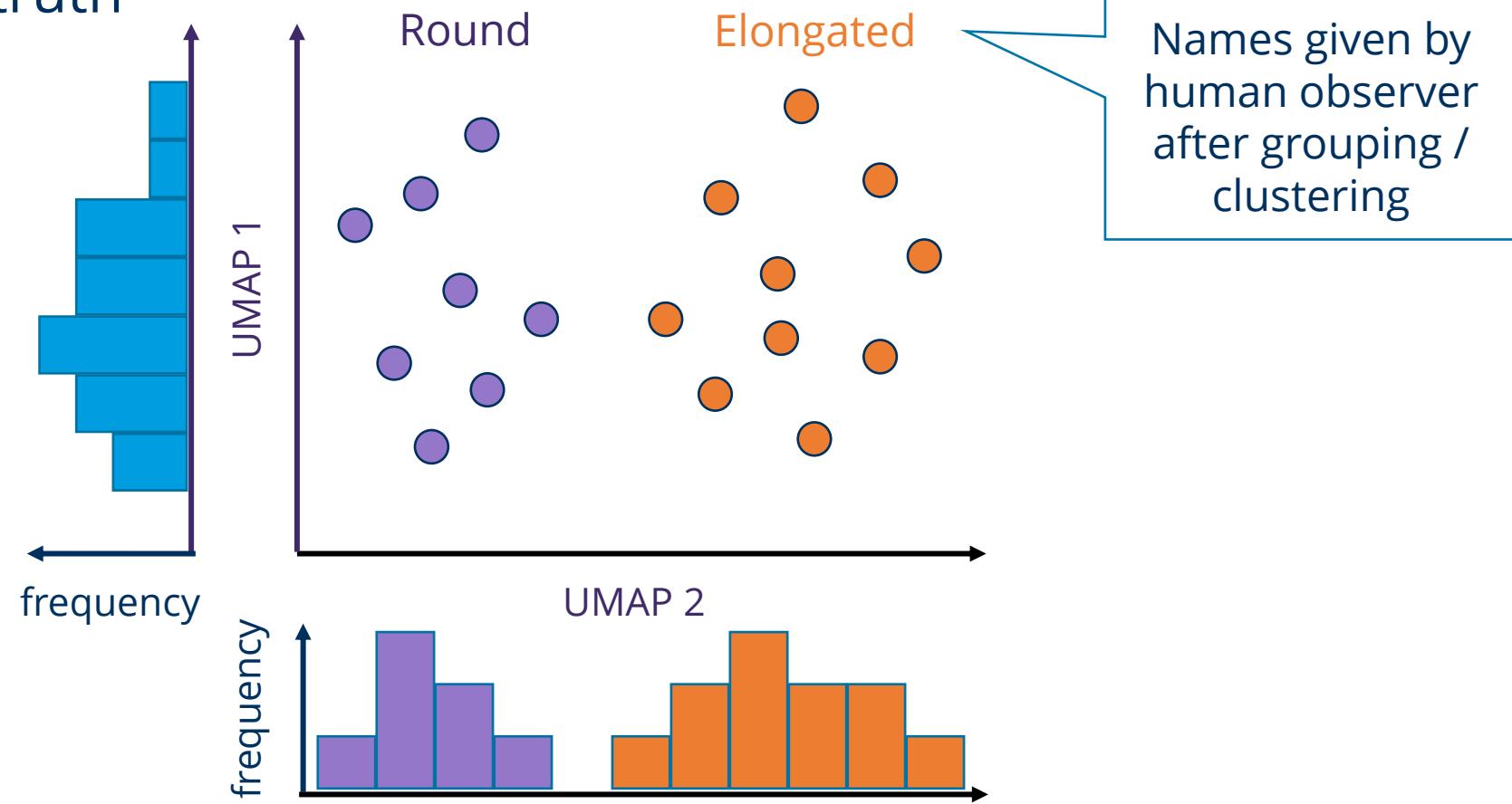
Clustering

Unsupervised machine learning may include grouping objects without given ground truth



Clustering

Unsupervised machine learning may include grouping objects without given ground truth



K-Means Clustering

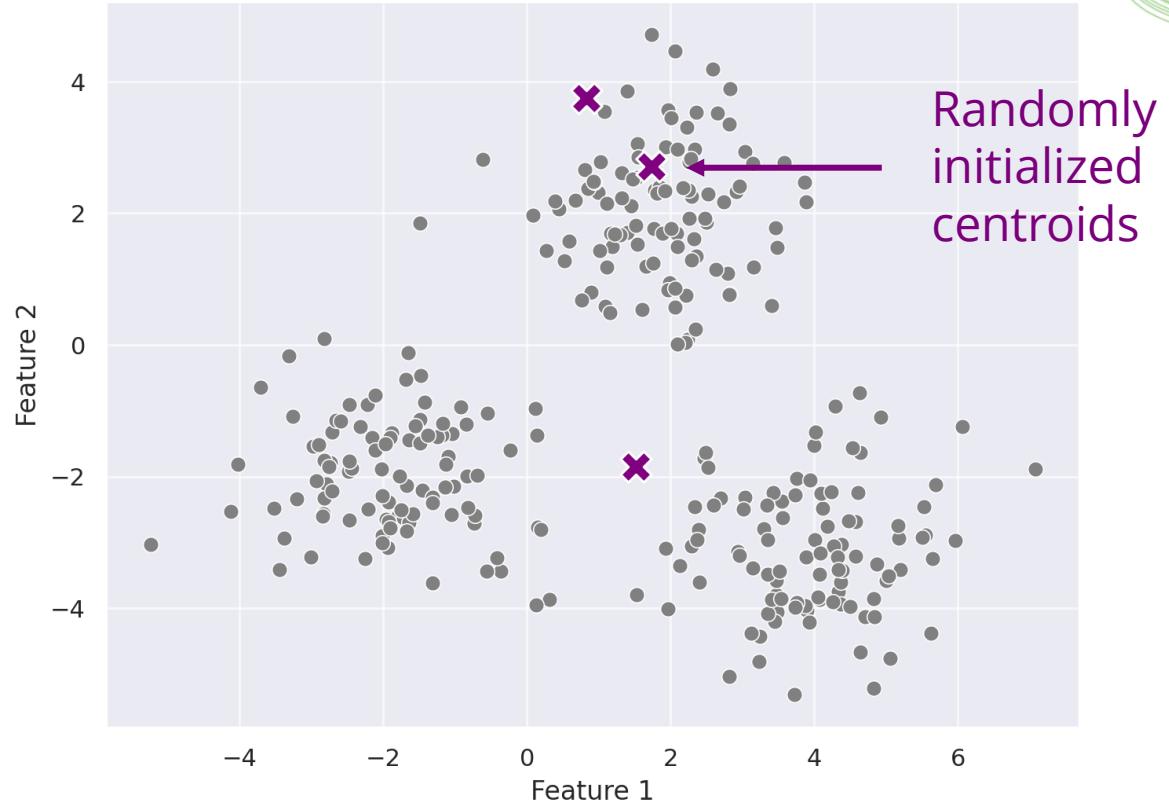
Goal: group data points into k groups so that variance within group is minimal.

STEP 1: Seed k initial cluster centroids randomly

STEP 2: Assign all points to nearest centroid

$$d(p, q) = \sqrt{\sum_{i=1}^n (p_i - q_i)^2} = \sqrt{(p_1 - q_1)^2 + (p_2 - q_2)^2}$$

n – dimensionality, in this example = 2



K-Means Clustering

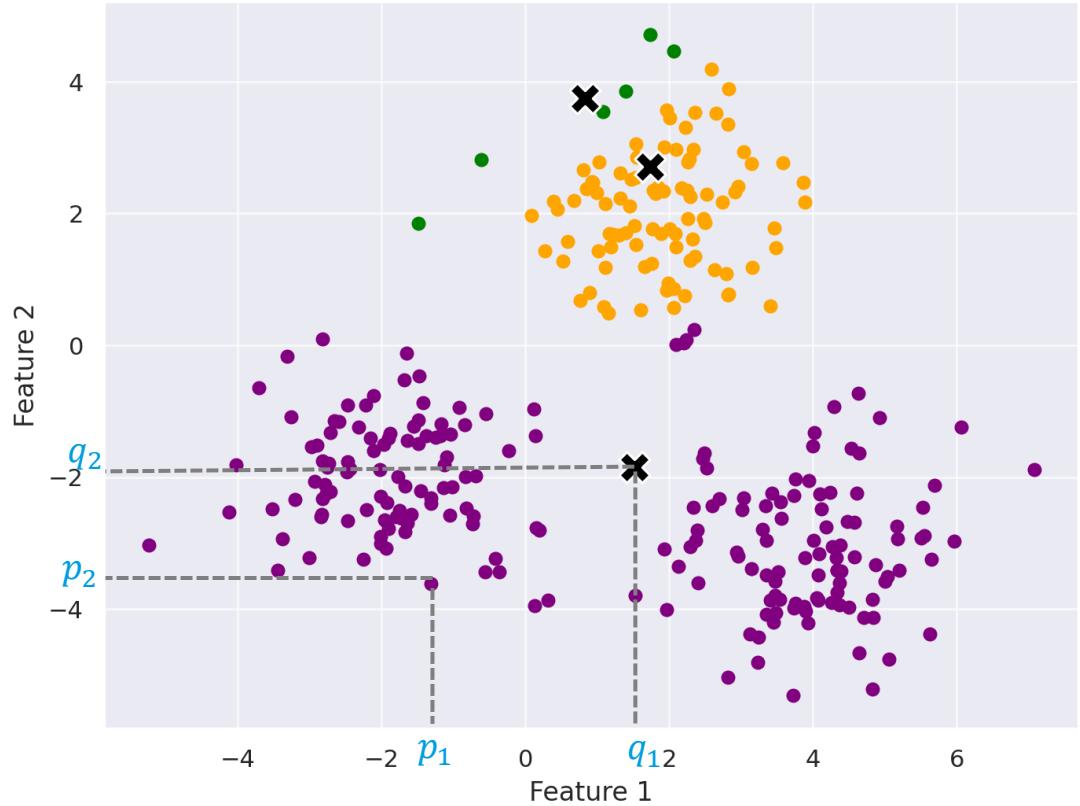
Goal: group data points into k groups so that variance within group is minimal.

STEP 1: Seed k initial cluster centroids randomly

STEP 2: Assign all points to nearest centroid

$$d(p, q) = \sqrt{\sum_{i=1}^n (p_i - q_i)^2} = \sqrt{(p_1 - q_1)^2 + (p_2 - q_2)^2}$$

n – dimensionality, in this example = 2



K-Means Clustering

Goal: group data points into k groups so that variance within group is minimal.

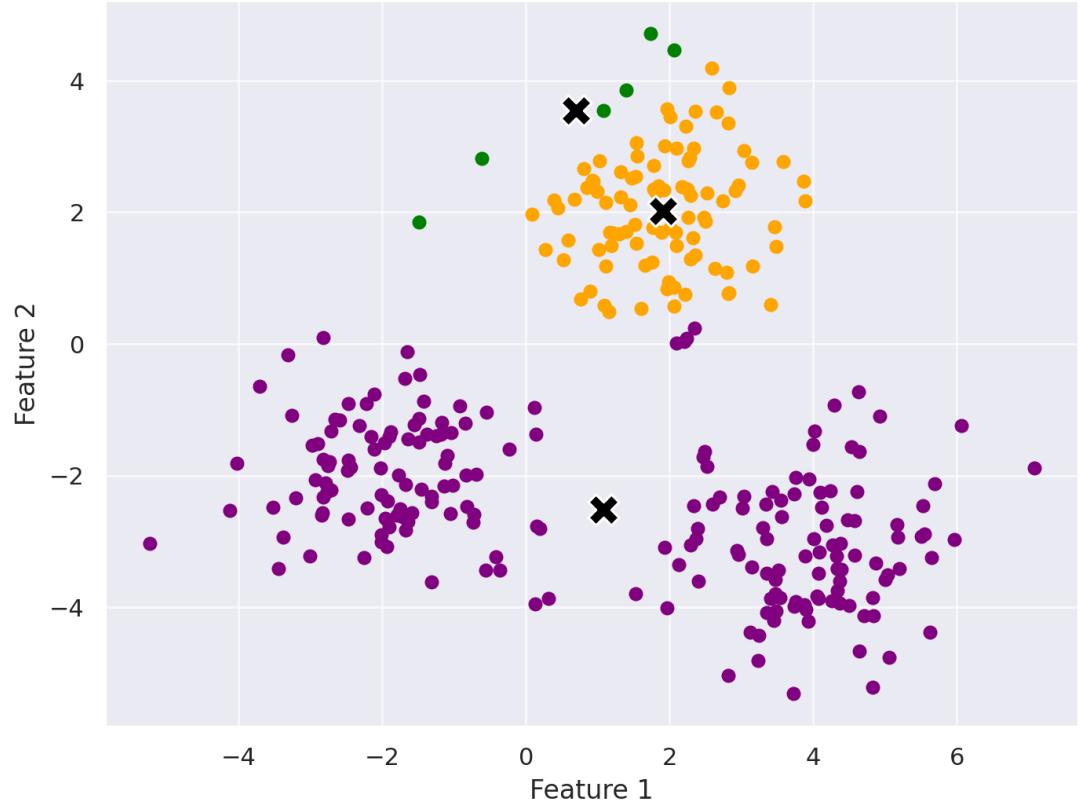
STEP 3: Determine new centroid positions as mean position of all assigned points.

$$\text{New centroid}_i = \frac{1}{|C_i|} \sum_{x \in C_i} x$$

C_i - the number of data points in cluster i

Repeat steps 2-3: the assignment and update steps are repeated iteratively until:

- Centroids not changing anymore,
- Point assignments not changing anymore or
- Maximum number of iterations reached



K-Means Clustering

Goal: group data points into k groups so that variance within group is minimal.

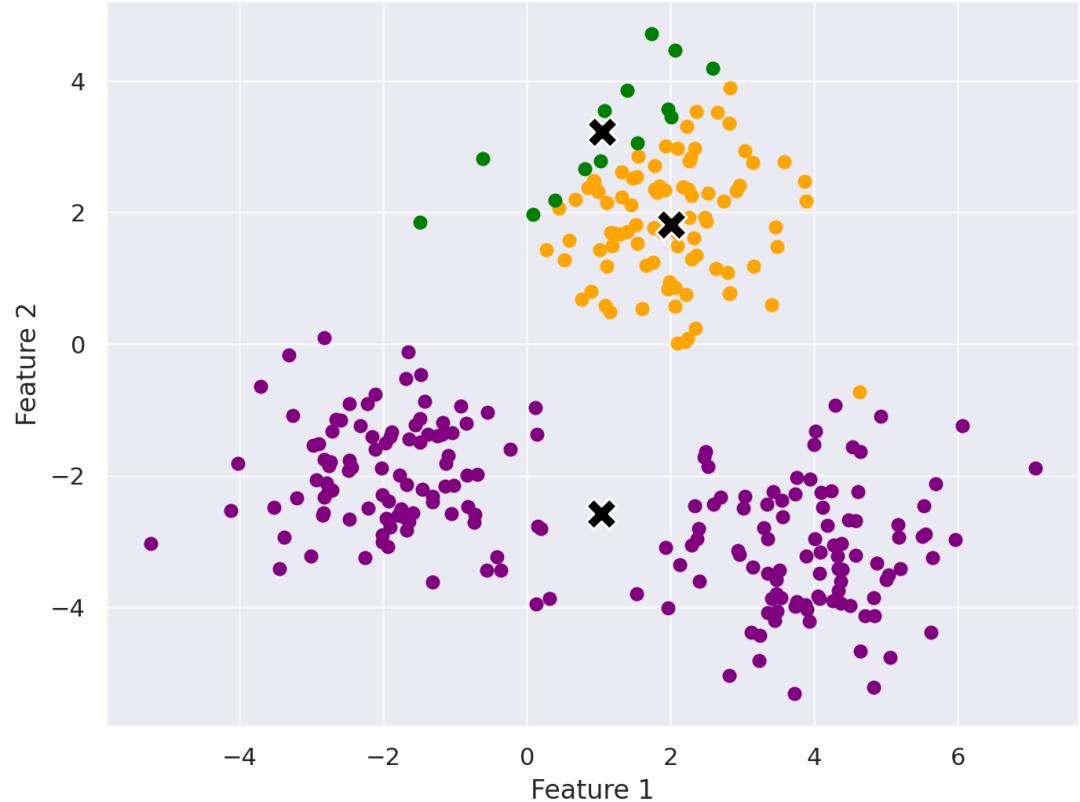
STEP 3: Determine new centroid positions as mean position of all assigned points.

$$\text{New centroid}_i = \frac{1}{|C_i|} \sum_{x \in C_i} x$$

C_i - the number of data points in cluster i

Repeat steps 2-3: the assignment and update steps are repeated iteratively until:

- Centroids not changing anymore,
- Point assignments not changing anymore or
- Maximum number of iterations reached



K-Means Clustering

Goal: group data points into k groups so that variance within group is minimal.

In Python:

```
from sklearn import cluster
```

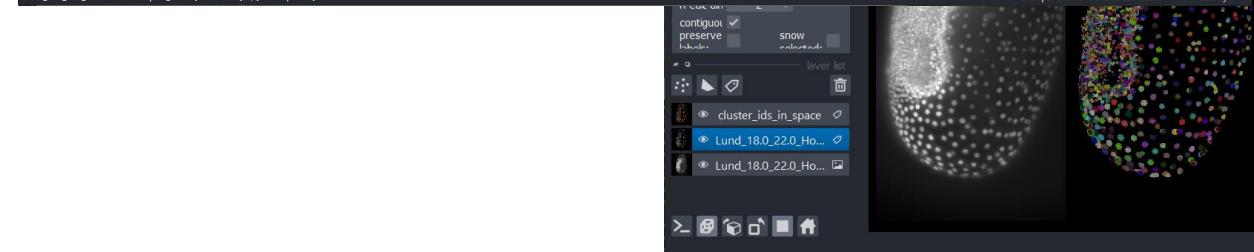
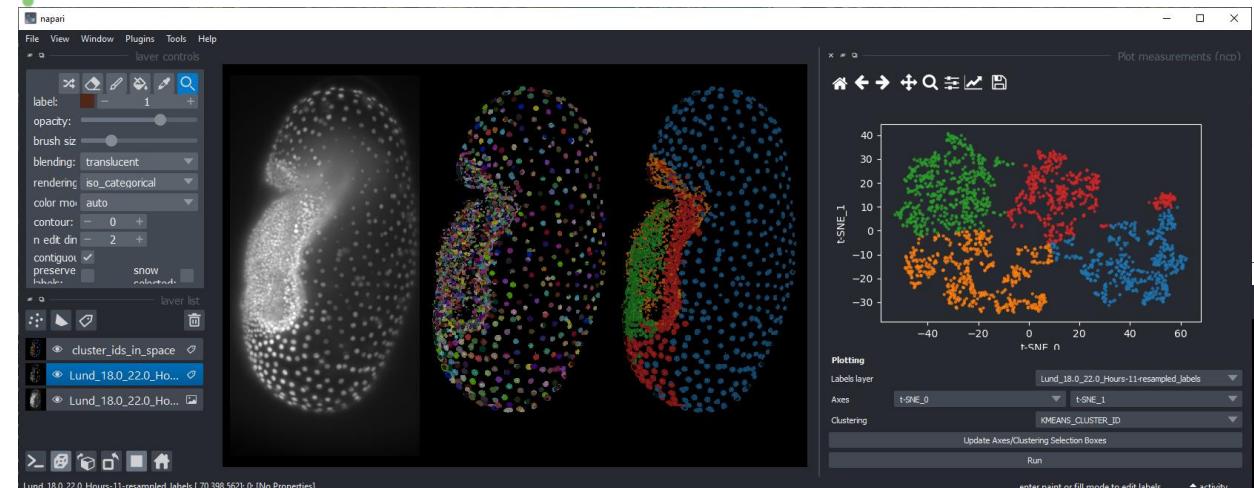
Create

```
clusterer = cluster.KMeans(n_clusters=3)  
clusterer.fit(X)
```

Predict

```
predicted_class = clusterer.predict(X)
```

Clustering



Laura Žigutytė
@zigutyte

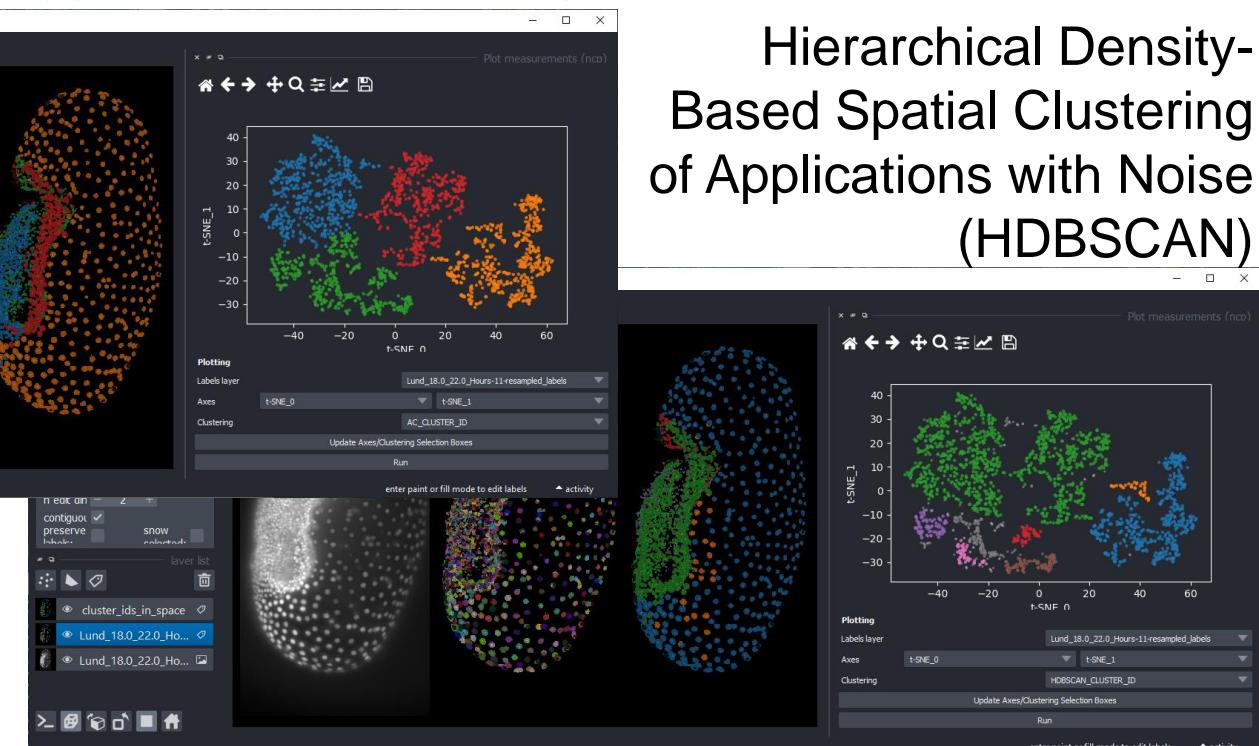
Ryan Savill
@RyanSavill4

Marcelo Zoccoler
@zoccolermarcelo

K-means clustering

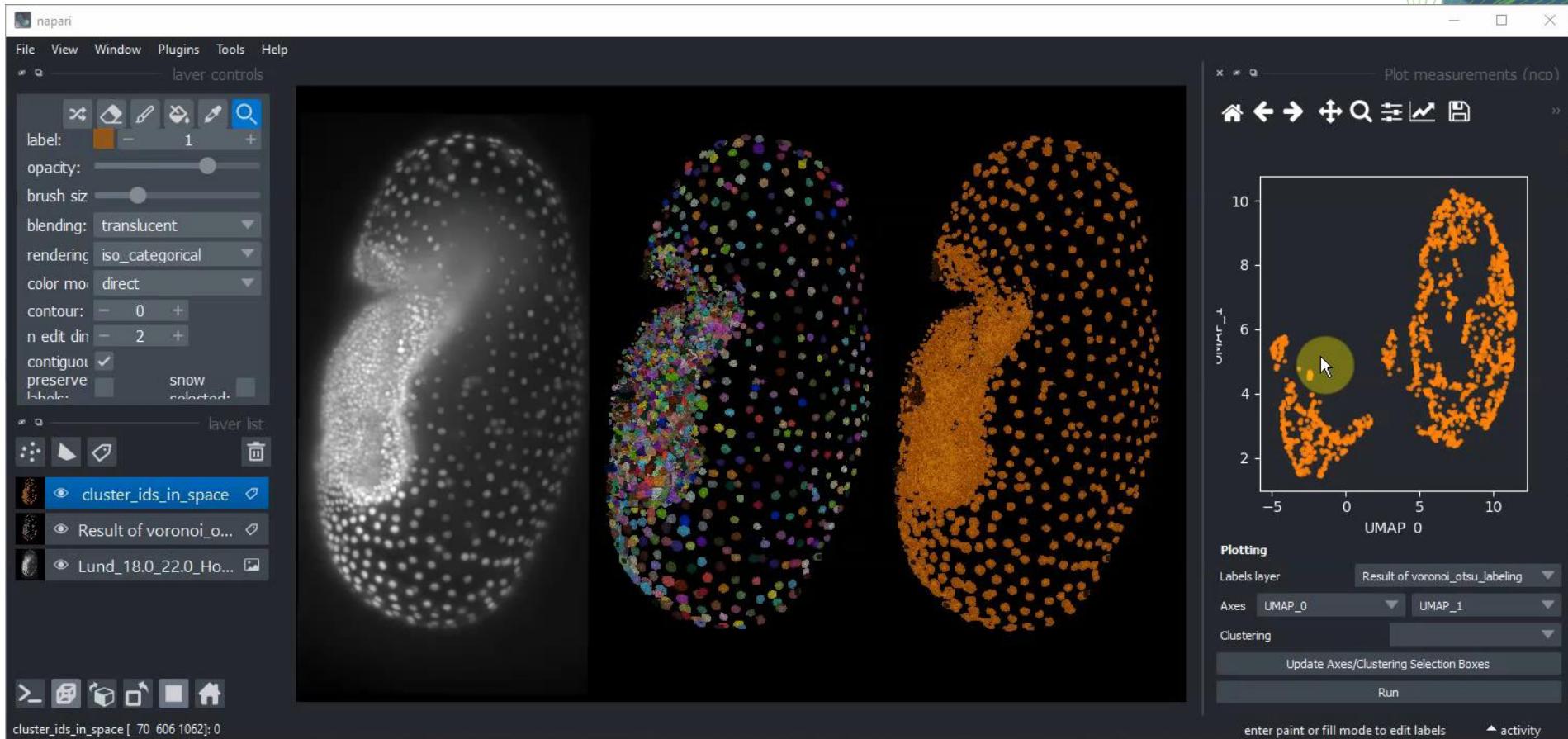
Agglomerative clustering

Hierarchical Density-Based Spatial Clustering of Applications with Noise (HDBSCAN)



Manual clustering

To better understand relationships between data



Laura Žigutytė
@zigutyte



Ryan Savill
@RyanSavill4



Marcelo Zoccoler
@zoccolermarcelo

Exercises

Robert Haase

Funded by



Bundesministerium
für Bildung
und Forschung



Diese Maßnahme wird gefördert durch die Bundesregierung
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der Grundlage des von den Abgeordneten des Sächsischen
Landtags beschlossenen Haushaltes.

Exercise: Feature exploration

Use dimensionality reduction to elaborate features that might allow round and elongated objects

The image shows a Jupyter Notebook interface with several open cells and their outputs.

- Cell 2:** Displays a scatter plot of colored blobs (blobs.tif) on a black background. The blobs are various colors (red, green, blue, yellow, purple) and have different shapes (round, elongated).
- Cell 16:** Displays a correlation matrix heatmap titled "Correlation statistics". The matrix shows correlations between parameters: label, area, bbox_area, equivalent_diameter, convex_area, max_intensity, and mean_intensity. The diagonal is 1.000000. Correlations are generally high between area, bbox_area, and equivalent_diameter.
- Cell 16:** Displays a UMAP plot titled "TERACTION". It shows a 2D embedding of cell shapes. A lasso annotation is drawn around a cluster of orange and yellow elongated shapes. A tooltip for "lasso(annotation)" provides instructions: "pari clusters plotter using the menu Tools > Visualization > Plot measurements can use a lasso-annotation to identify the cluster that corresponds to the 8-shaped objects. After notebook cells once you have a visualization that is similar to the screen shown below."
- Cell 1:** Displays a UMAP plot titled "Exercise". The caption asks: "The UMAP-generation above is done without parameters such as centroid and orientation. Why?"
- Cell 1:** Displays a UMAP plot titled "Exercise". The caption asks: "Repeat the procedure above with the dataset human_mitosis. Identify parameters for differentiating the small bright cells from the others. (hint)"
- Cell 1:** Contains the code:

```
image = human_mitosis()  
stackview.insight(image)
```

Pixel classification / object segmentation

Use Napari to segment objects

Interactive pixel classification and object segmentation in Napari

In this exercise we will train a [Random Forest Classifier](#) for pixel classification and convert the result in an instance segmentation. We will use the napari plugin [napari-accelerated-pixel-and-object-classification](#).

Getting started

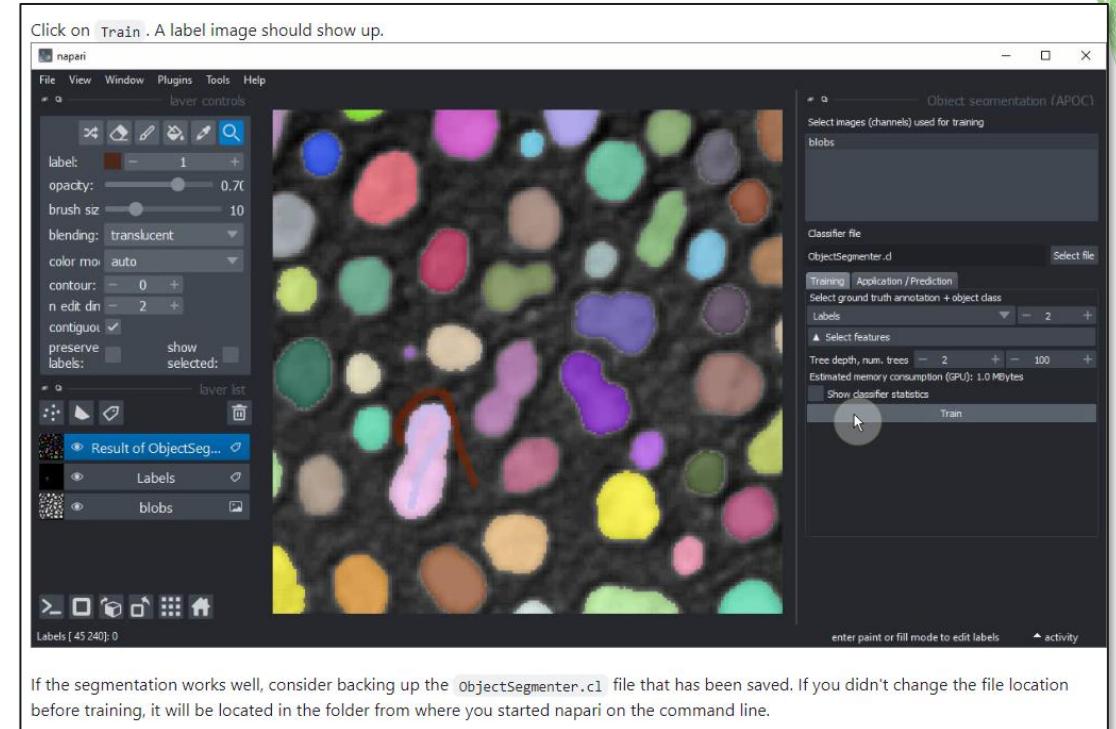
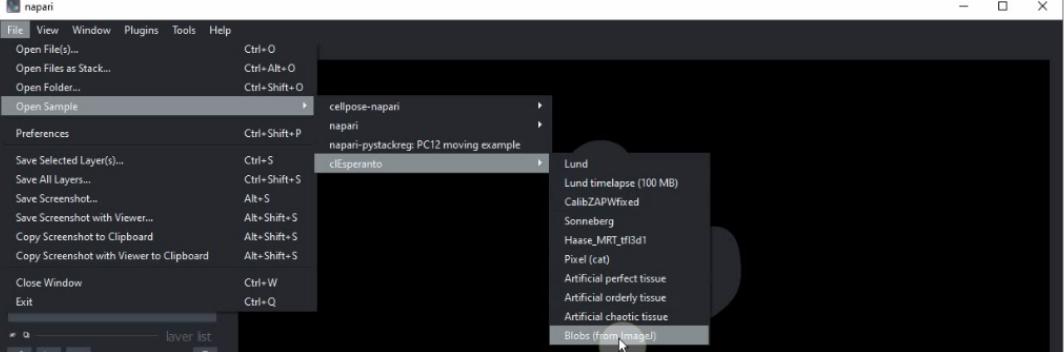
Open a terminal window and activate your conda environment:

```
conda activate devbio-napari-env
```

Afterwards, start up Napari:

```
napari
```

Load the "Blobs" example dataset from the menu File > Open Sample > c1Esperanto > Blobs (from ImageJ)



Object classification

Use Napari to classify round and elongated objects

Interactive object classification in Napari

In this exercise we will train a [Random Forest Classifiers](#) for classifying segmented objects. We will use the napari plugin [napari-accelerated-pixel-and-object-classification](#).

Getting started

Open a terminal window and activate your conda environment:

```
conda activate devbio-napari-env
```

Afterwards, start up Napari:

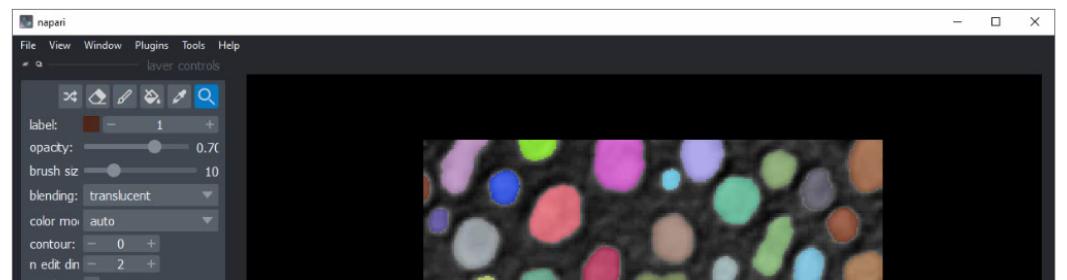
```
napari
```

Load the "Blobs" example dataset from the menu [File > Open Sample > c1Esperanto > Blobs \(from ImageJ\)](#)

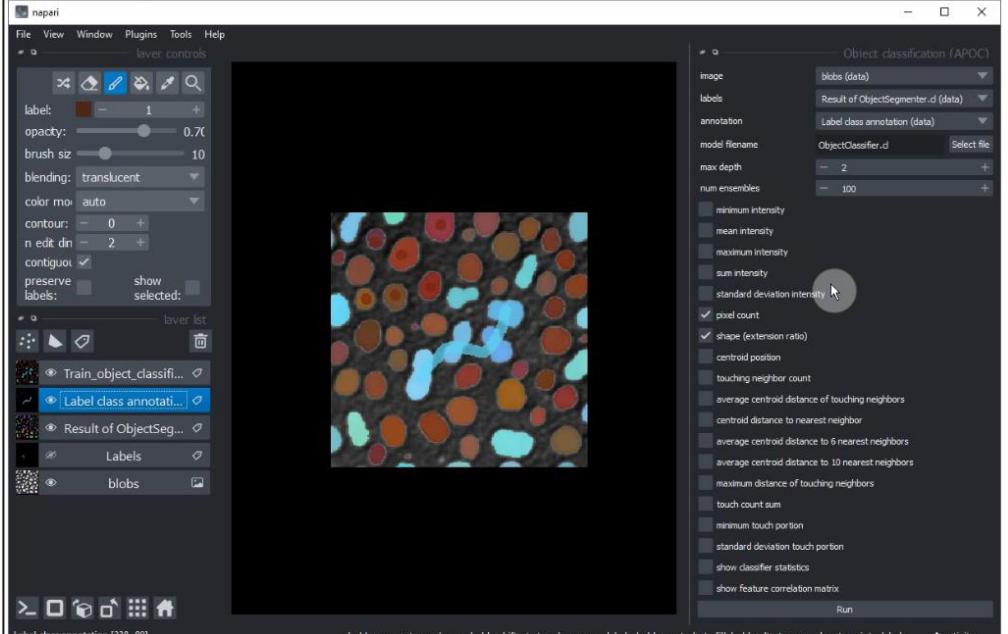
We furthermore need a label image. You can create it using the pixel classifier trained earlier or using the menu [Tools > Segmentation / labeling > Gauss-Otsu Labeling \(clesperanto\)](#).

Object classification

Our starting point is a loaded image and a label image with segmented objects. The following procedure is also shown in [this video](#).



Train the classifier again.



If you are happy with the trained classifier, copy the file to a safe place. When training the next classifier this one might be overwritten.

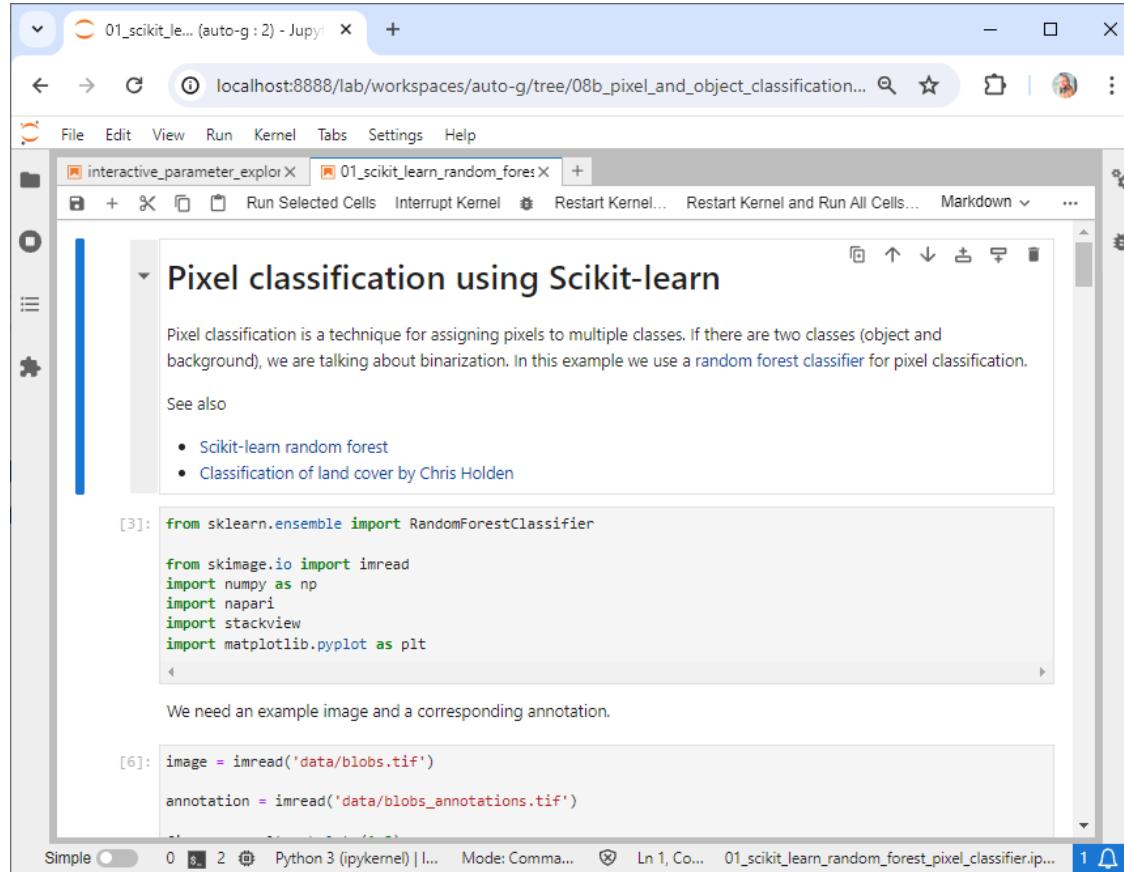
Extra exercise

Retrain the classifier so that it can differentiate three different classes:

- Small round objects
- Large round objects
- Large elongated objects

Supervised machine learning using Python

Use scikit-learn and napari in Jupyter Notebooks to train and apply Random Forest Classifiers



Pixel classification using Scikit-learn

Pixel classification is a technique for assigning pixels to multiple classes. If there are two classes (object and background), we are talking about binarization. In this example we use a random forest classifier for pixel classification.

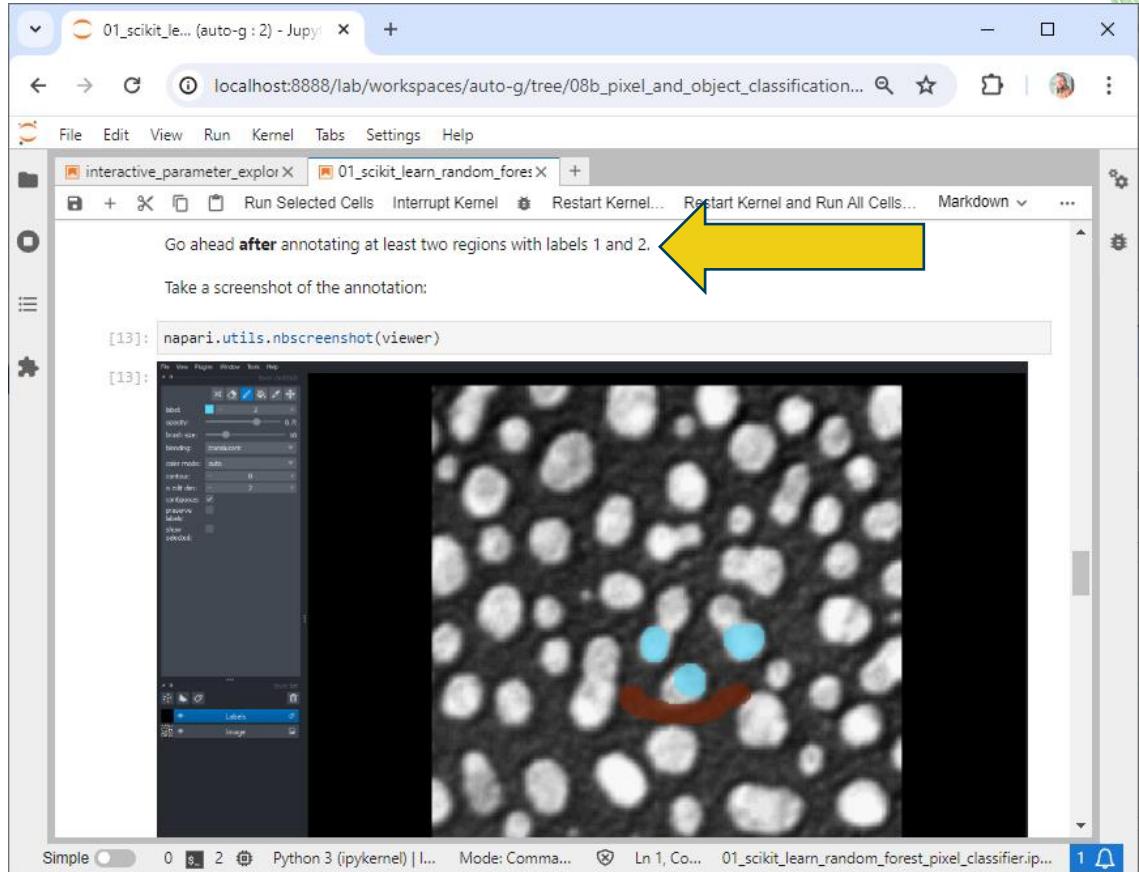
See also

- Scikit-learn random forest
- Classification of land cover by Chris Holden

```
[3]: from sklearn.ensemble import RandomForestClassifier  
  
from skimage.io import imread  
import numpy as np  
import napari  
import stackview  
import matplotlib.pyplot as plt
```

We need an example image and a corresponding annotation.

```
[6]: image = imread('data/blobs.tif')  
  
annotation = imread('data/blobs_annotations.tif')
```



Go ahead after annotating at least two regions with labels 1 and 2.

Take a screenshot of the annotation:

```
[13]: napari.utils.nbscreenshot(viewer)
```

```
[13]:
```

The napari interface shows a grayscale image of blobs with two regions annotated: one blue and one red.

Configuring Random Forest Classifiers

The image shows three side-by-side Jupyter Notebook interfaces, each with a title bar reading "05_configuri... (auto-g : 4) - Jupyter Notebook".

Left Notebook: The title is "Classifier statistics". It contains the following text:
After training, we can print out some statistics from the classifier. It gives us a table of used features and how important the features were for making the pixel classification decision.

```
[6]: shares, counts = classifier.statistics()  
  
def colorize(styler):  
    styler.background_gradient(axis=None, cmap="PiYG")  
    return styler  
  
df = pd.DataFrame(shares).T  
df.style.pipe(colorize)
```

Below the code is a table:

| | 0 | 1 | 2 | 3 | 4 |
|---------------------------------|----------|----------|----------|----------|----------|
| original | 0.138000 | 0.046423 | 0.042312 | 0.037281 | 0.062112 |
| gaussian_blur=1 | 0.228000 | 0.092846 | 0.074303 | 0.105263 | 0.055901 |
| difference_of_gaussian=1 | 0.000000 | 0.108828 | 0.095975 | 0.074561 | 0.086957 |
| laplace_box_of_gaussian_blur=1 | 0.000000 | 0.105784 | 0.089783 | 0.081140 | 0.099379 |
| gaussian_blur=5 | 0.096000 | 0.064688 | 0.118679 | 0.096491 | 0.130435 |
| difference_of_gaussian=5 | 0.254000 | 0.182648 | 0.112487 | 0.120614 | 0.118012 |
| laplace_box_of_gaussian_blur=5 | 0.209000 | 0.194064 | 0.121775 | 0.118421 | 0.124224 |
| gaussian_blur=25 | 0.004000 | 0.061644 | 0.113519 | 0.127193 | 0.080745 |
| difference_of_gaussian=25 | 0.031000 | 0.072298 | 0.122807 | 0.127193 | 0.130435 |
| laplace_box_of_gaussian_blur=25 | 0.040000 | 0.070776 | 0.108359 | 0.111842 | 0.111801 |

Middle Notebook: The title is "Classifier statistics". It contains the following text:
The new classifier still produces a very similar result. It takes less features into account, which makes it faster, but potentially also less robust again differences between images and imaging conditions. We just take another look at the classifier statistics:

```
[8]: shares, counts = classifier.statistics()  
df = pd.DataFrame(shares).T  
df.style.pipe(colorize)
```

Below the code is a table:

| | 0 | 1 | 2 |
|--------------------------------|----------|----------|----------|
| gaussian_blur=1 | 0.331000 | 0.349194 | 0.344620 |
| difference_of_gaussian=5 | 0.356000 | 0.329839 | 0.337096 |
| laplace_box_of_gaussian_blur=5 | 0.313000 | 0.320968 | 0.318284 |

Right Notebook: The title is "Classifier statistics". It contains the following text:
cl_filename = 'data/blobs_object_segmenter_3.cl'

apoc.erase_classifier(cl_filename)
classifier = apoc.ObjectSegmenter(opencl_filename=cl_filename,
positive_class_identifier=2,
max_depth=3,
num_ensembles=1000)

classifier.train(features, manual_annotation, image)

segmentation_result = classifier.predict(features=features, image=image)
stackview.imshow(segmentation_result, labels=True)