

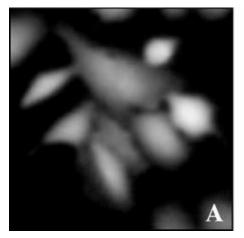
Introduction to Instance Segmentation

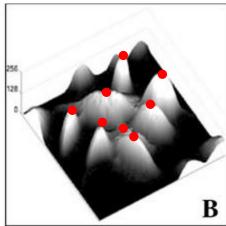
Beth Cimini, PhD Associate Director, Bioimage Analysis Imaging Platform, Broad Institute of MIT and Harvard

Slides available at broad.io/InstanceDL_2024

Instance segmentation

We next need to distinguish multiple objects contained in the same "clump"





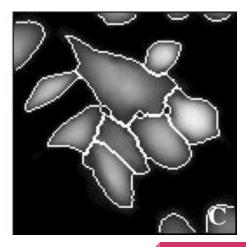


Image Processing Modules

- All classical image analysis segmentation works off of the principle- "my image should have my objects of interest bright and everywhere else dark"
- For segmentation purposes, basically any transformation you can think of to achieve this is "legal"
 - <u>Emphatically</u>, this does not apply to images you are actively measuring, just images to be fed into a segmentation
- To achieve this, you may need to enhance dim signal that really is there and/or mask out bright signal that truly is (but is not supposed to be segmented at the current time)
- The more transformations you do before segmentation, the more critical it is to check your segmentations against your raw images

Classical Image Processing Options

(Not exhaustive, just inspirational)

- EnhanceEdges
- EnhanceOrSuppressFeatures- works on tubes, speckles, and/or dark holes
- Smooth
- Crop
- ColorToGray / GrayToColor

- UnmixColors
- GaussianFilter
- MedialAxis
- RemoveHoles
- MatchTemplate

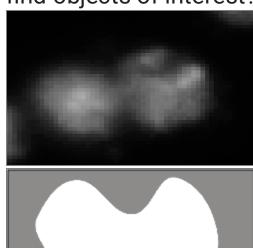
PLUS, classical (such as Random Forest based) semantic segmenters

Object identification

Once the images are loaded, how do you find objects of interest?

 Step 1: Distinguish the foreground from the background (thresholding)

 Step 2: Split/merge "objects" properly

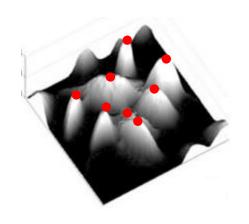


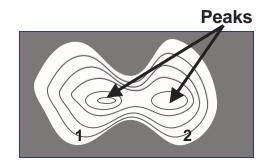


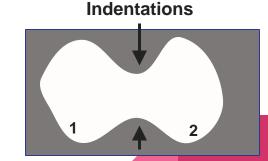
Separating touching objects – clump identification

Intensity: Works best if objects are brighter at center, dimmer at edges

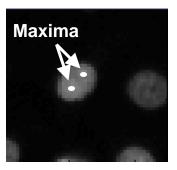
Shape: Works best if objects have indentations where clumps touch (esp. if objects are round)







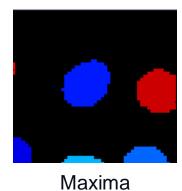
Separating touching objects – clump identification



Original image



distance = 4



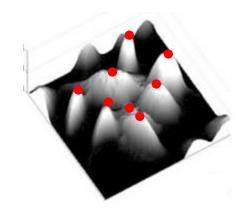
distance = 8

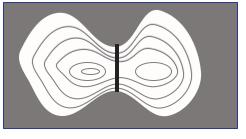
- . Suppress Local Maxima
- . Smallest distance allowed between object intensity peaks to be considered one object rather than a clump
- . Decrease to reduce improper merging of objects in clumps

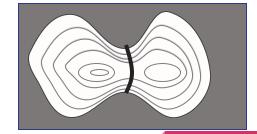
Separating touching objects

Drawing boundaries: Two options

- Shape: Drawsboundary lines at the indentations
- boundary lines at dimmest line between objects







So to sum up, here's how classical instance segmentation works

- Optionally, do SOMETHING to make your image "everything I care about is bright, and everything else is dark"
- Decide on a threshold for that image
- Then either
 - Say every pixel that touches its neighbors after thresholding is part of the same object (not shown above)
- OR
 - Figure out the centers (and thereby also, the count) of objects you have, based on some rational principle
 - Follow some rational principle to trace from the centers to the edges

Sorry, excuse me Beth, this is a deep learning course, why are you telling us irrelevant stuff?

Sorry, my bad, here's how deep learning instance segmentation (typically) works

- Optionally, do SOMETHING to make your image "everything I care about is bright, and everything else is dark"
- Decide on a threshold for that image
- Also, either
 - Say every pixel that touches its neighbors after thresholding is part of the same object (not shown above)
- OR (one or both, in either order)
 - Figure out the approximate locations (centers and/or bounding boxes) of objects you have,
 based on some rational principle
 - Follow some rational principle to pull pixels together

Deep learning for object locations

- Optionally, do SOMETHING to make your image "everything I care about is bright, and everything else is dark"
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DL for approximate location finding – Region Proposal Networks (RPNs)

- Take in an image, put out bounding boxes of likely objects
- Used in things like Faster R-CNN (object detection) Mask R-CNN
- Can share convolutional layers with downstream tasks, reducing extra time required

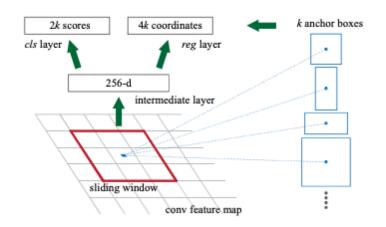




Figure 1: Different schemes for addressing multiple scales and sizes. (a) Pyramids of images and feature maps are built, and the classifier is run at all scales. (b) Pyramids of filters with multiple scales/sizes are run on the feature map. (c) We use pyramids of reference boxes in the regression functions.

Mask-R-CNN

- Supports overlapping objects
- RPN (gets class and box), PLUS n-classes binary semantic segmenters

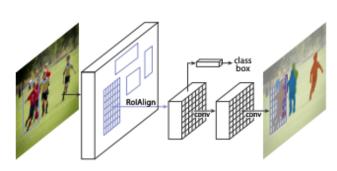
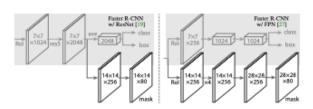
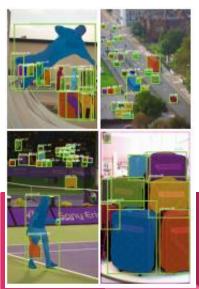


Figure 1. The Mask R-CNN framework for instance segmentation.





From each pixel, a distance polygon - StarDist

- Predict for each pixel how far it is from the edge of its object in each of 32 directions, making n-pixels polygons; also predict object probability
- Keep the highest-probability polygons
 - Later versions add some "majorityrules" shape refinement, to make the final thing less polygonal

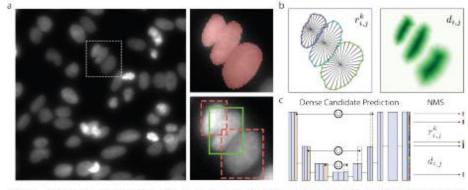
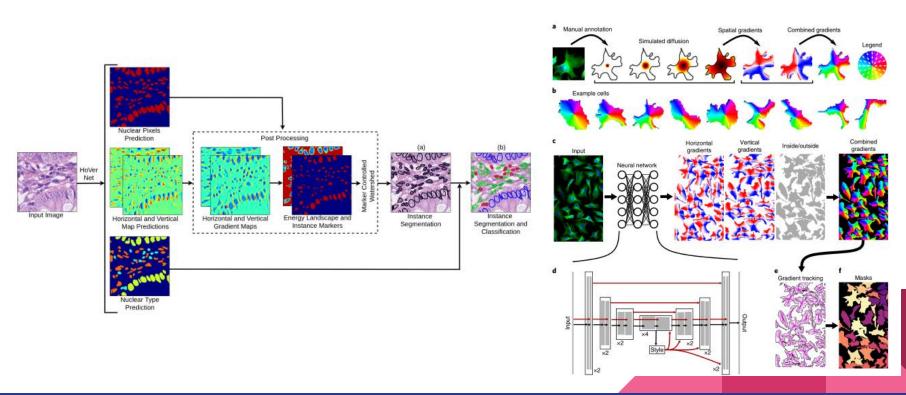


Fig. 1: (a) Potential segmentation errors for images with crowded nuclei: Merging of touching cells (upper right) or suppression of valid cell instances due to large overlap of bounding box localization (lower right). (b) The proposed StarDist method predicts object probabilities $d_{i,j}$ and star-convex polygons parameterized by the radial distances $r_{i,j}^k$. (c) We densely predict $r_{i,j}^k$ and $d_{i,j}$ using a simple U-Net architecture [15] and then select the final instances via non-maximum suppression (NMS).

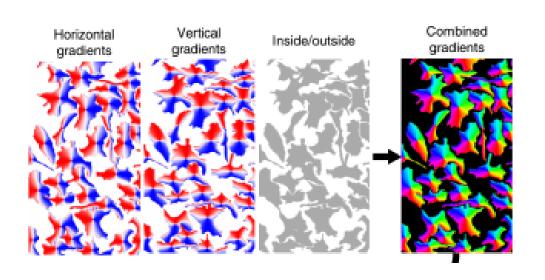
Deep learning for together-ness

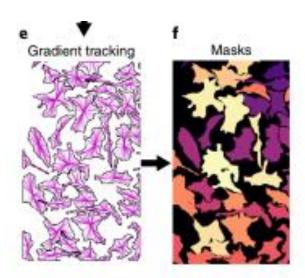
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Gradient flows - Cellpose



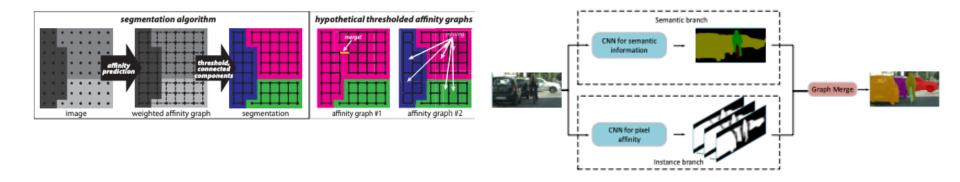
Gradient flows - Cellpose





What belongs together – affinity learning

Figure out what belongs with what, and group it together



What belongs together – affinity learning

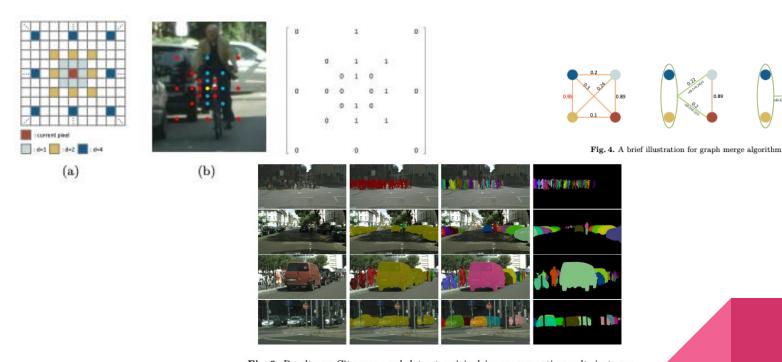
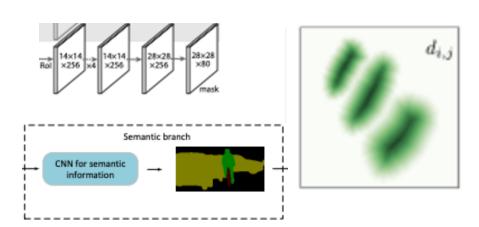


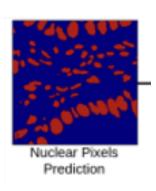
Fig. 6. Results on Cityscapes val dataset, original image, semantic result, instance result and ground truth from left to right. Results in last two rows are cropped from original ones for better visualization.

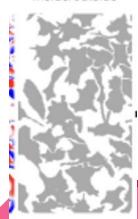
Deep learning for probability

- Optionally, do SOMETHING to make your image "everything I care about is bright, and everything else is dark"
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- OR (one or both, in either order)
 - Figure out the approximate locations (centers and/or bounding boxes) of objects you have, based on some rational principle
 - Follow some rational principle to pull pixels together

We've already seen this in the other things we've looked at too – can be combined with more or less fancy post-processing – typically (though not always) 2-class "inside the object" probabilities

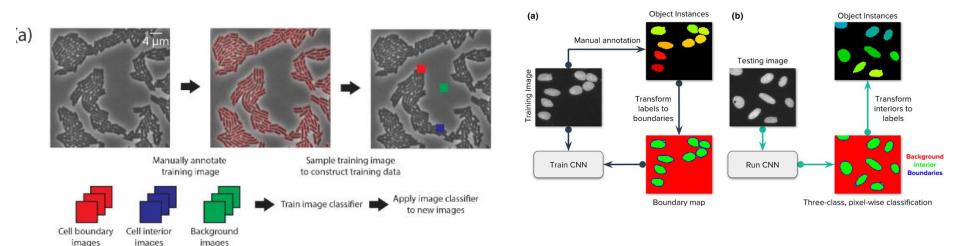




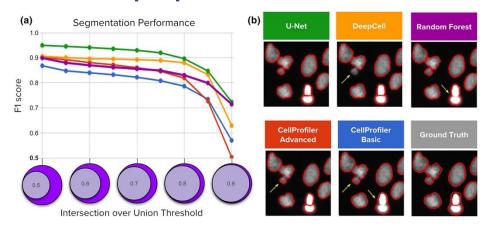


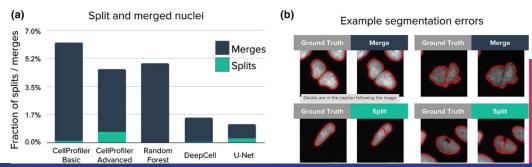
Inside/outside

Semantic models based on 3 classes (cell boundary, interior, background) are also popular choices

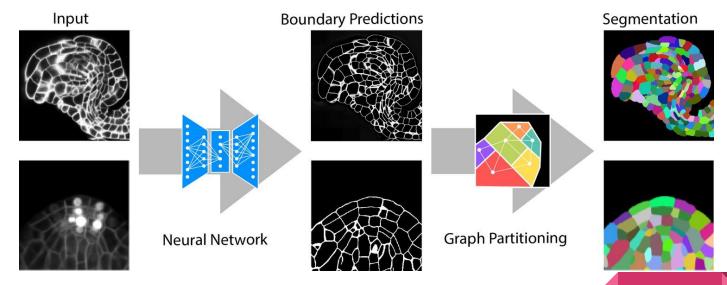


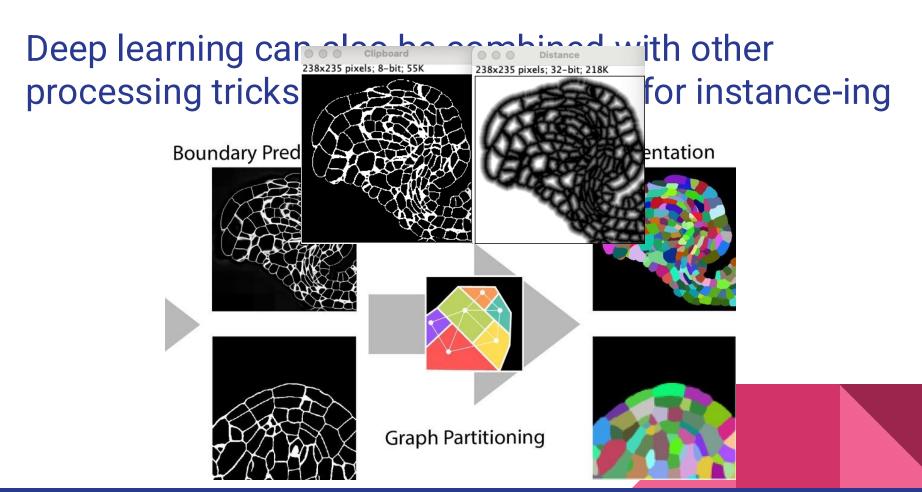
Models based on 3 classes (cell boundary, interior, background) are also popular choices





Deep learning can also be combined with other processing tricks to get the right input for instance-ing





So it sounds like I have a lot of options on how to put deep learning into my instance segmentation. Too many options maybe?

Tips for creating a good high content analysis workflow

- When finding the objects that you care about, ask yourself for your whole experiment:
 - Do I generally agree with most of the object segmentations from my analysis workflow?
 - Do I have an approximately equal number of regions/images where the threshold chosen by the algorithm for this image is a bit too low vs a bit too high?
 - Do I have an approximately equal number of oversegmentations/splits and undersegmentations/merges?
 - Very important: Do both the second and third bullet points hold true for both my negative control images and my positive control (or most extreme expected phenotype(s) sample) images?

https://carpenter-singh-lab.broadinstitute.org/blog/when-to-say-good-enough

Based on that and other things, here are some simple considerations

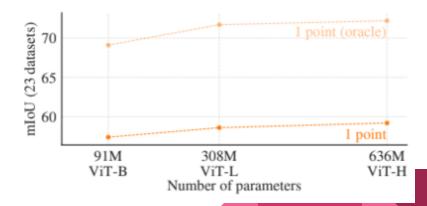
Think about... how much compute do you need (and how much do you have)?

 The more types of DL, and the presence of certain types of DL, the more parameters you'll have to tune

U-Net (2 class): We use the popular U-Net architecture [15] as a baseline to predict 2 output classes (cell, background). We use 3 down/up-sampling blocks, each consisting of 2 convolutional layers with $32 \cdot 2^k (k=0,1,2)$ filters of size 3×3 (approx. 1.4 million parameters in total). We apply a threshold σ on the cell probability map and retain the connected components as final result (σ is optimized on the validation set for every dataset).

U-Net (3 class): Like U-Net (2 class), but we additionally predict the boundary pixels of cells as an extra class. The purpose of this is to differentiate crowded cells with touching borders (similar to [4,5]). We again use the connected components of the thresholded cell class as final result.

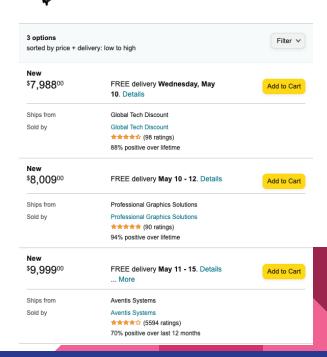
Mask R-CNN: A state-of-the-art instance segmentation method combining a bounding-box based region proposal network, non-maximum-suppression (NMS), and a final mask segmentation (approx. 45 million parameters in total). We use a popular open-source implementation⁶. For each dataset, we perform a gridsearch over common hyper-parameters, such as detection NMS threshold, region proposal NMS threshold, and number of anchors.



Think about... how much compute do you need (and how much do you have)? NVIDIA Tesla A100 Ampere 40 GB Graphics Card - PCle...

Segment Anything



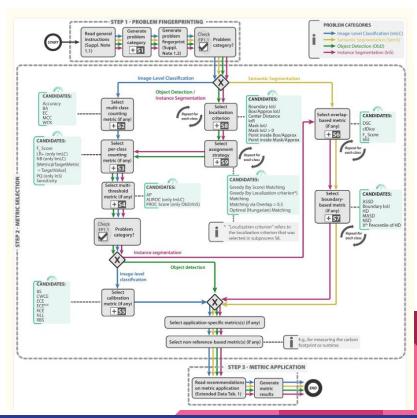


** 2 ratings

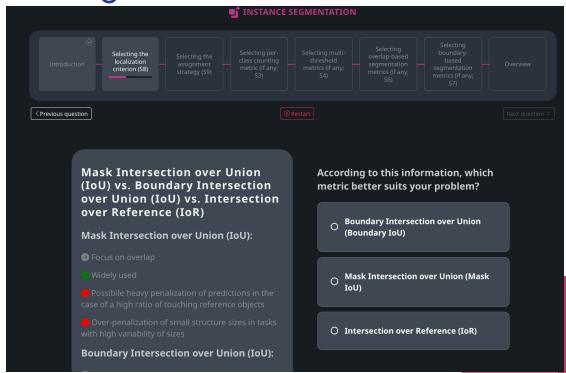
Think about... how will you actually determine if the

model is working?

 What is the right way to assess accuracy? Which parts of assessing accuracy are fast enough to incorporate into your loss function? Can you train your model in parts?



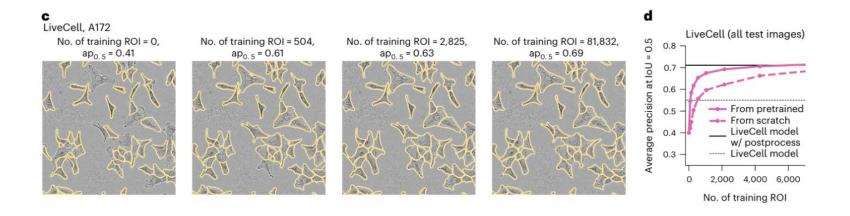
Think about... how will you actually determine if the model is working?



Think about... where will I get my data?

- Certain kinds of models (especially transformers) need a LOT of data
- Some sets:
 - CytolmageNet
 - o Broad Bioimage Benchmark Collection
 - Cell Painting Gallery
 - Cell Tracking Challenge
 - Image Data Resource
 - Human Protein Atlas
 - Open Cell
 - Bioimage Archive
- But all still dwarfed by natural images (we never made cell Facebook, darn it)
 - Pretraining!

Think about... how can I make it easy to get more data if I need to?

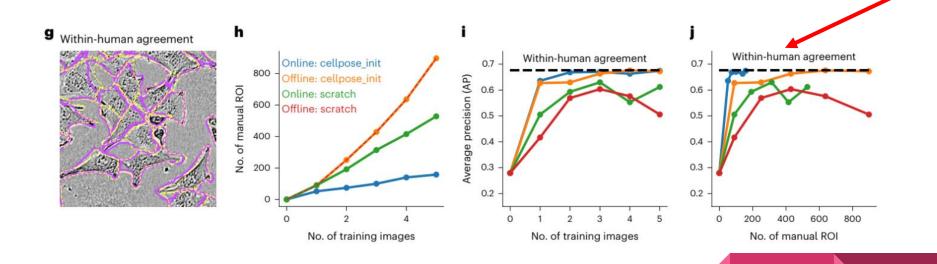


Think about... how can I make it easy to get more data if I need to?

Training a new LiveCell model

Manual ROI 52/127 22/173 26/200 40/241 18/293

Think about... how can I make it easy to get more data if I need to?



The world's best instance segmenter is still - but maybe not for much longer!





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