

Lecture 14b: Beyond classification – segmentation and autoencoders

Machine Learning and Imaging

BME 590L
Roarke Horstmeyer

Class project option – work with a new image dataset from Kenya!



- Collaboration with Dr. Wendy Prudhomme-O'Meara at the Duke Global Health Institute
- Certain species of mosquitos carry the malaria parasite
- Classification of different mosquitos into different species at different locations/villages can offer some really useful information!

Dataset:

- **4 species imaged:** 1710 images identified as *gambiae*, 402 as unidentified, 107 images as *funestus* but **only 17** as *demeilloni*
- Each species imaged 4 times from 4 directions
- In one of 4 states: unfed, blood-fed, gravid, half gravid

Task: Prep dataset, develop a classification (or other) network to establish some preliminary findings

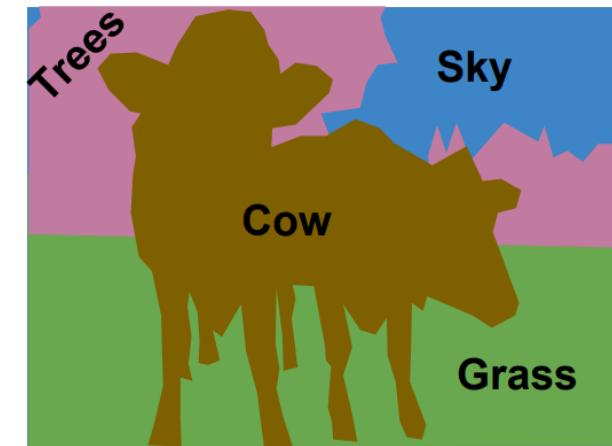
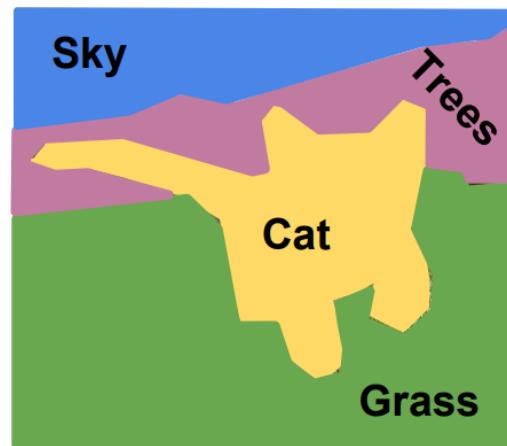
Semantic Segmentation

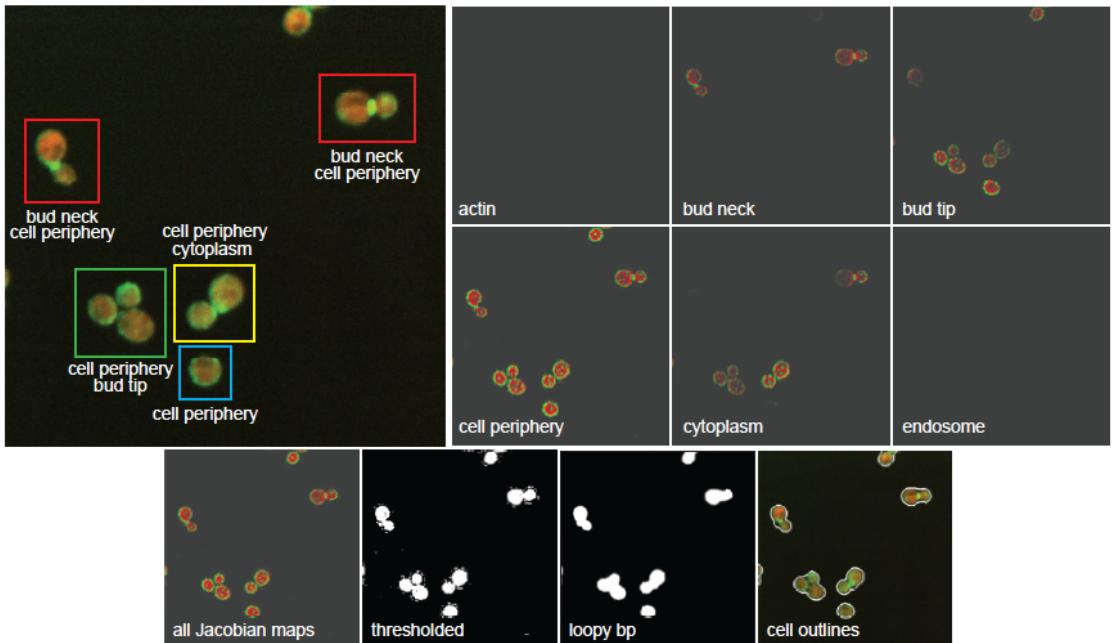
Label each pixel in the image with a category label

Don't differentiate instances, only care about pixels

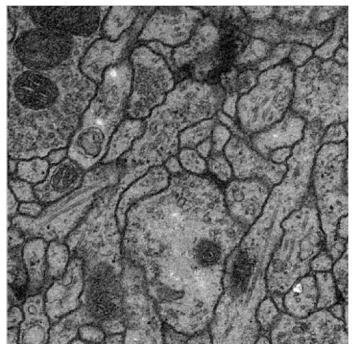


This image is CC0 public domain





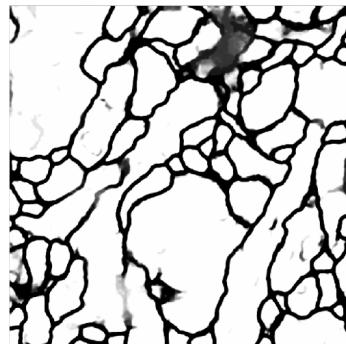
Oren Z. Kraus et al., “Classifying and Segmenting Microscopy Images Using Convolutional Multiple Instance Learning,” arXiv 2015



(a) Input image

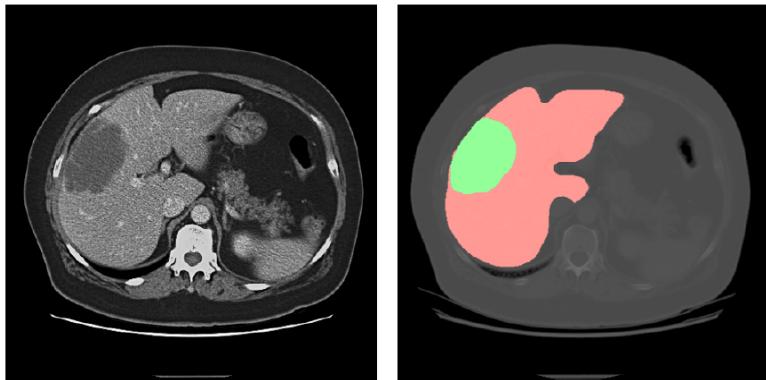
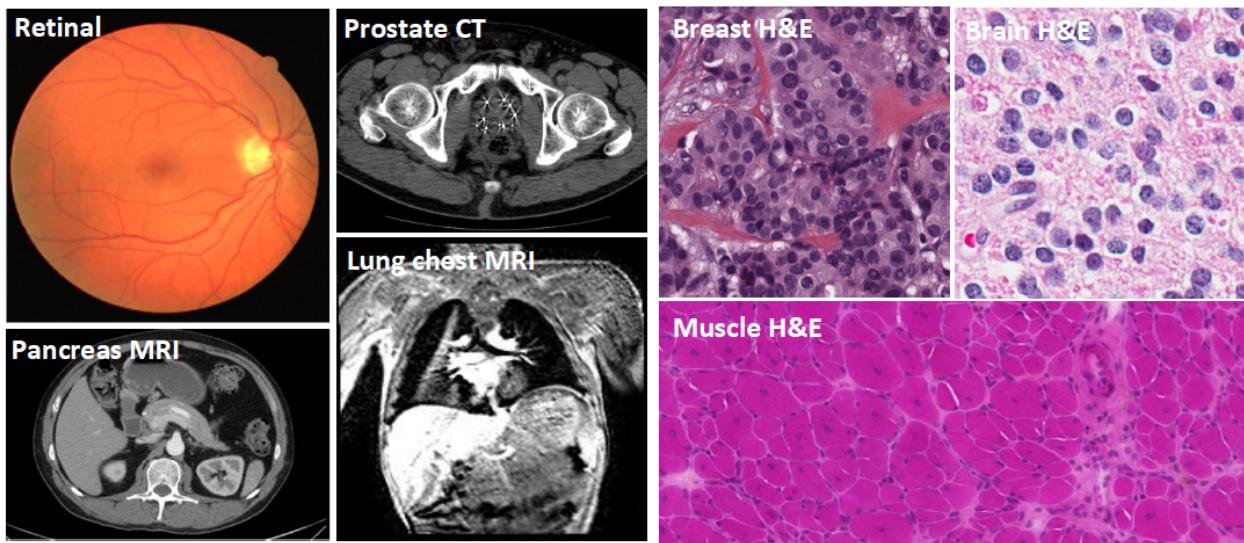


(b) FC-ResNet with dropout at test time [17]



(c) Segmentation result of our pipeline

Other possible examples:

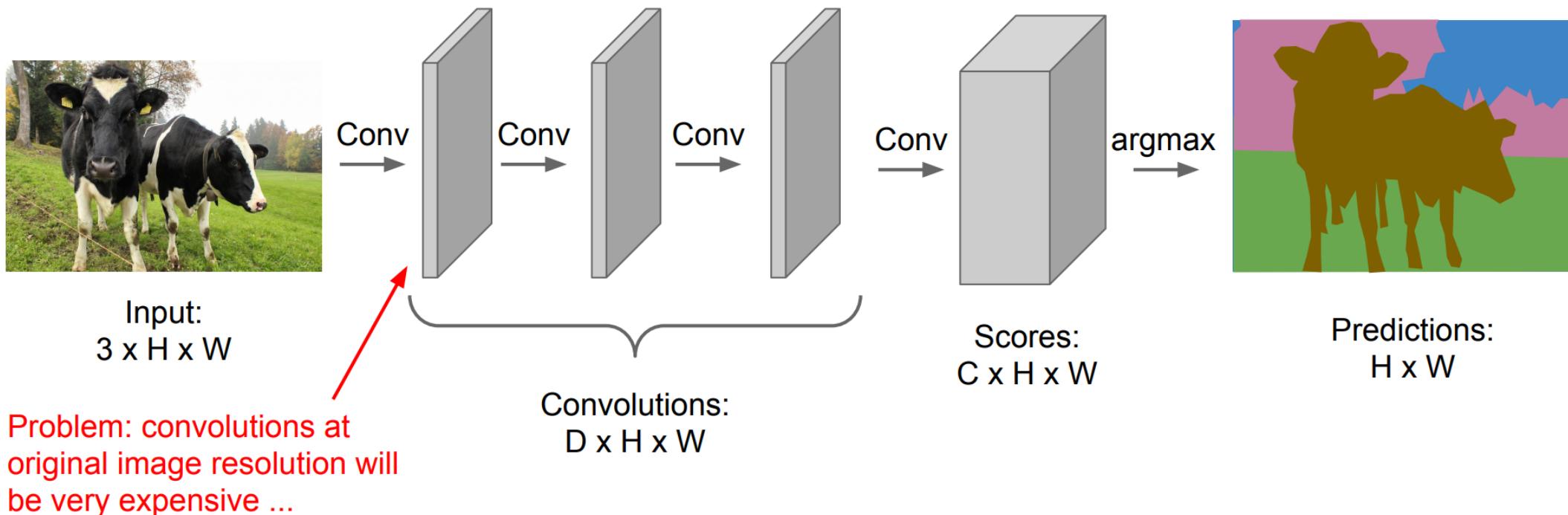


M. Drozdzal et al., Learning Normalized Inputs for Iterative Estimation in Medical Image Segmentation (2017)

Z. Zhang et al., Recent Advances in the Applications of Convolutional Neural Networks to Medical Image Contour Detection (2017)

Semantic Segmentation Idea: Fully Convolutional ?

Design a network as a bunch of convolutional layers
to make predictions for pixels all at once!



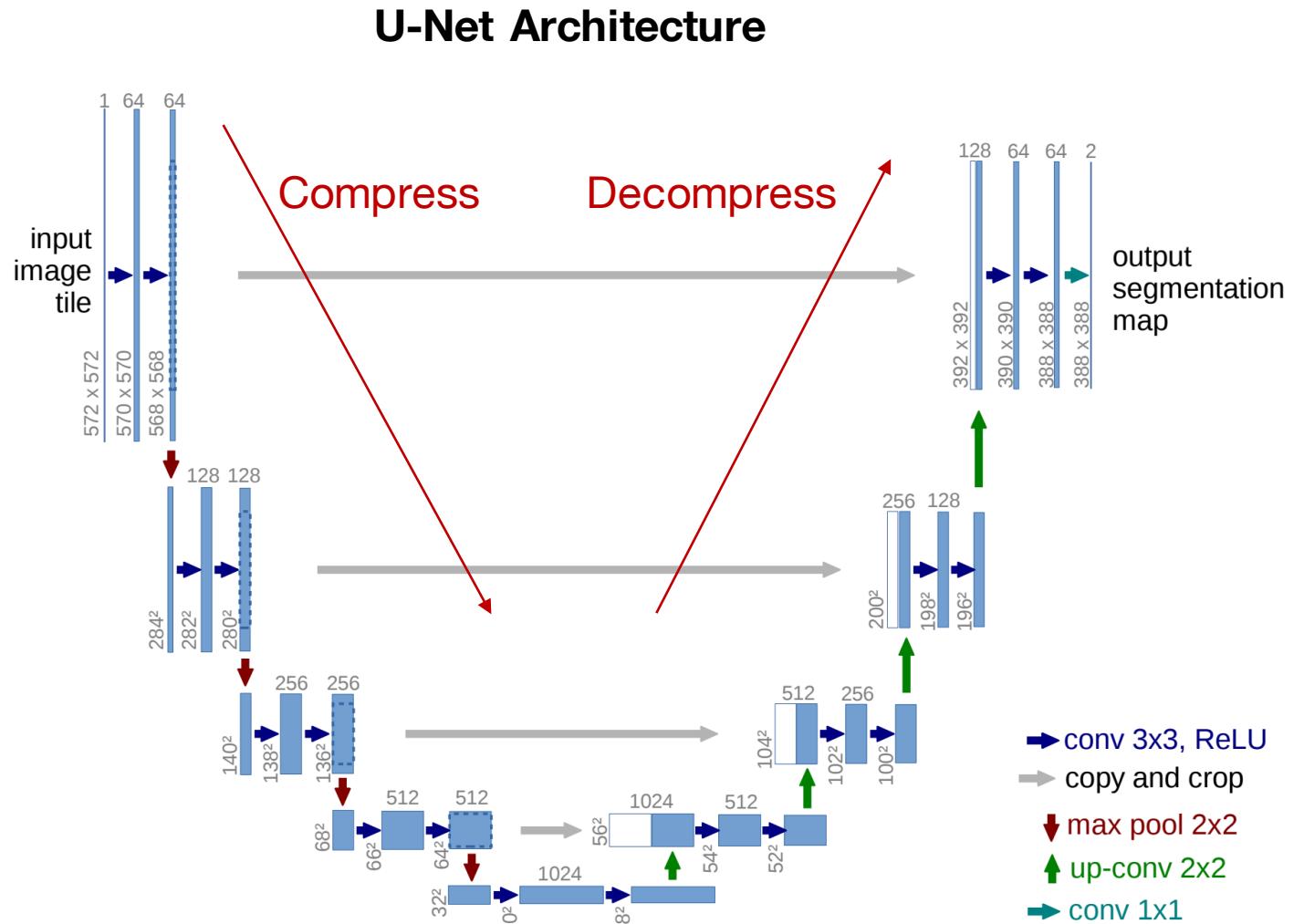
Instead, compress x-y dimensions of input image

- Compress spatial features into learned filters
- Then, decompress learned filters back into same spatial dimensions

U-Net: Convolutional Networks for Biomedical Image Segmentation

Olaf Ronneberger, Philipp Fischer, and Thomas Brox

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WWW home page: <http://lmb.informatik.uni-freiburg.de/>



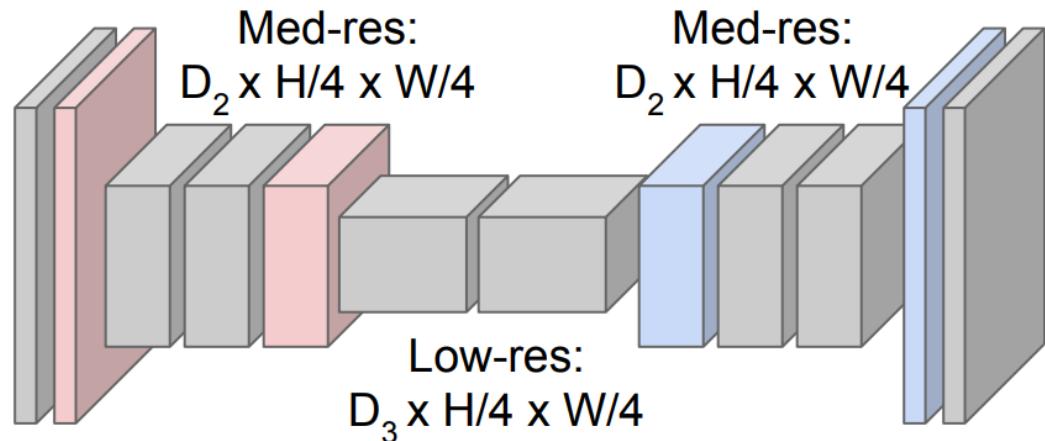
Semantic Segmentation Idea: Fully Convolutional

Downsampling:
Pooling, strided
convolution



Input:
 $3 \times H \times W$

Design network as a bunch of convolutional layers, with
downsampling and **upsampling** inside the network!



Upsampling:
???



Predictions:
 $H \times W$

Long, Shelhamer, and Darrell, "Fully Convolutional Networks for Semantic Segmentation", CVPR 2015

Noh et al, "Learning Deconvolution Network for Semantic Segmentation", ICCV 2015

In-Network upsampling: “Unpooling”

Nearest Neighbor

1	2
3	4



1	1	2	2
1	1	2	2
3	3	4	4
3	3	4	4

Input: 2 x 2

Output: 4 x 4

“Bed of Nails”

1	2
3	4



1	0	2	0
0	0	0	0
3	0	4	0
0	0	0	0

Input: 2 x 2

Output: 4 x 4

In-Network upsampling: “Max Unpooling”

Max Pooling

Remember which element was max!

1	2	6	3
3	5	2	1
1	2	2	1
7	3	4	8

Input: 4 x 4

5	6
7	8

Output: 2 x 2

Max Unpooling

Use positions from pooling layer

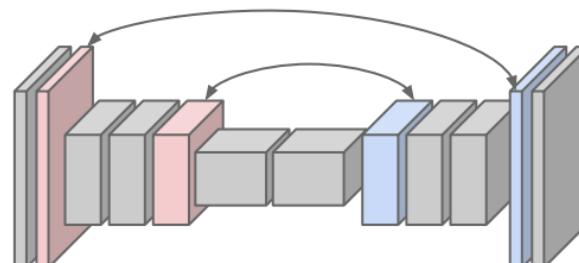
1	2
3	4

Input: 2 x 2

0	0	2	0
0	1	0	0
0	0	0	0
3	0	0	4

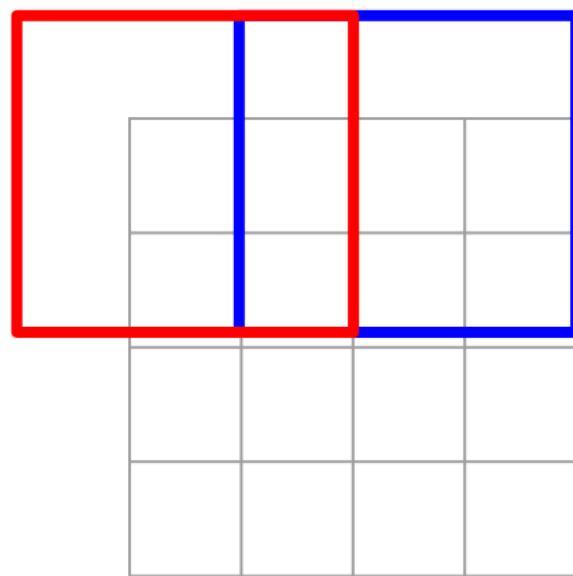
Output: 4 x 4

Corresponding pairs of
downsampling and
upsampling layers



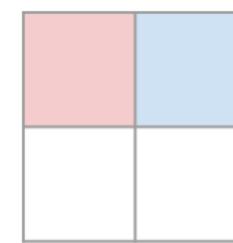
Learnable Upsampling: Transpose Convolution

Recall: Normal 3×3 convolution, stride 2 pad 1



Input: 4×4

Dot product
between filter
and input

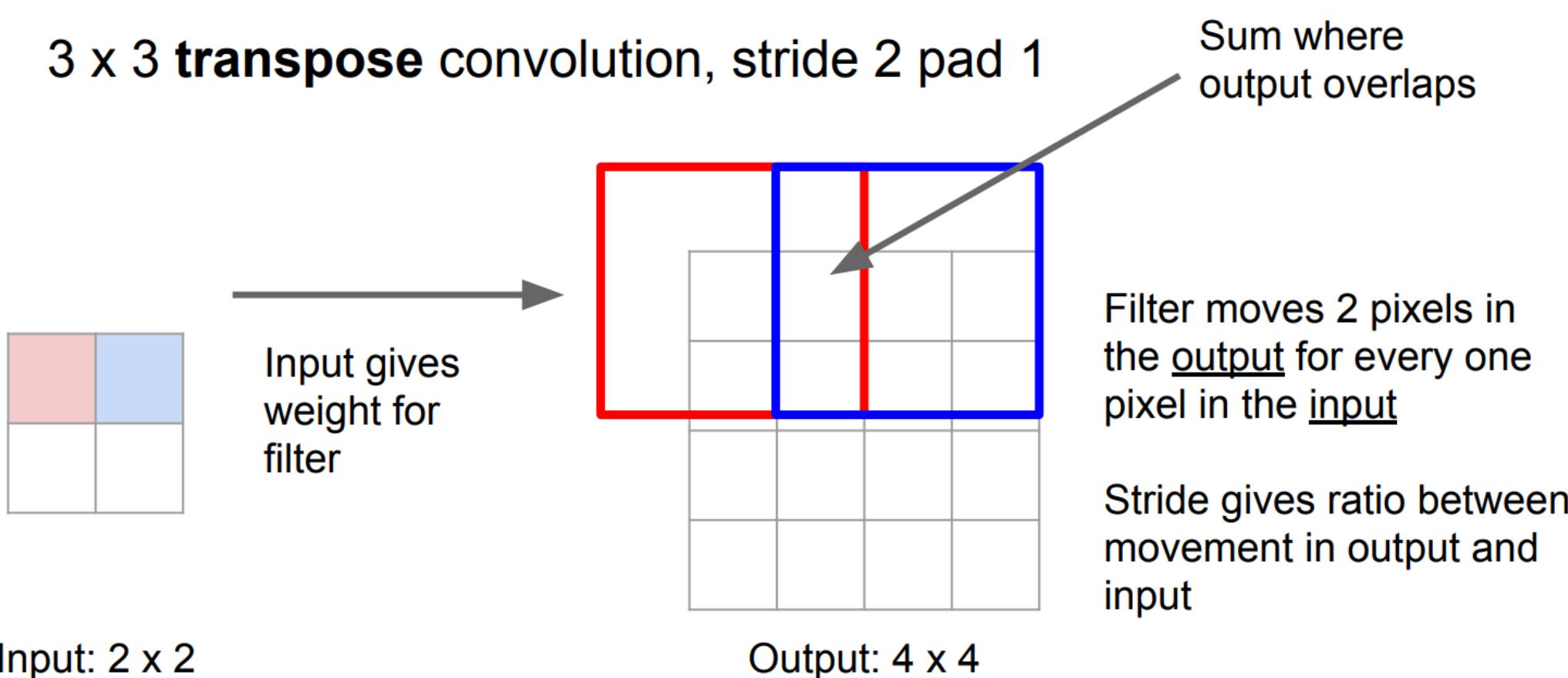


Output: 2×2

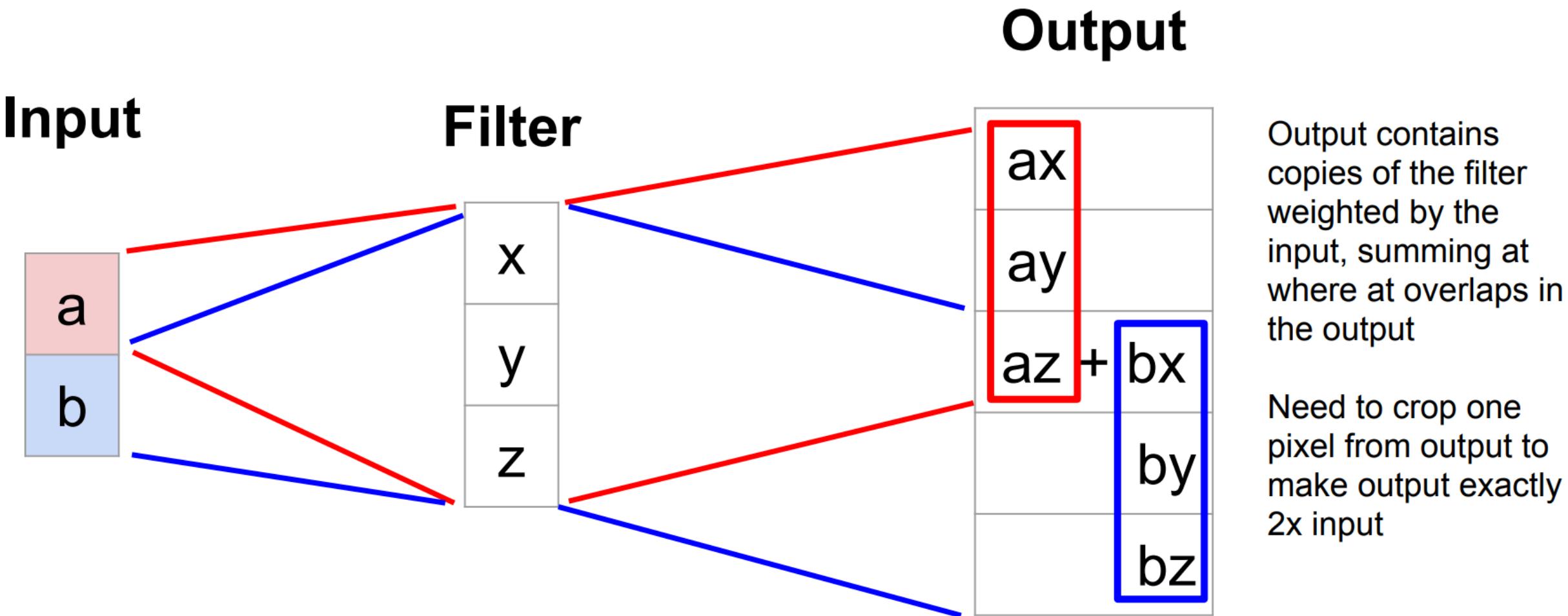
Filter moves 2 pixels in
the input for every one
pixel in the output

Stride gives ratio between
movement in input and
output

Learnable Upsampling: Transpose Convolution



Learnable Upsampling: 1D Example



Convolution as Matrix Multiplication (1D Example)

We can express convolution in terms of a matrix multiplication

$$\vec{x} * \vec{a} = X\vec{a}$$

$$\begin{bmatrix} x & y & x & 0 & 0 & 0 \\ 0 & x & y & x & 0 & 0 \\ 0 & 0 & x & y & x & 0 \\ 0 & 0 & 0 & x & y & x \end{bmatrix} \begin{bmatrix} a \\ b \\ c \\ d \\ 0 \end{bmatrix} = \begin{bmatrix} ay + bz \\ ax + by + cz \\ bx + cy + dz \\ cx + dy \end{bmatrix}$$

Example: 1D conv, kernel size=3, stride=1, padding=1

Convolution transpose multiplies by the transpose of the same matrix:

$$\vec{x} *^T \vec{a} = X^T\vec{a}$$

$$\begin{bmatrix} x & 0 & 0 & 0 \\ y & x & 0 & 0 \\ z & y & x & 0 \\ 0 & z & y & x \\ 0 & 0 & z & y \\ 0 & 0 & 0 & z \end{bmatrix} \begin{bmatrix} a \\ b \\ c \\ d \end{bmatrix} = \begin{bmatrix} ax \\ ay + bx \\ az + by + cx \\ bz + cy + dx \\ cz + dy \\ dz \end{bmatrix}$$

When stride=1, convolution transpose is just a regular convolution (with different padding rules)

Convolution as Matrix Multiplication (1D Example)

We can express convolution in terms of a matrix multiplication

$$\vec{x} * \vec{a} = X\vec{a}$$

$$\begin{bmatrix} x & y & z & 0 & 0 & 0 \\ 0 & 0 & x & y & z & 0 \end{bmatrix} \begin{bmatrix} 0 \\ a \\ b \\ c \\ d \\ 0 \end{bmatrix} = \begin{bmatrix} ay + bz \\ bx + cy + dz \end{bmatrix}$$

Example: 1D conv, kernel size=3, stride=2, padding=1

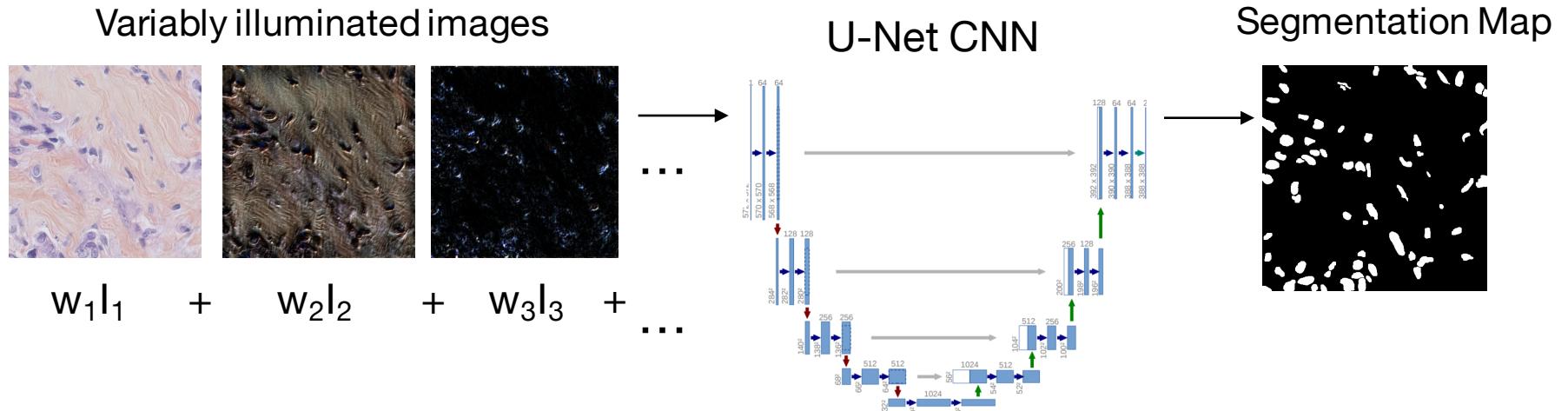
Convolution transpose multiplies by the transpose of the same matrix:

$$\vec{x} *^T \vec{a} = X^T \vec{a}$$

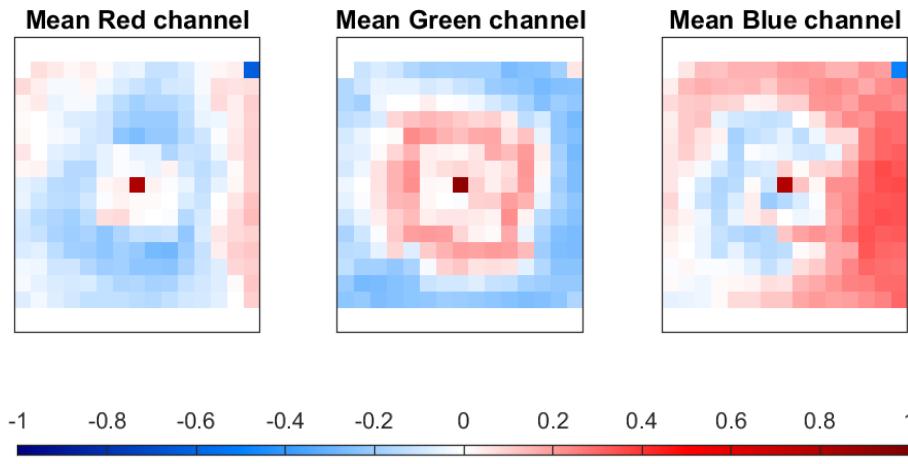
$$\begin{bmatrix} x & 0 \\ y & 0 \\ z & x \\ 0 & y \\ 0 & z \\ 0 & 0 \end{bmatrix} \begin{bmatrix} a \\ b \end{bmatrix} = \begin{bmatrix} ax \\ ay \\ az + bx \\ by \\ bz \\ 0 \end{bmatrix}$$

When stride>1, convolution transpose is no longer a normal convolution!

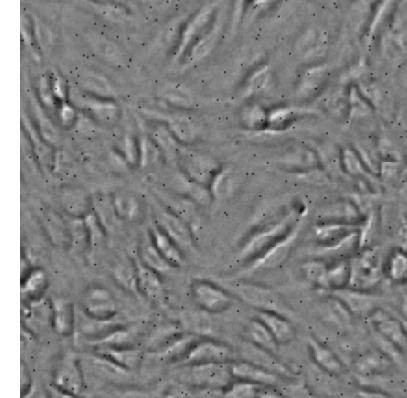
Learned sensing for improved image segmentation



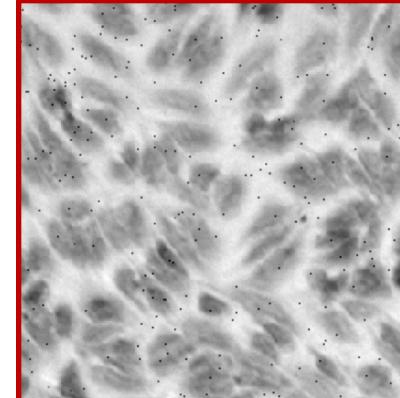
Optimized illumination for nuclei segmentation



Standard illumination



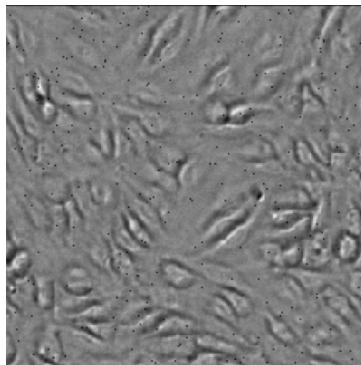
Learned illumination



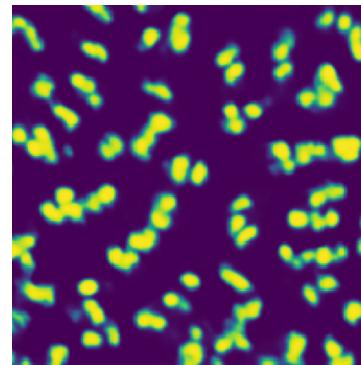
+5-10% accuracy

Image segmentation –current workflow

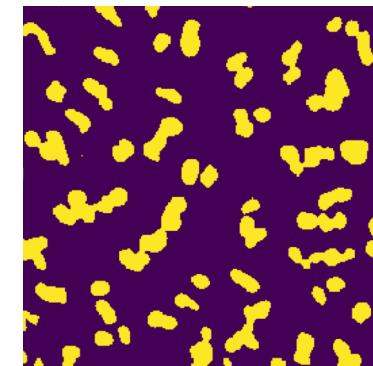
Capture: BF images



Capture: Fluorescence



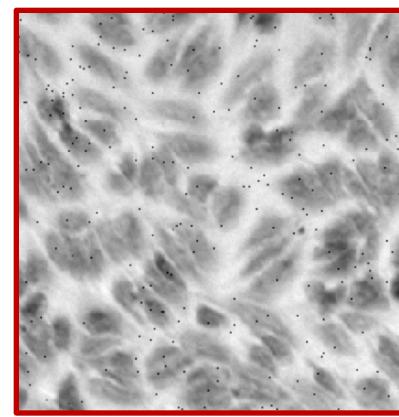
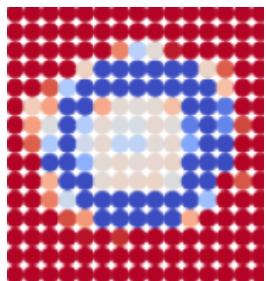
Segmentation Mask



Threshold

(e.g., DAPI-stained nuclei)

Learned
illumination



Inference via a trained U-Net

Optimally illuminated

in silico labeling: fluorescence image inference from bright-field data

In Silico Labeling: Predicting Fluorescent Labels in Unlabeled Images

Eric M. Christiansen,^{1,11,*} Samuel J. Yang,¹ D. Michael Ando,^{1,9} Ashkan Javaherian,^{2,9} Gaia Skibinski,^{2,9} Scott Lipnick,^{3,4,8,9} Elliot Mount,^{2,10} Alison O'Neil,^{3,10} Kevan Shah,^{2,10} Alicia K. Lee,^{2,10} Piyush Goyal,^{2,10} William Fedus,^{1,6,10} Ryan Poplin,^{1,10} Andre Esteva,^{1,7} Marc Berndt,¹ Lee L. Rubin,³ Philip Nelson,^{1,*} and Steven Finkbeiner^{2,5,*}

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²Taube/Koret Center for Neurodegenerative Disease Research and DaedalusBio, Gladstone Institutes, San Francisco, CA 94158, USA

³Department of Stem Cell and Regenerative Biology, Harvard University, Cambridge, MA 02138, USA

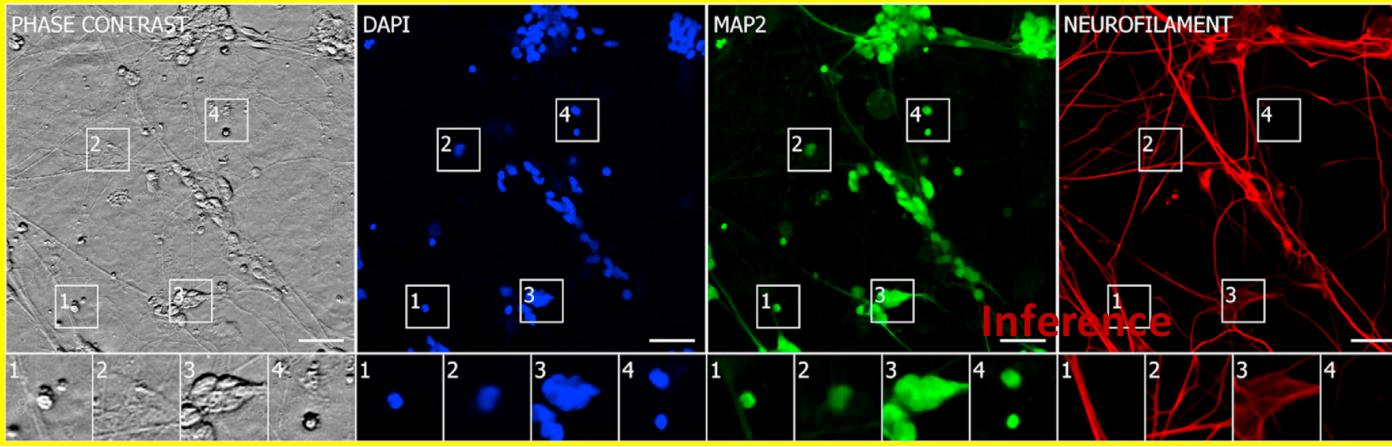
⁴Department of Biomedical Informatics, Harvard Medical School, Boston, MA 02115, USA

⁵Departments of Neurology and Physiology, University of California, San Francisco, 94158, USA

⁶Montreal Institute of Learning Algorithms, University of Montreal, Montreal, QC, Canada

⁷Department of Electrical Engineering, Stanford University, Stanford, CA 94305, USA

⁸Center for Assessment Technology and Continuous Health, Massachusetts General Hospital, Boston, MA 02114, USA

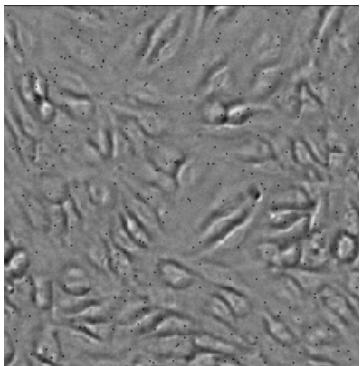


**BF Focal stack
(26+ images)**

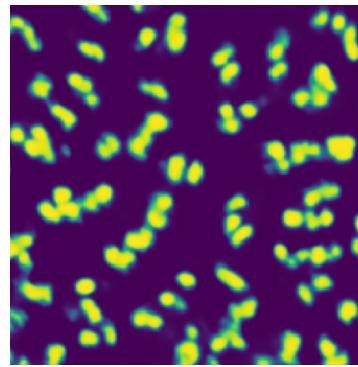
Task: bright-field to fluorescence image Inference

Image segmentation versus *in silico* labeling (fluorescence inference)

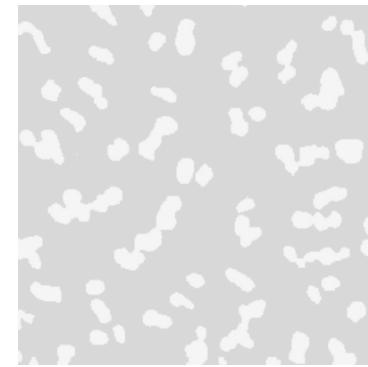
Capture: Bright field



Capture: Fluorescence



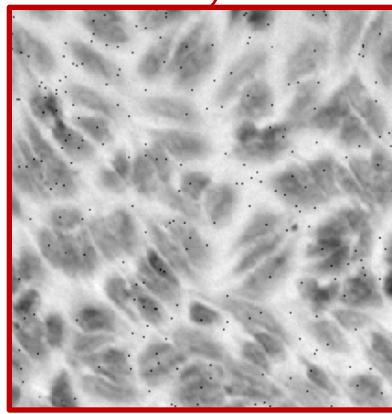
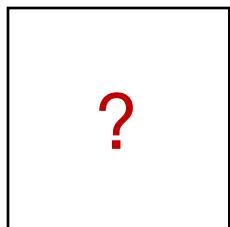
Segmentation Mask



Threshold

We can just Inference the fluorescence image itself...

Learned
illumination for
fluorescent image
inference?



Optimally illuminated

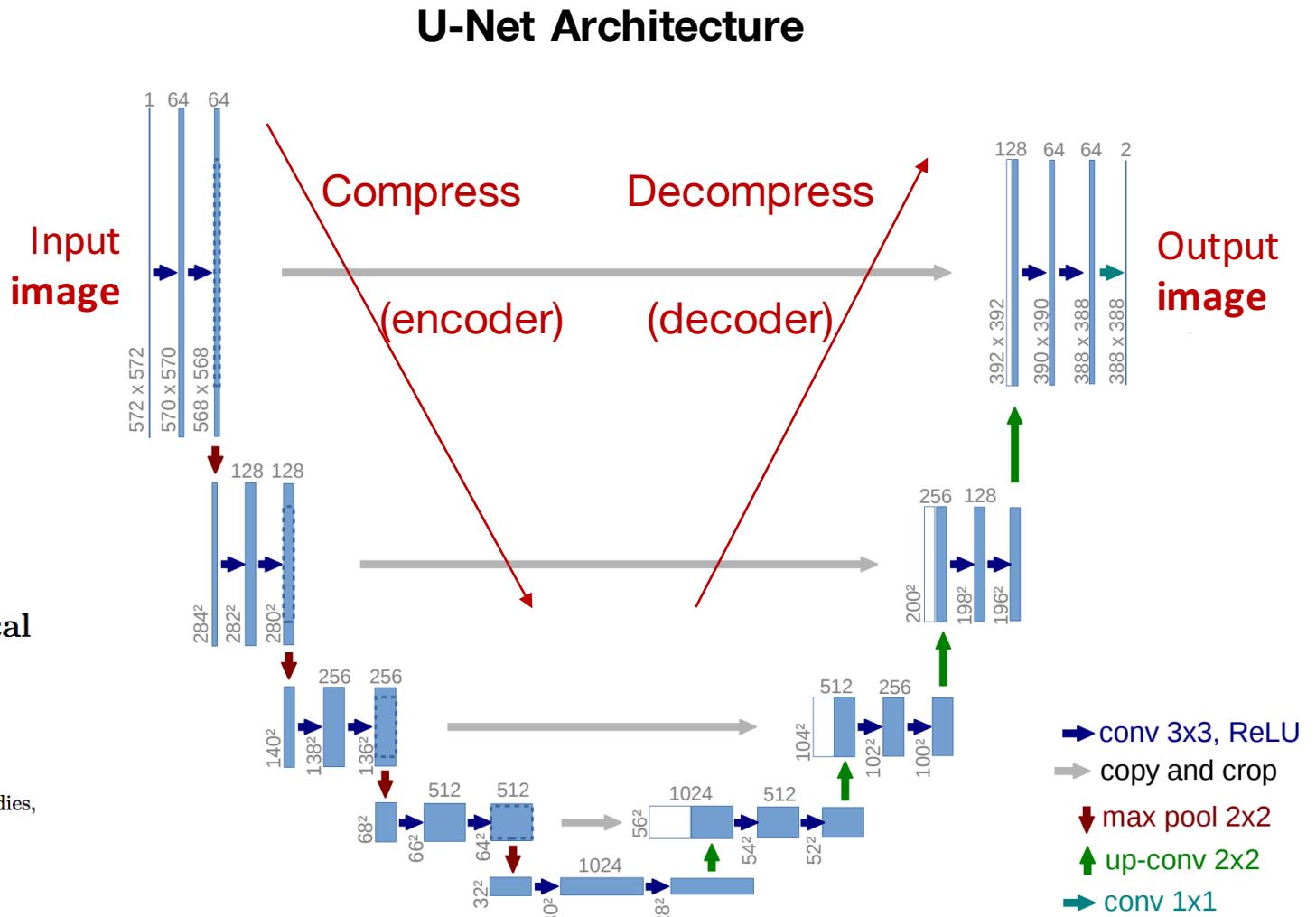
Instead, compress x-y dimensions of input image

- Compress spatial features into learned filters
- Then, decompress learned filters back into same spatial dimensions
- **Can be an autoencoder**
- Analogous to image compression
- A very powerful idea...

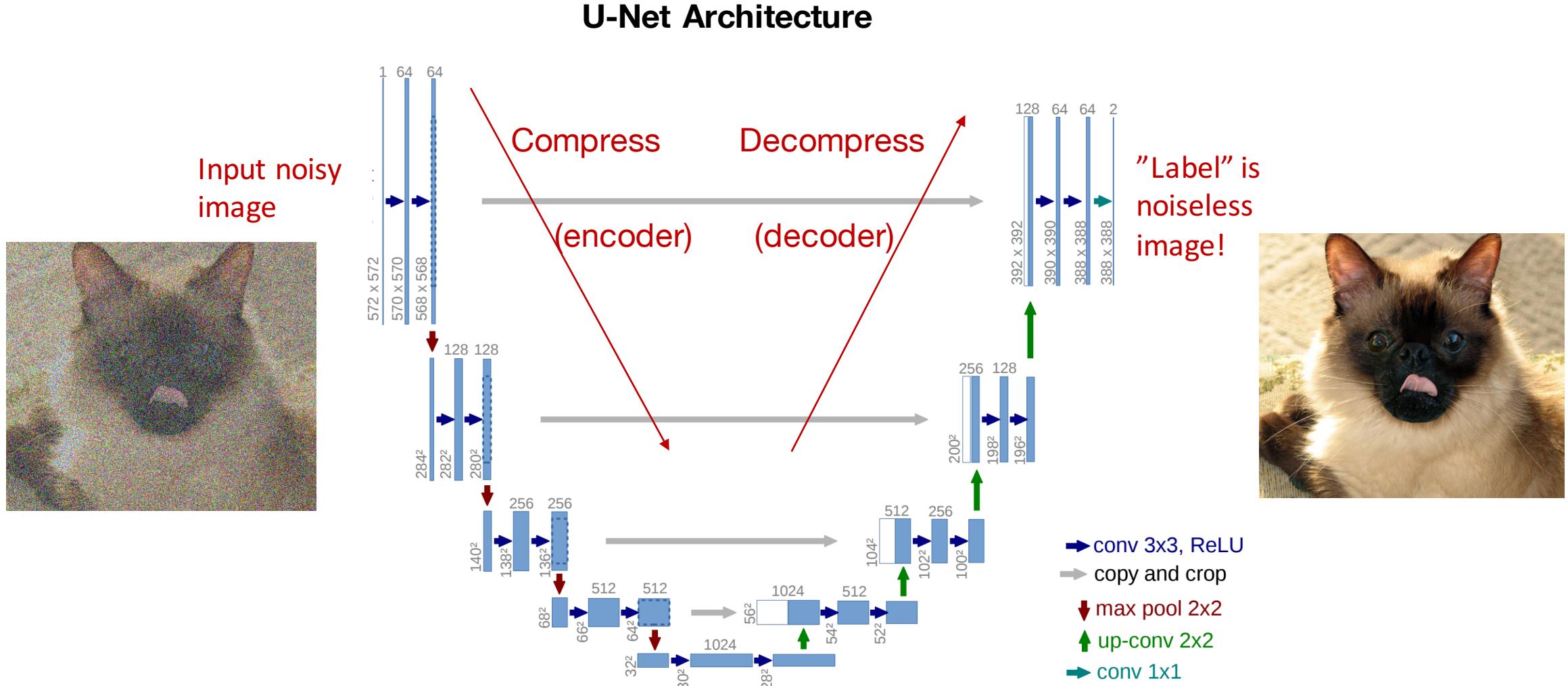
U-Net: Convolutional Networks for Biomedical Image Segmentation

Olaf Ronneberger, Philipp Fischer, and Thomas Brox

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University of Freiburg, Germany
ronneber@informatik.uni-freiburg.de,
WWW home page: <http://lmb.informatik.uni-freiburg.de/>

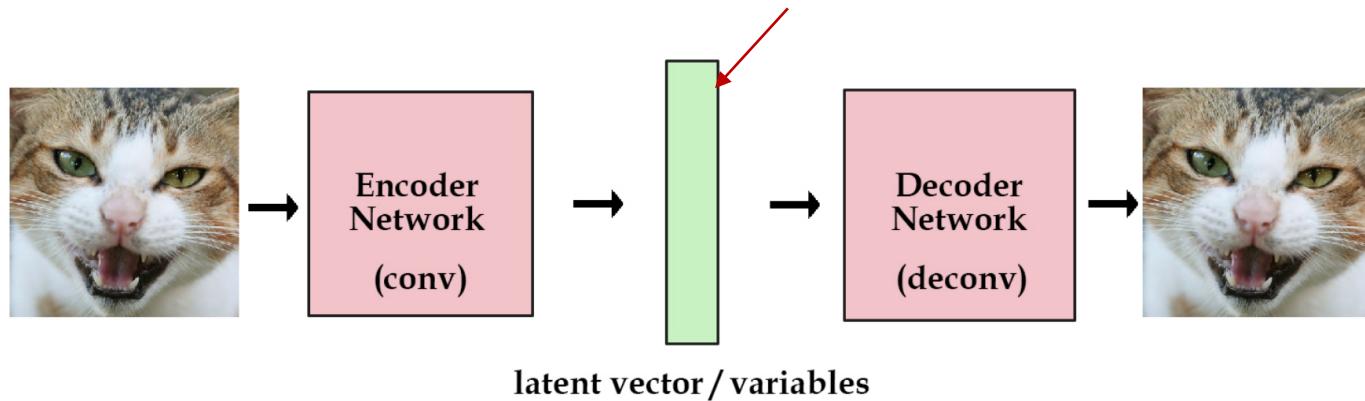


Another example: Denoising Autoencoder



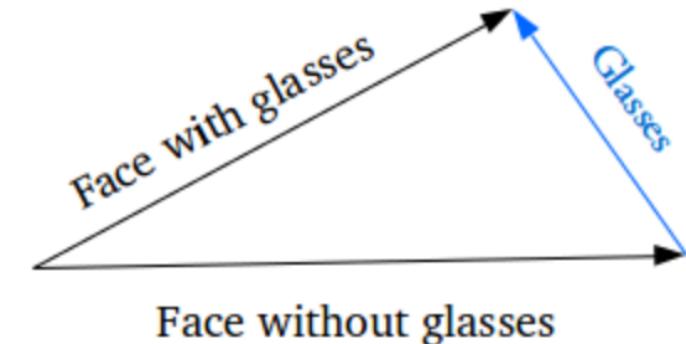
Example: Variational Autoencoder (VAE)

Force this vector to follow a Gaussian PDF



Minimize (KL) distance between latent
vector and Gaussian normal

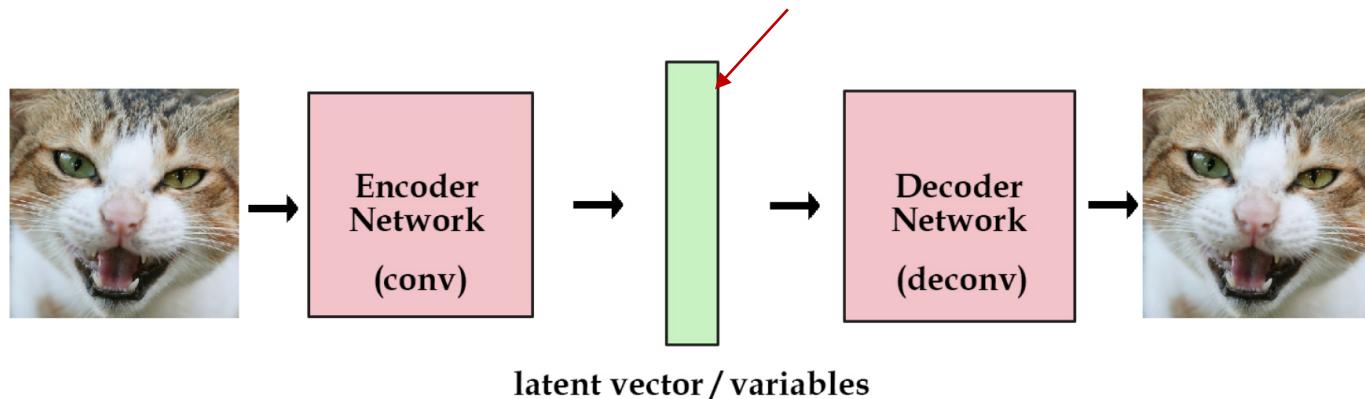
- With Gaussian PDF, can start to add/subtract latent vector in a normalized vector space



Adding new features to samples

Example: Variational Autoencoder (VAE)

Force this vector to follow a Gaussian PDF

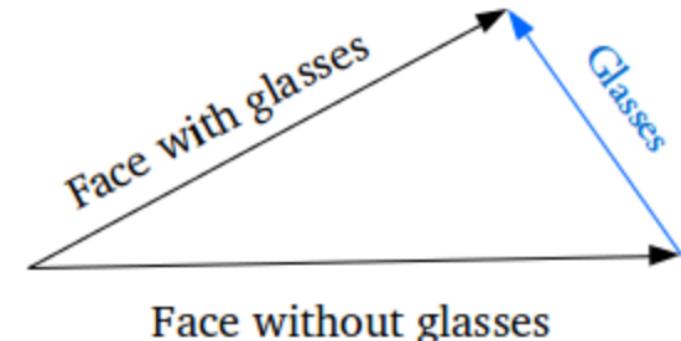


Minimize (KL) distance between latent
vector and Gaussian normal

Generative Example (once trained):

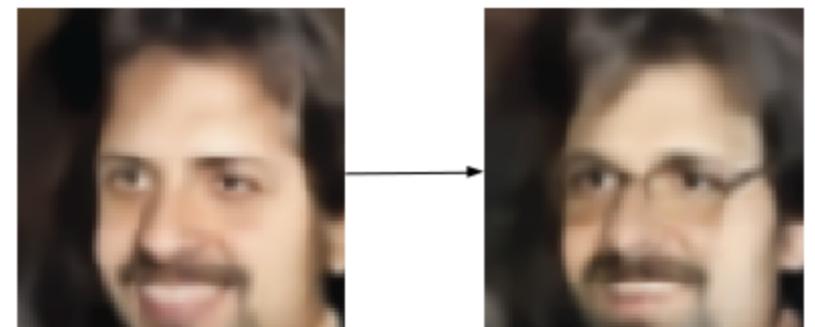
- Encode image with glasses, obtain latent vector PDF P_g
- Encode image without glasses, obtain PDF P_{ng}
- Compute $\text{diff} = P_g - P_{ng}$
- Encode new image to obtain P_{new} , add in diff
- Decode $P_{\text{new}} + \text{diff}$ to get guy with glasses!

- With Gaussian PDF, can start to add/subtract latent vector in a normalized vector space



Adding new features to samples

Glasses



Exploring a specific variation of input data[1]

Code review: See the following:

[Jupyter Notebook: A simple Autoencoder in Tensorflow/Keras](#)

<https://deepimaging.github.io/lectures/>