

U-Net: Convolutional Networks for Biomedical Image Segmentation

Sujeong Han,

0001entropy@gmail.com

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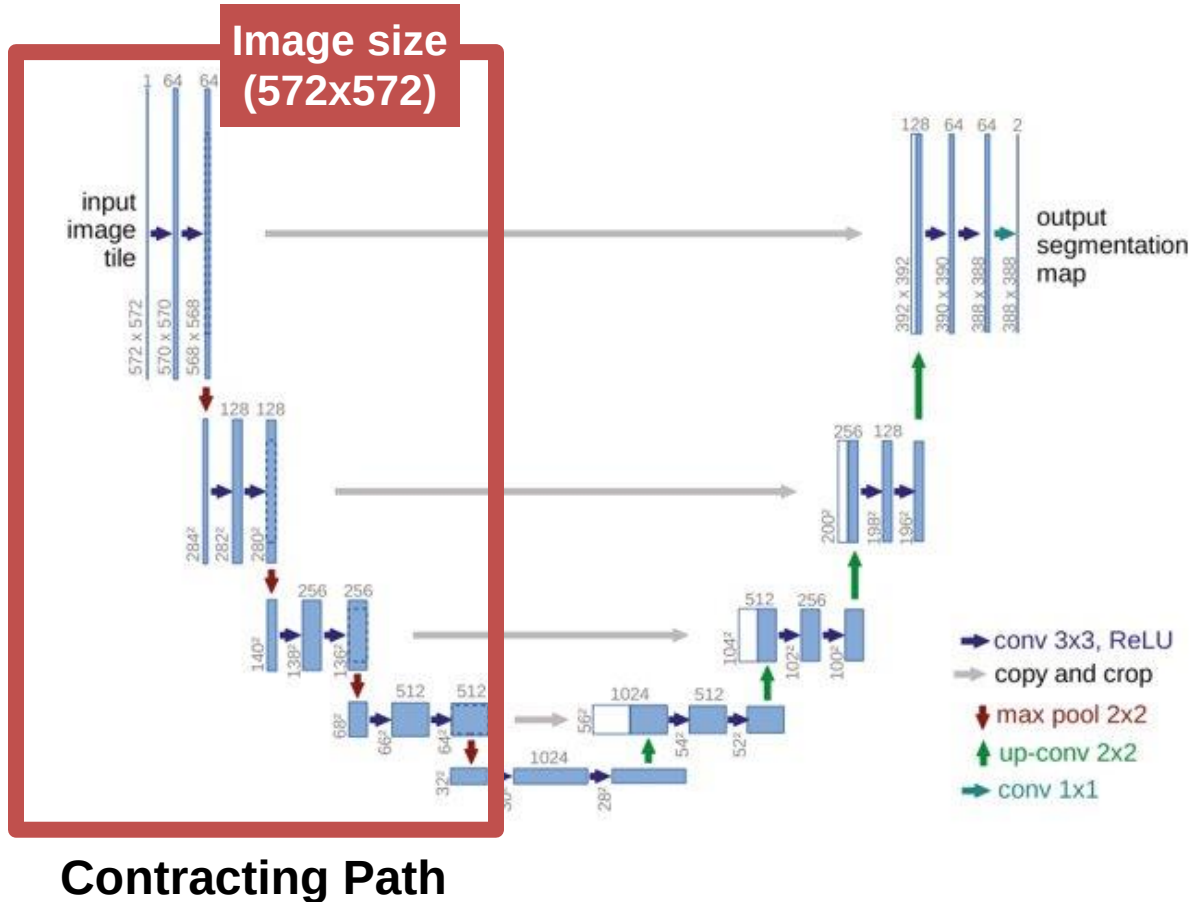
Overview

- **Network Architecture**
 - Contracting Path
 - Bottle Neck
 - Expanding Path
- **Learning method**
 - Overlap-tite strategy
- **Experiments and Result**
 - EM Segmentation challenge Dataset
 - ISBI cell tracking challenge
- **Conclusion**

Network Architecture

- U-Net (Contracting Path)

- Contracting Path that extracts context information by looking at a wide range of image pixels gradually



Repeat the downsampling process to create a feature map

Extracting contextual information from the image by expanding the reference range of surrounding pixels

When performing 3×3 convolution, **there is no padding**, so the size of the feature map is reduced

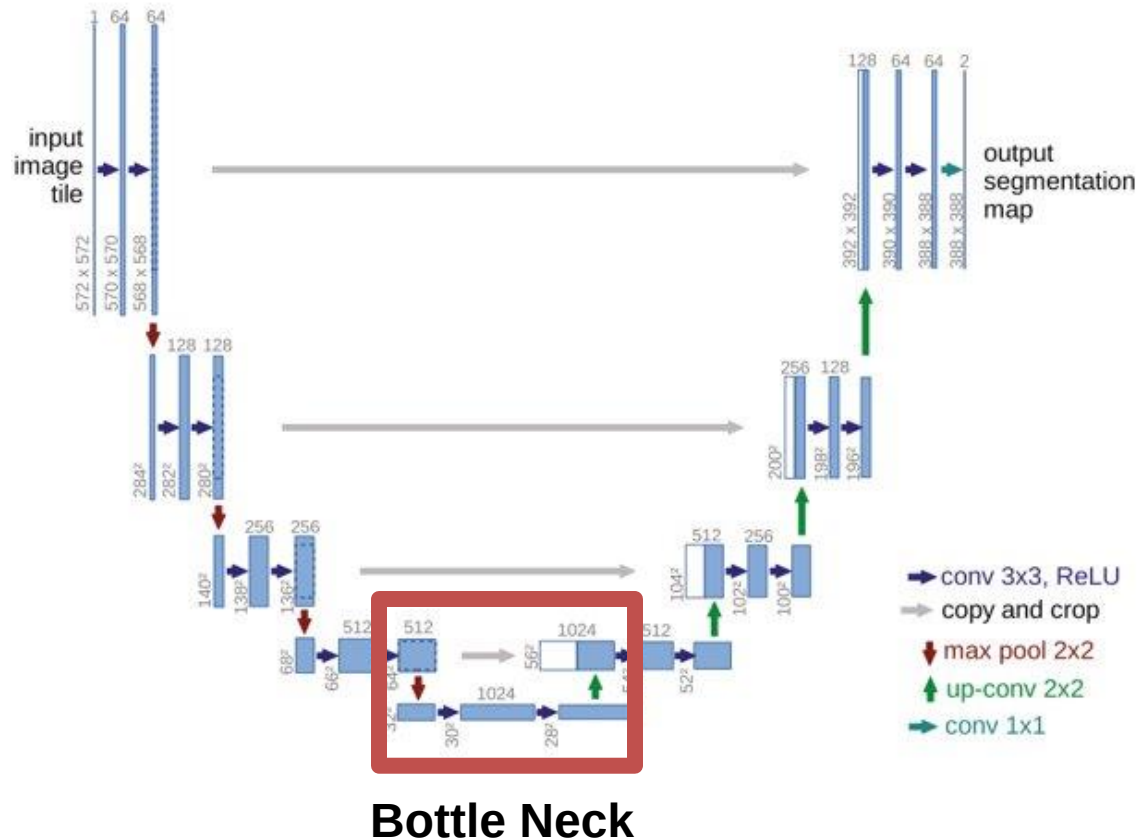
Each time downsampling, the number of channels is doubled

1 -> 64 -> 128 -> 256 -> 512 -> 1024

Network Architecture

- U-Net (Bottle Neck)

- Transition from contraction path to expansion path

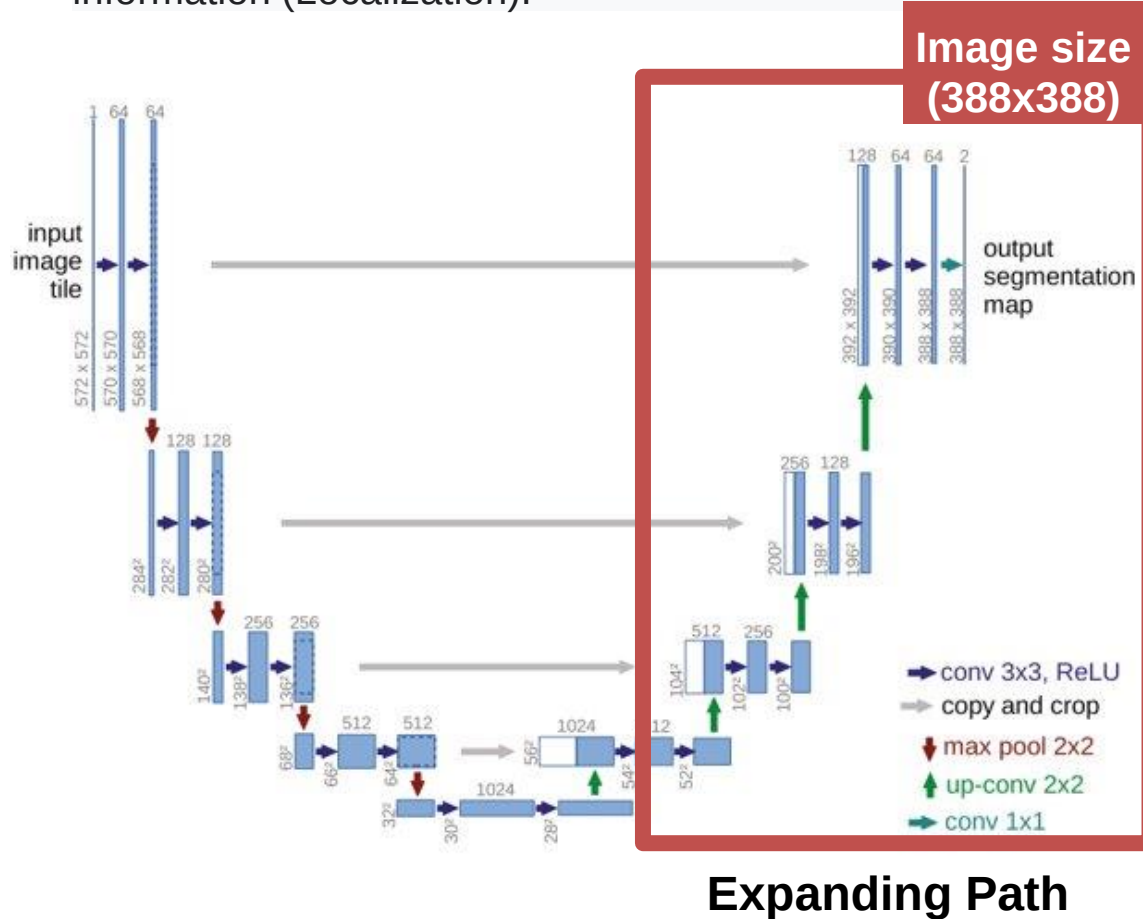


The Dropout Layer applied at the end is a device that **generalizes** the model and makes it robust to noise.

Network Architecture

- **U-Net (Expanding Path)**

- An extension path that identifies which object each pixel belongs to by combining semantic information with pixel location information (Localization).



Repeat the Upsampling process to create a feature map

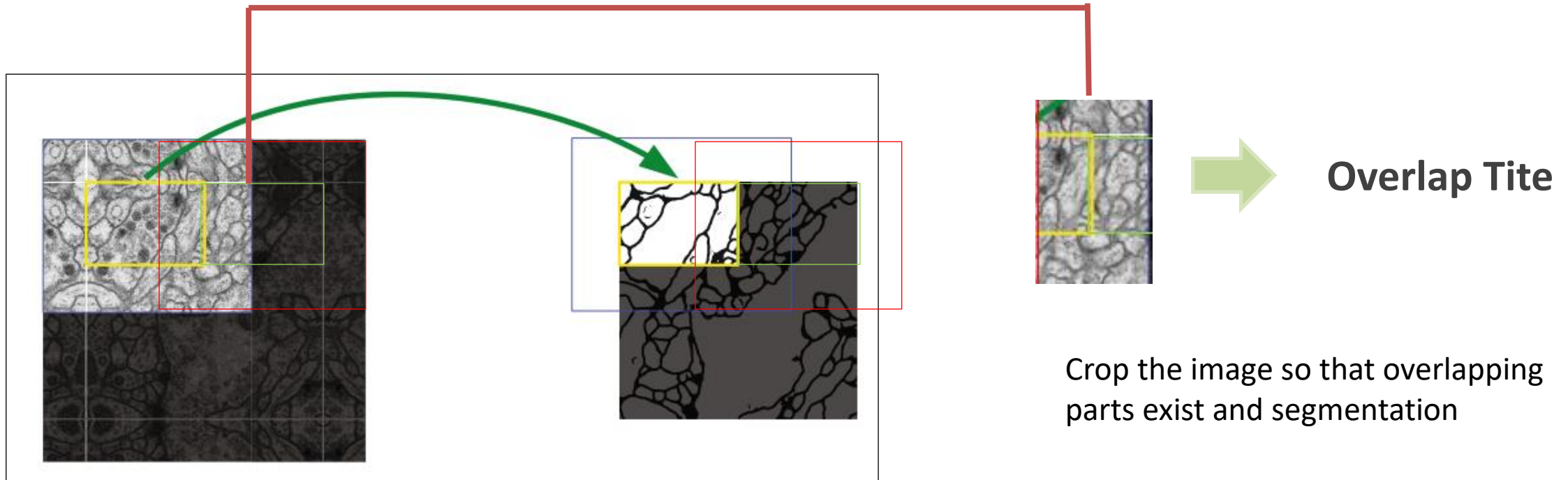
Extract the feature map of the same level from the shrinkage path and **cut it to fit the size (Cropping)** and connect it with the feature map created in the previous layer (**Concatenation**)

The role of combining **contextual information and positional information** generated in the contraction path through Skip Connection

After passing through the **1x1 convolution layer**, a **three-dimensional (Width x Height x Class) vector** representing information about which **class each pixel corresponds to** is generated

Learning method

- Overlap-tite strategy



the second closest boundary and distance from the position of pixel x

Learning method

- **Touching cells separation**

- Main challenges in cell segmentation is to separate contact objects of the same class.

Softmax
$$p_k(\mathbf{x}) = \exp(a_k(\mathbf{x})) / \left(\sum_{k'=1}^K \exp(a_{k'}(\mathbf{x})) \right)$$

Cross-Entropy
$$E = \sum_{\mathbf{x} \in \Omega} w(\mathbf{x}) \log(p_{\ell(\mathbf{x})}(\mathbf{x}))$$

$$\ell : \Omega \rightarrow \{1, \dots, K\}$$
$$w : \Omega \rightarrow \mathbb{R}$$

- Weight map for Ground-Truth

$$w(\mathbf{x}) = w_c(\mathbf{x}) + w_0 \cdot \exp \left(-\frac{(d_1(\mathbf{x}) + d_2(\mathbf{x}))^2}{2\sigma^2} \right)$$

$$w_0 = 10$$

$$\sigma \approx 5 \text{ pixels}$$

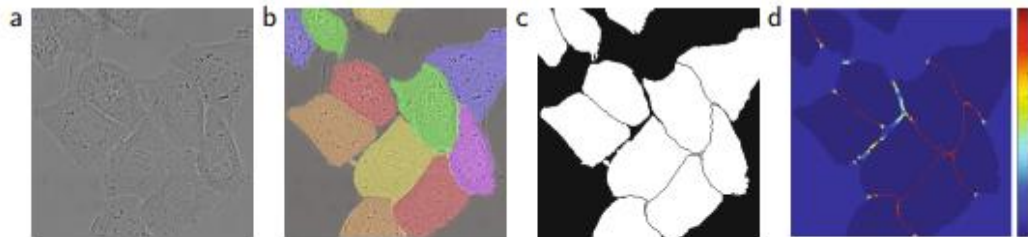


Fig. 3. HeLa cells on glass recorded with DIC (differential interference contrast) microscopy. (a) raw image. (b) overlay with ground truth segmentation. Different colors indicate different instances of the HeLa cells. (c) generated segmentation mask (white: foreground, black: background). (d) map with a pixel-wise loss weight to force the network to learn the border pixels.

Use a pre-trained weight map to separate the boundaries between contacting cells.

- $a_k(x)$ - Value where pixel x is class k (output of model per pixel)
- $p_k(x)$ - Probability that pixel x is class k (0~1)
- $\ell(x)$ - Actual Label of pixel x
- w_0 - Weight hyper-parameter of the paper, set to 10 in the paper
- σ - Weight hyper-parameter of the paper, set to 5 in the paper
- $d_1(x)$ - The nearest boundary and distance from the position of pixel x
- $d_2(x)$ - The second closet boundary and distance from the posirion of pixel x

- Use the image as large as possible
- Adjust the optimizer to learn consistently by giving Momentum (0.99)

Experiments and Result

- **EM Segmentation challenge Dataset**

- EM Segmentation challenge Dataset provides 30 training data
- Each data includes a Ground Truth Segmentation Map with separate objects and backgrounds (0 or 1) with images.

- **Data augmentation methods** : Rotation, Shift, Elastic distortion

Table 1. Ranking on the EM segmentation challenge [14] (march 6th, 2015), sorted by warping error.

Rank	Group name	Warping Error	Rand Error	Pixel Error
	** human values **	0.000005	0.0021	0.0010
1.	u-net	0.000353	0.0382	0.0611
2.	DIVE-SCI	0.000355	0.0305	0.0584
3.	IDSIA [2]	0.000420	0.0504	0.0613
4.	DIVE	0.000430	0.0545	0.0582
	⋮			
10.	IDSIA-SCI	0.000653	0.0189	0.1027

Experiments and Result

- ISBI cell tracking challenge

- “PhC-U373” data provided as training data from 35 images recorded with a phase-contrast microscope
- “DIC-HeLa” data were recorded from HeLa cells under a microscope and 20 images were provided as training data.

Table 2. Segmentation results (IOU) on the ISBI cell tracking challenge 2015.

Name	PhC-U373	DIC-HeLa
IMCB-SG (2014)	0.2669	0.2935
KTH-SE (2014)	0.7953	0.4607
HOUS-US (2014)	0.5323	-
second-best 2015	0.83	0.46
u-net (2015)	0.9203	0.7756

Conclusion

- **Conclusion**

- “U-Net is a model that applies Up-sampling and Skip Architecture of an expanded concept than FCNs.
- As a result, the structure of U-Net exhibits excellent performance in various biomedical image segmentation problems by utilizing Data Augmentation with only a very small amount of training data.
- Skip Architecture makes it possible to stack layers deeply, so that complex tasks can be performed well.
- Reducing the loss of information from the bottle neck

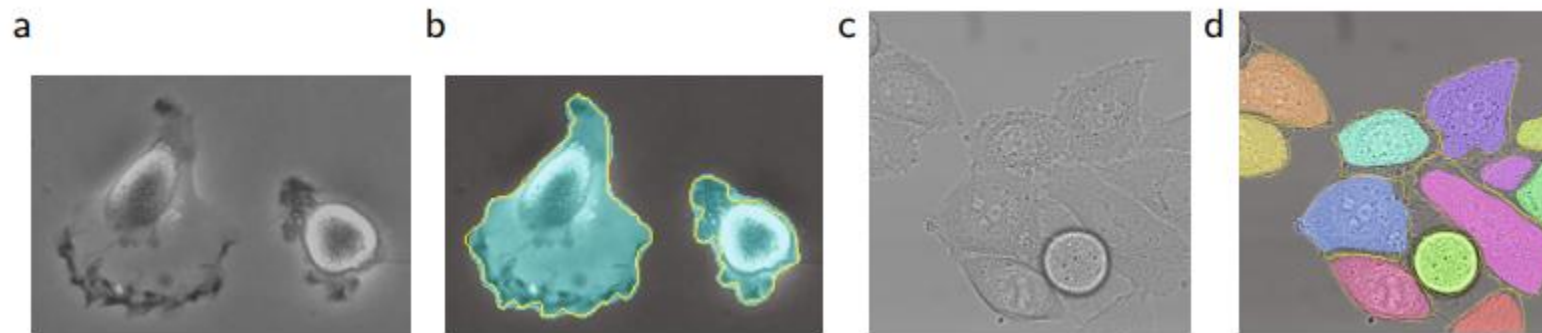


Fig. 4. Result on the ISBI cell tracking challenge. (a) part of an input image of the “PhC-U373” data set. (b) Segmentation result (cyan mask) with manual ground truth (yellow border) (c) input image of the “DIC-HeLa” data set. (d) Segmentation result (random colored masks) with manual ground truth (yellow border).

Thank you