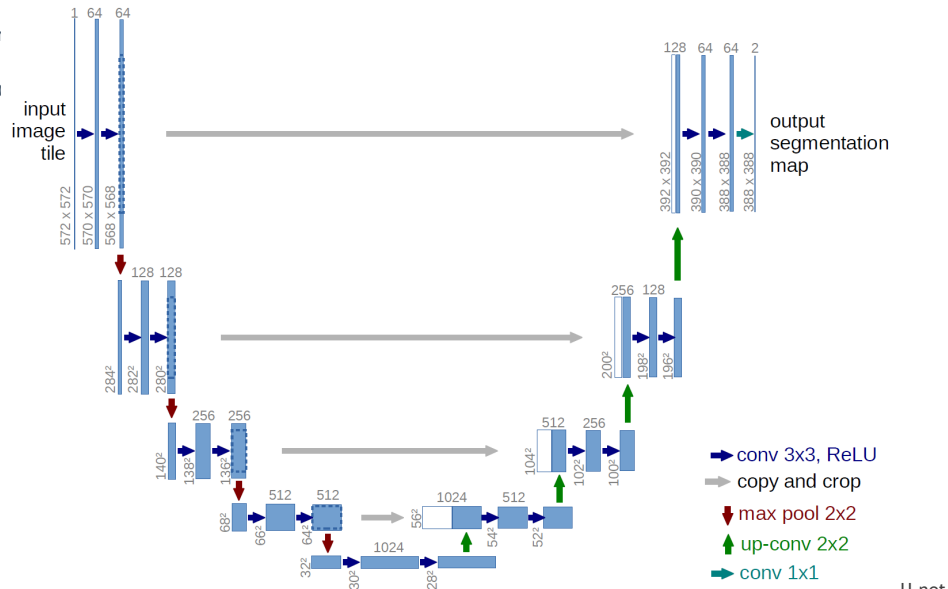


U-Net: Convolutional Networks for Biomedical Image Segmentation

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- The u-net is convolutional network architecture for fast and precise segmentation of images. Up to now it has outperformed the prior best method (a sliding-window convolutional network) on the [ISBI challenge for segmentation of neuronal structures in electron microscopic stacks](#). It has won the [Grand Challenge for Computer-Automated Detection of Caries in Bitewing Radiography at ISBI 2015](#), and it has won the [Cell Tracking Challenge at ISBI 2015](#) on the two most challenging transmitted light microscopy categories (Phase contrast and DIC microscopy) by a large margin (See also [our announcement](#)).



architecture (example for 32x32 pixels in the lowest resolution). Each blue box corresponds to a multi-channel feature map. The number of channels is denoted on top of the box. The x-y-size is provided at the lower left edge of the box. White boxes represent copied feature maps. The arrows denote the different operations.

Article describing U-net



[U-Net: Convolutional Networks for Biomedical Image Segmentation](#) [doi>](#)

Olaf Ronneberger, Philipp Fischer, Thomas Brox

Medical Image Computing and Computer-Assisted Intervention (MICCAI), Springer, LNCS, Vol.9351: 234--241, 2015, available at [arXiv:1505.04597 \[cs.CV\]](#)

5 Minute Teaser Presentation of the U-net

U-net: Convolutional Networks for Biomedical Image Segmentation

Olaf Ronneberger, Philipp Fischer, and Thomas Brox

Computer Science Department and
BIOSS Centre for Biological Signalling Studies
University of Freiburg, Germany

▶ 0:00 / 5:03    
Download video: [u-net-teaser.mp4](#) (68MB)

Download

We provide the u-net for download in the following archive: [u-net-release-2015-10-02.tar.gz](#) (185MB). It contains the ready trained network, the source code, the matlab binaries of the modified caffe network, all essential third party libraries, the matlab-interface for overlap-tile segmentation and a greedy tracking algorithm used for our submission for the ISBI cell tracking challenge 2015. Everything is compiled and tested only on Ubuntu Linux 14.04 and Matlab 2014b (x64)

To apply the segmentation and the tracking to the images in "PhC-C2DH-U373/01" simply run the shell script

```
./segmentAndTrack.sh
```

The resulting segmentation masks will be written to "PhC-C2DH-U373/01_RES"

If you do not have a CUDA-capable GPU or your GPU is smaller than mine, edit `segmentAndTrack.sh` accordingly (see there for documentation). If you have any questions, you may contact me at ronneber@informatik.uni-freiburg.de, but be aware that I can not provide any support.

Olaf Ronneberger

Webmaster Disclaimer

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