



Scientific IT Services (SIS): Team and Mission

 Section of ETH ITS and a pool of experts (~45 members)

Main tasks:

- Enable research
- Improve efficiency
- Scientific computing experts & data scientists from various scientific backgrounds
- Scientific software developers with computer science background and industry experience
- System administrators & DevOps for the HPC infrastructure and research IT platforms



Working model

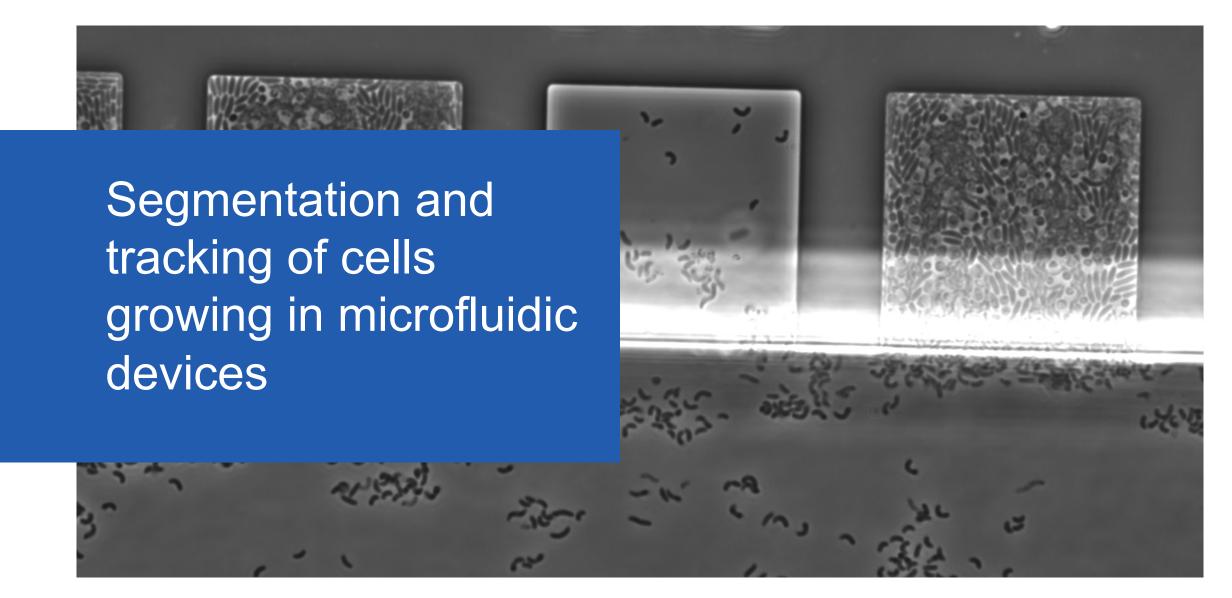
- Contracts with ETH researchers to the extent of:
 - a few days full-time support
 - one or several years part-time (20-50%) support



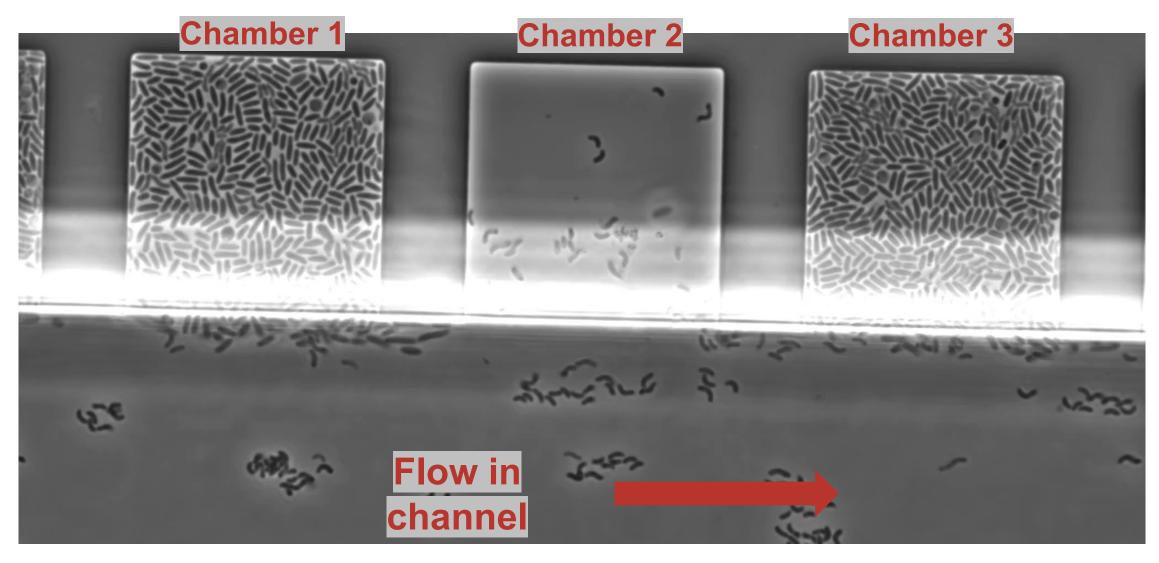
SIS Services





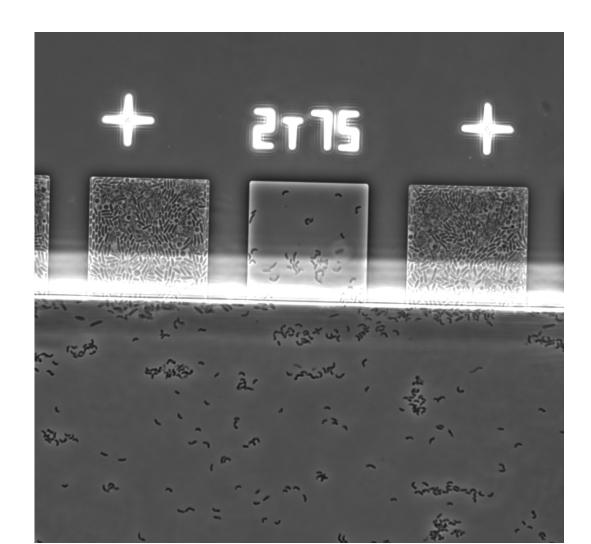


Cell growth in microfluidic devices





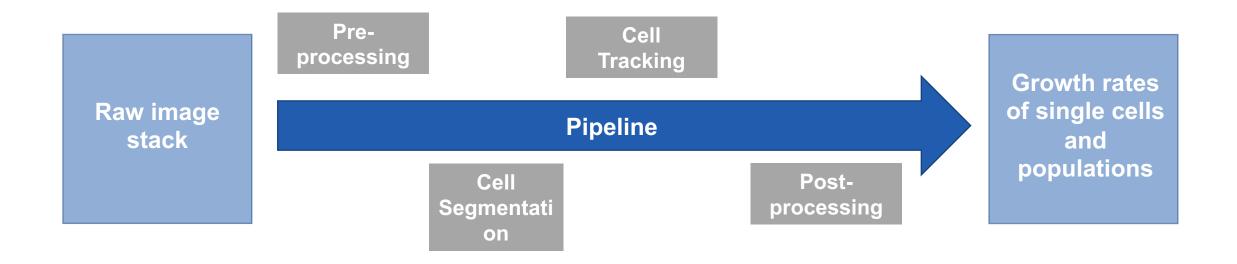
Quantification of population growth and dynamics



Required results:

- Number of cells
- Position of cells
- Relationship between cells
- Movement of cells
- Interaction between cell populations

Requirements for image analysis



Methods used before:

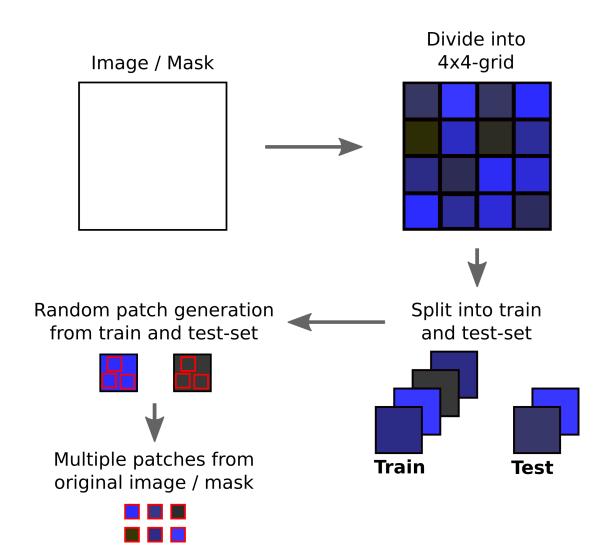
 State-of-the-art tools (ImageJ, Ilastik, SuperSegger etc.)

Problems:

- Methods don't work properly for their datasets
- No software development or Machine Learning expertise in research group
- Data analysis is bottleneck of research work
- → Contacted SIS for support

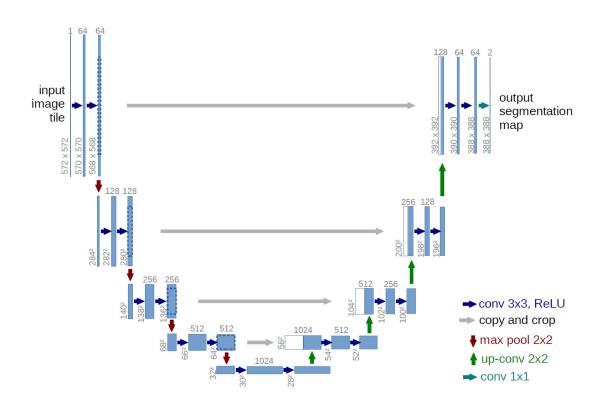


Dataset generation



- Genenration of training dataset with minimal manual effort
- Original image size: ~500 x 500 pixels
- Size of single cell: ~30 x 30 pixels
- → allows split of original image in multiple smaller patches
- Data augmentation of each pach
- Approach generates ~ 8000 training images based on one original image

Deep Neural Network for segmentation: U-Net



- Convolution block: Learning of cell shape
- Downstream branch: 'what'-information
- **Upstream branch**: 'where'-information

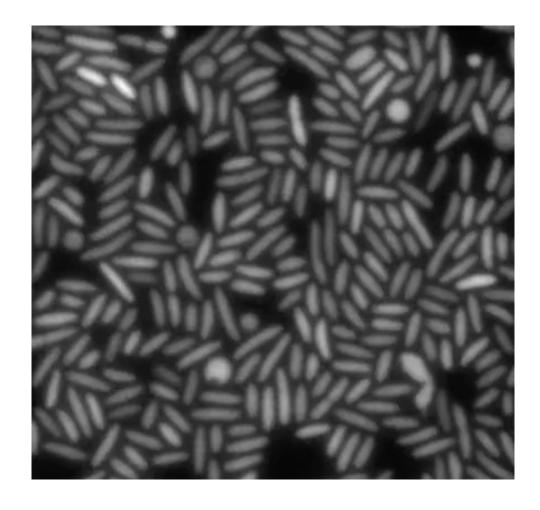
Reasons for using the U-Net:

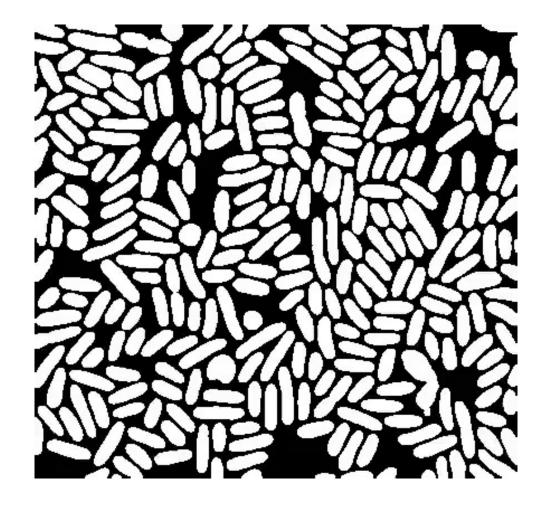
- State of the art machine learning model for segmentations (> 15 000 citations)
- Winner in several image segmentation competitions

Ronneberger et al, MICCAI 2015



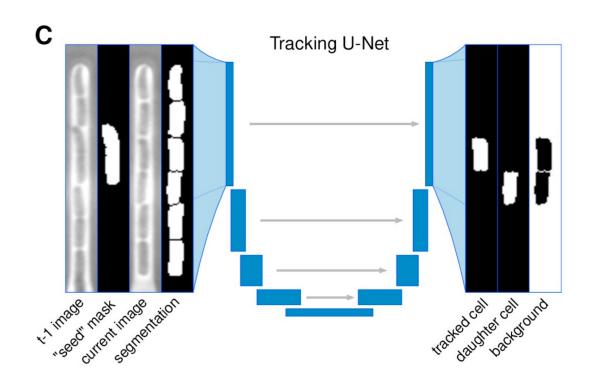
Segmentation results







Deep Neural Network for tracking: U-Net

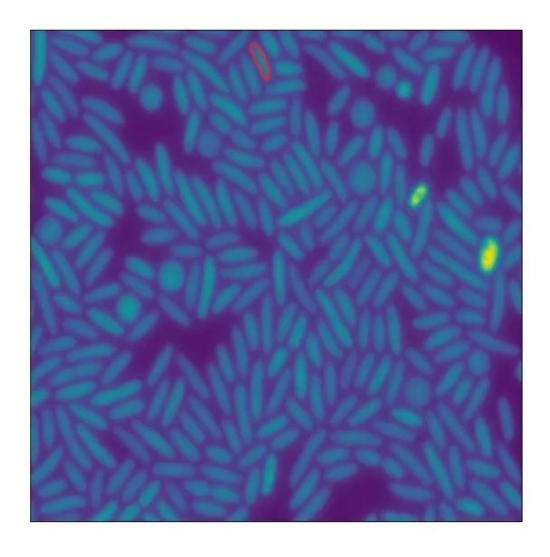


- Modified input and output compared to original U-Net
- Tracking is computed per time frame and per cell

Lugagne et al, Plos Computational Biology, 2020



Tracking results



- Application of pretrained network
- First results look promising, but fine-tuning with own training-dataset is necessary



Requirements for upscaling

- Increasing interest from collaboration partners in pipeline
- Adaptations of pipeline are required:

1. Neural Networks:

Building general models for different datasets and cell types

2. **GUI**:

Manual correction of segmentation and tracking

3. Workflow managers:

- Enable modular usage of pipeline
- Storage of metadata

4. HPC:

Processing of multiple datasets at the same time



From prototype to production : role of SIS in upscaling





Benefits, challenges and risks of this working model

Benefits:

- Pool of experts which work on one solution
- Easy access to knowledge which is not available in research group
- Continuous exchange between researchers and SIS members

Challenges and risks:

- Find a common language (IT & Biology)
- Expectation management (Al can not perform miracles)
- Not every field/problem is suitable for the application of ML



Thank you!

https://sis.id.ethz.ch/

