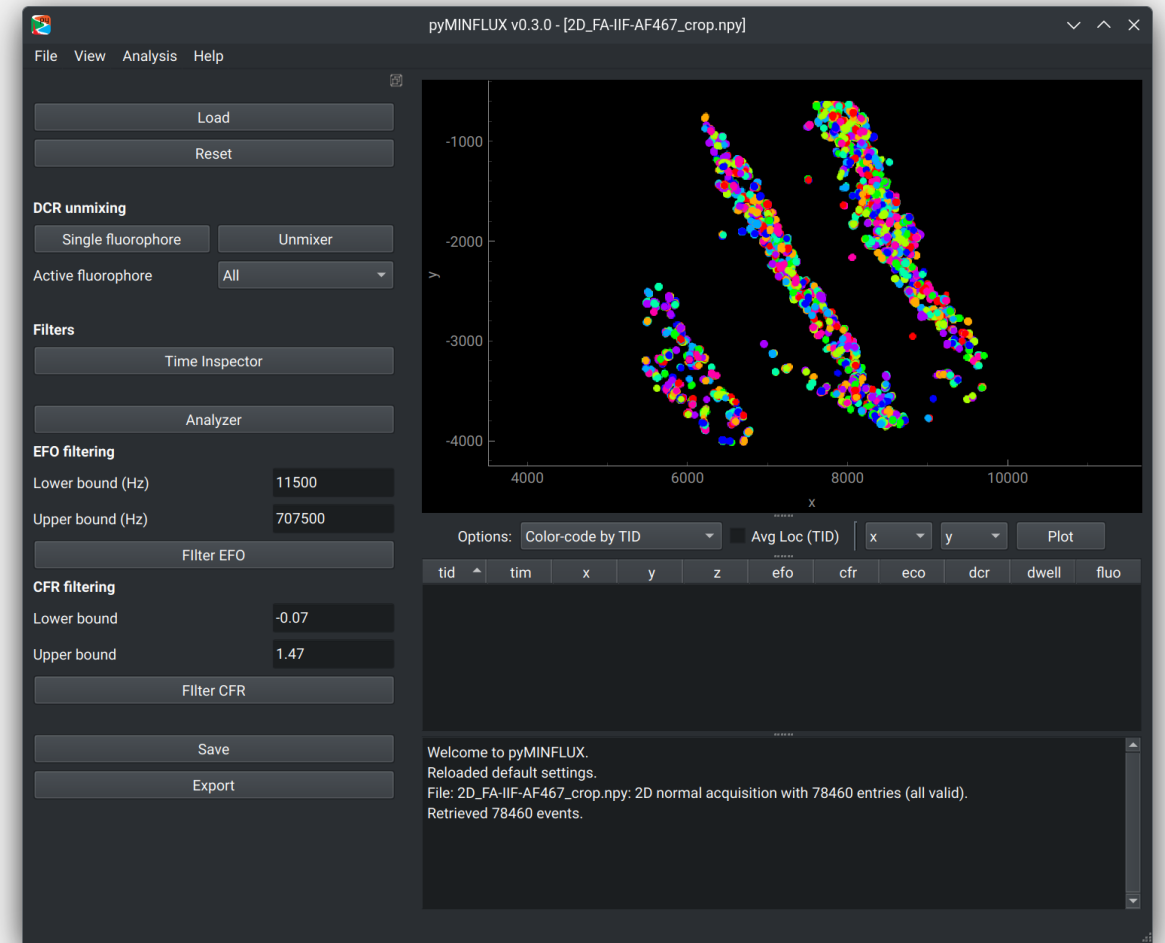


# Collaborative expertise

Synergies between multi-disciplinary experts and end users in software development

Aaron Ponti



# About me

## Education

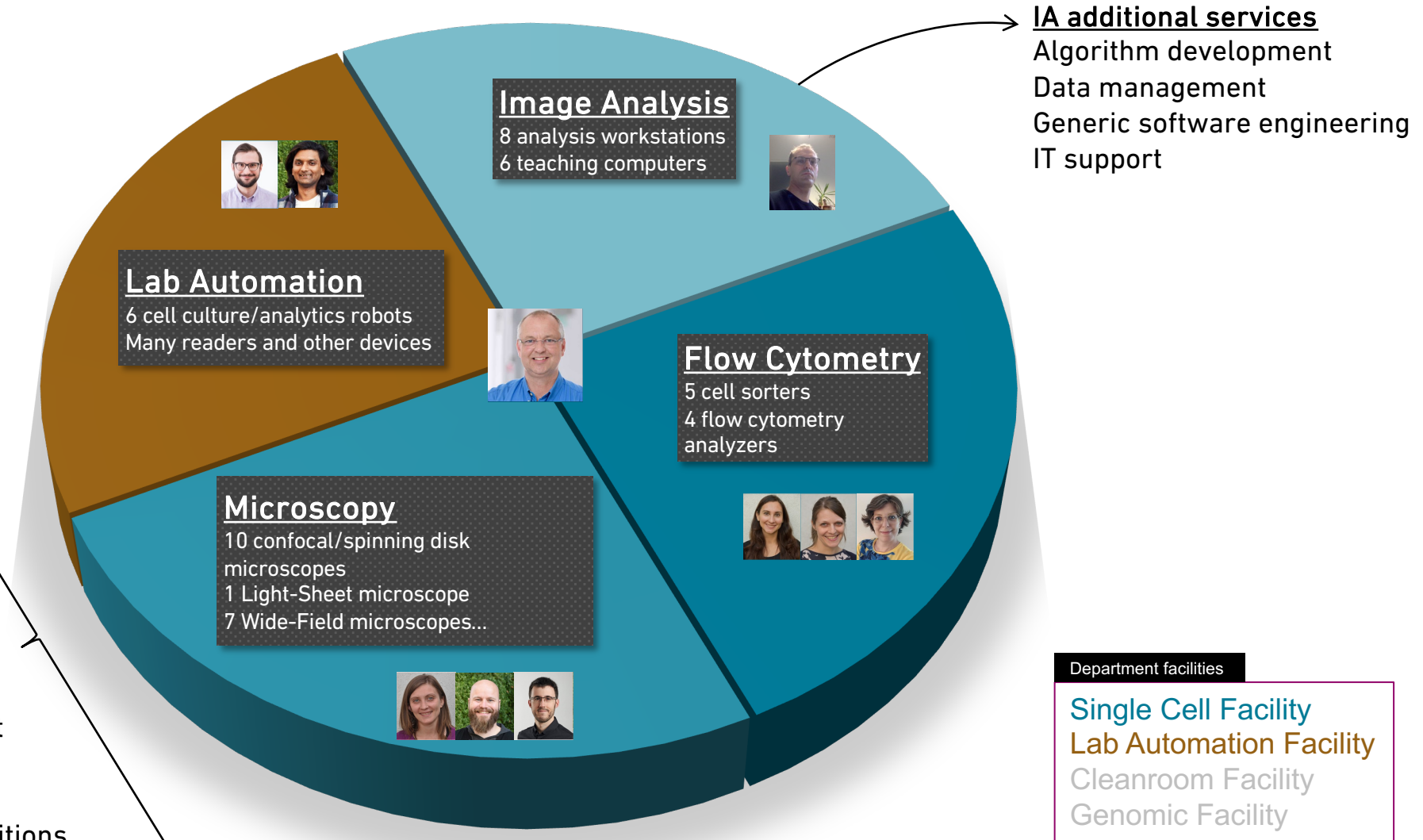
- M. Sc. Biotechnology  
D-BIOL ETH Zurich
- PhD Image Analysis  
D-MAVT ETH Zurich
- Post-doctoral fellow Image Analysis  
The Scripps Research Institute, San Diego, CA

Briefly  
D-INFK ETH Zurich

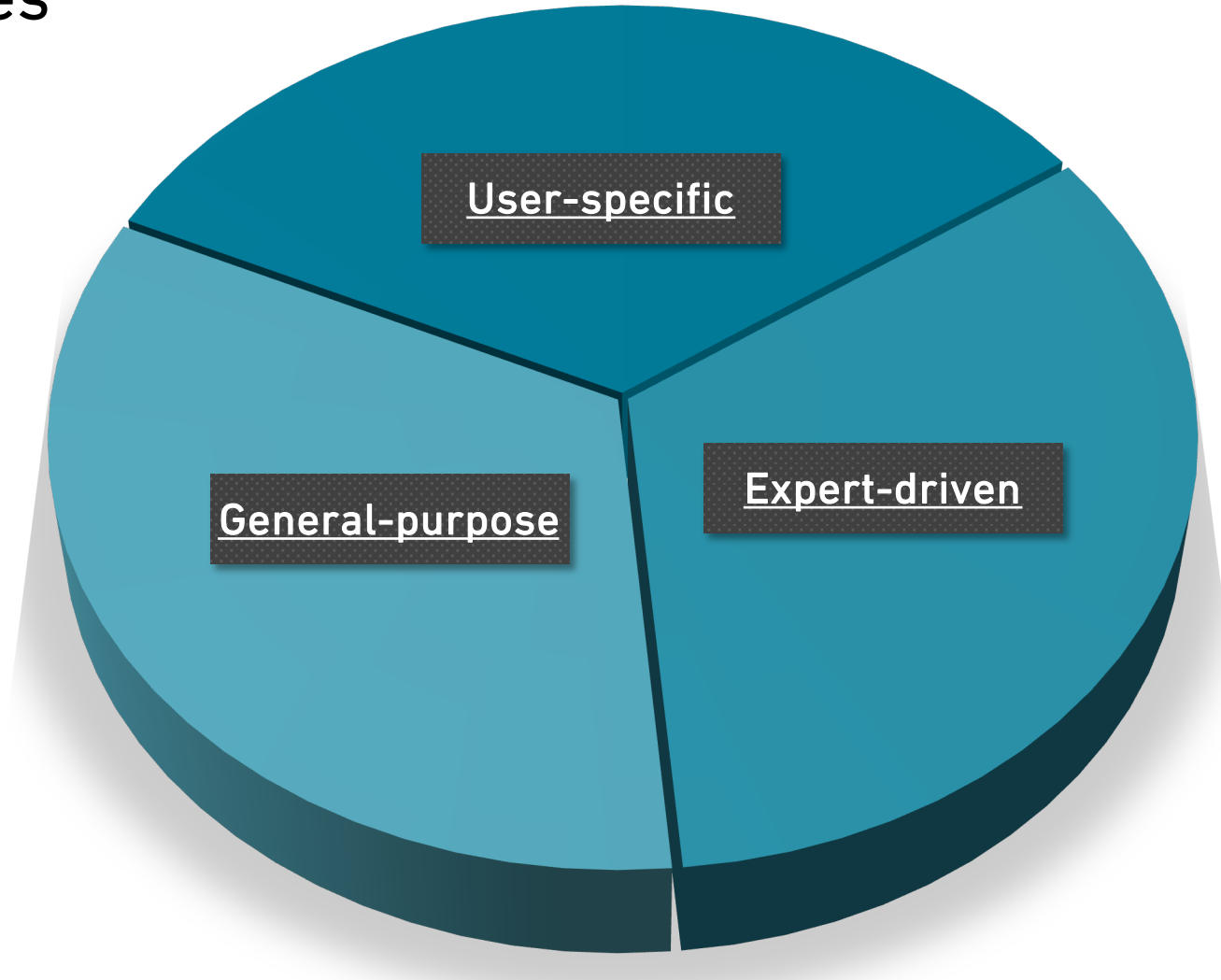
## As a grown-up

- Image analysis specialist  
Friedrich Miescher Institute :: Facility for Advanced Imaging and Microscopy
- Image Analysis Specialist & Software and Data Management Engineer  
D-BSSE ETH Zurich :: Single Cell Facility

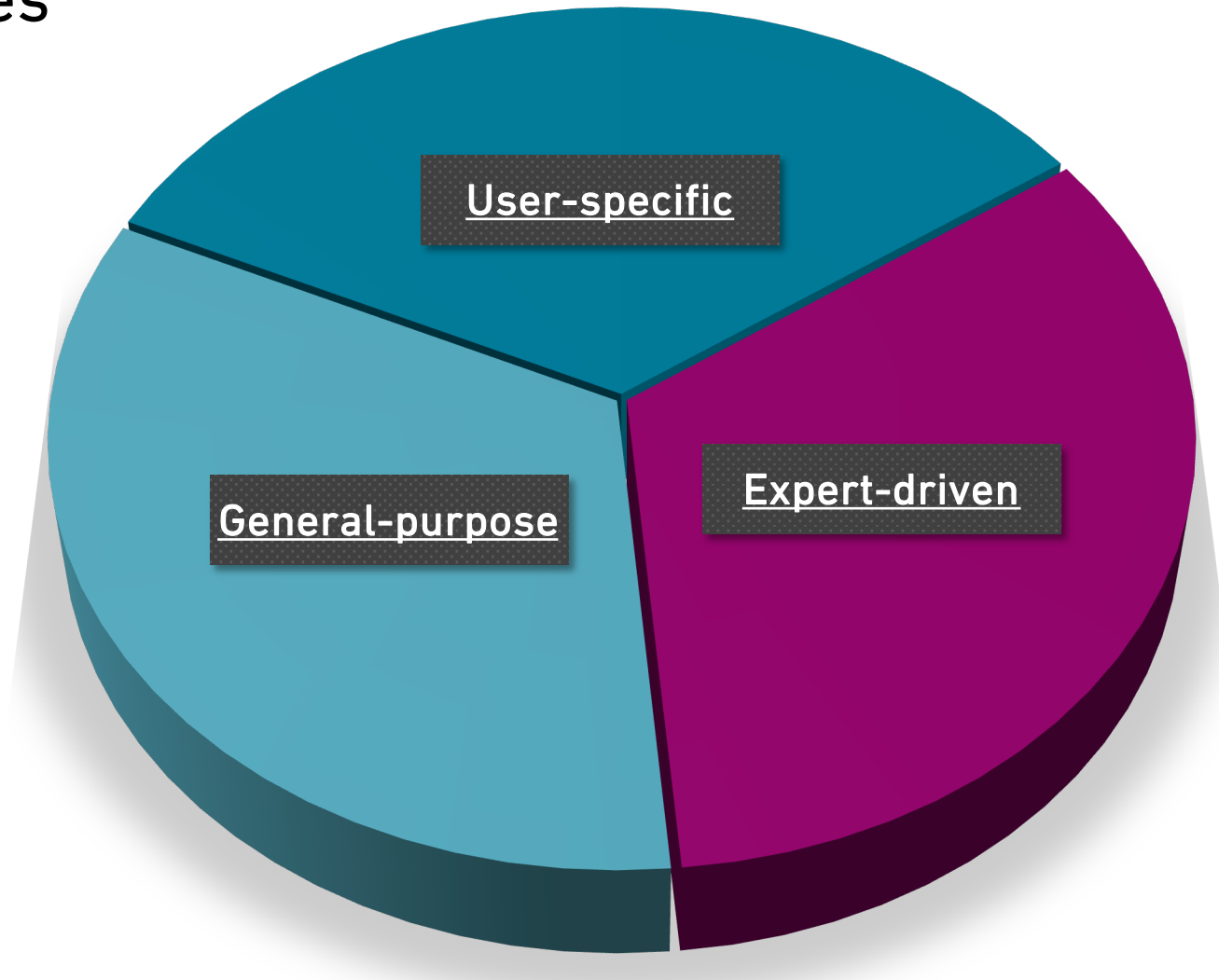
# About us



# IA project classes



# IA project classes



# Types of project

	User-specific projects	Expert-driven projects	General-purpose projects
Target audience	Single user	Several users	Many users
Specificity	Highly specialized	Specialized but scalable	Generalized
Team composition	Me (with user feedback)	Me and field experts (with user feedback)	Large and diverse team
Problem focus	Single, user-specific	Niche problems	Broad
Complexity	Varies	Complex	Moderate to complex
Scalability	Limited	Moderate	High
User input	Continuous	Initial and iterative (at release)	Initial and iterative (at release)
Project time	Short to moderate	Moderate to long	Long
Resource allocation	Moderate	High	Very high
Field knowledge	High (me)	Very high (experts)	Moderate
Code quality/testing	Minimal to moderate	High	Very high

# User-specific projects

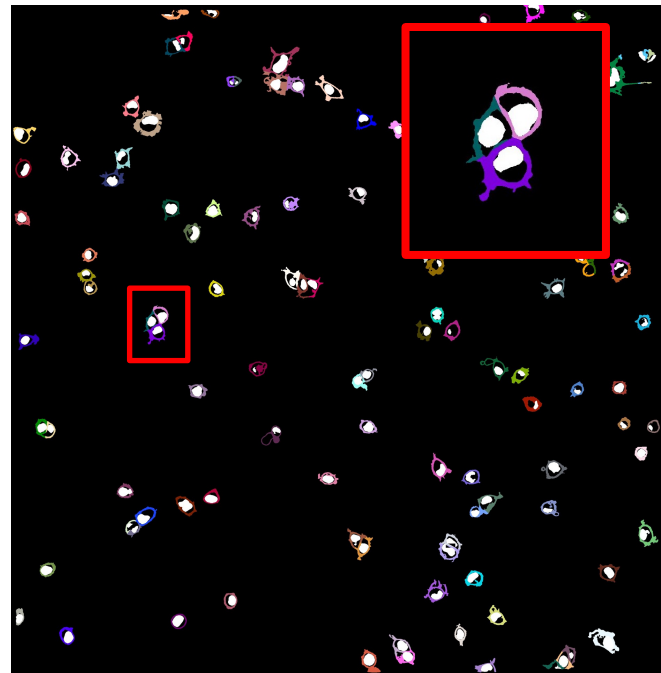
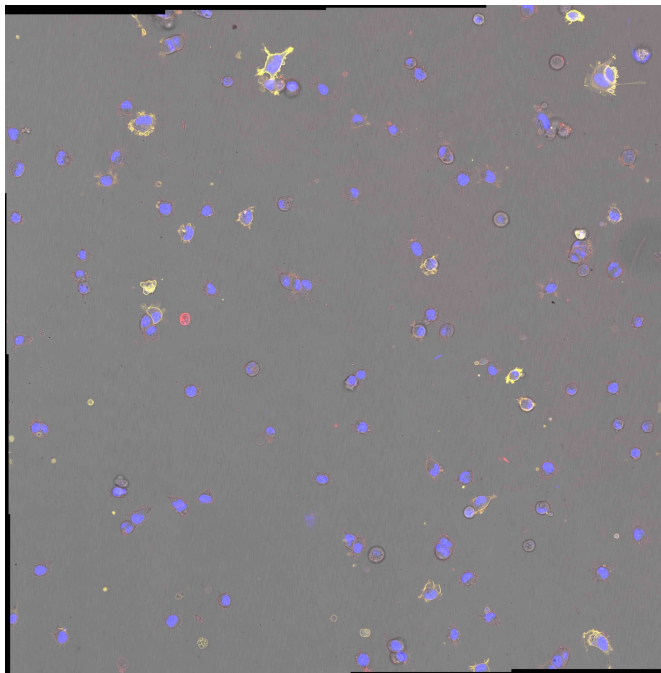
	User-specific projects
Target audience	Single user
Specificity	Highly specialized
Team composition	Me (with user feedback)
Problem focus	Single, user-specific
Complexity	Varies
Scalability	Limited
User input	Continuous
Project time	Short to moderate
Resource allocation	Moderate
Field knowledge	High (me)
Code quality/testing	Minimal to moderate

# User-specific projects :: Membrane localization study of a sensor



“

We designed a plasmid-based sensor consisting of two fusion-protein, that bind activated KRAS and can phosphorylate a synthetic transcription factor if it they dimerize. (...) As KRAS is a membrane protein and our sensor binds only to activated KRAS, membrane localization of our sensor-SYFP-fusion also tells us if KRAS is activated. (One of the) aim(s) of the study: investigate differences in membrane localization of the sensor between mutant and wild-type KRAS.



	A	B	C	D
1	cell_id	membrane_signal	total_signal	ratio
2	62	5740.9854	6643.087	0.86420447
3	72	26428.516	33384.64	0.79163694
4	81	3419.3042	4384.9595	0.7797801
5	99	3454.4004	4615.084	0.7485022
6	100	5038.2393	5553.0483	0.90729254
7	102	4730.7227	5524.704	0.8562853
8	103	4360.502	4901.9644	0.88954175
9	104	67296.74	69806.89	0.96404153
10	107	3924.6875	5026.2783	0.7808337
11	108	3036.5955	4003.891	0.7584111
12	111	11538.633	16132.592	0.7152374
13	112	5822.574	6326.4824	0.92034936
14	125	5634.9688	6411.24	0.87892026
15	129	17054.633	21817.885	0.7816813
16	131	5955.9385	7285.884	0.81746274
17	134	3460.176	3846.4526	0.8995759
18	135	2996.8965	3478.8113	0.8614714
19	138	3821.9531	4512.3306	0.847002
20	139	5966.1807	8770.289	0.68027186
21	144	3959.9258	4624.1426	0.85635895



Stats

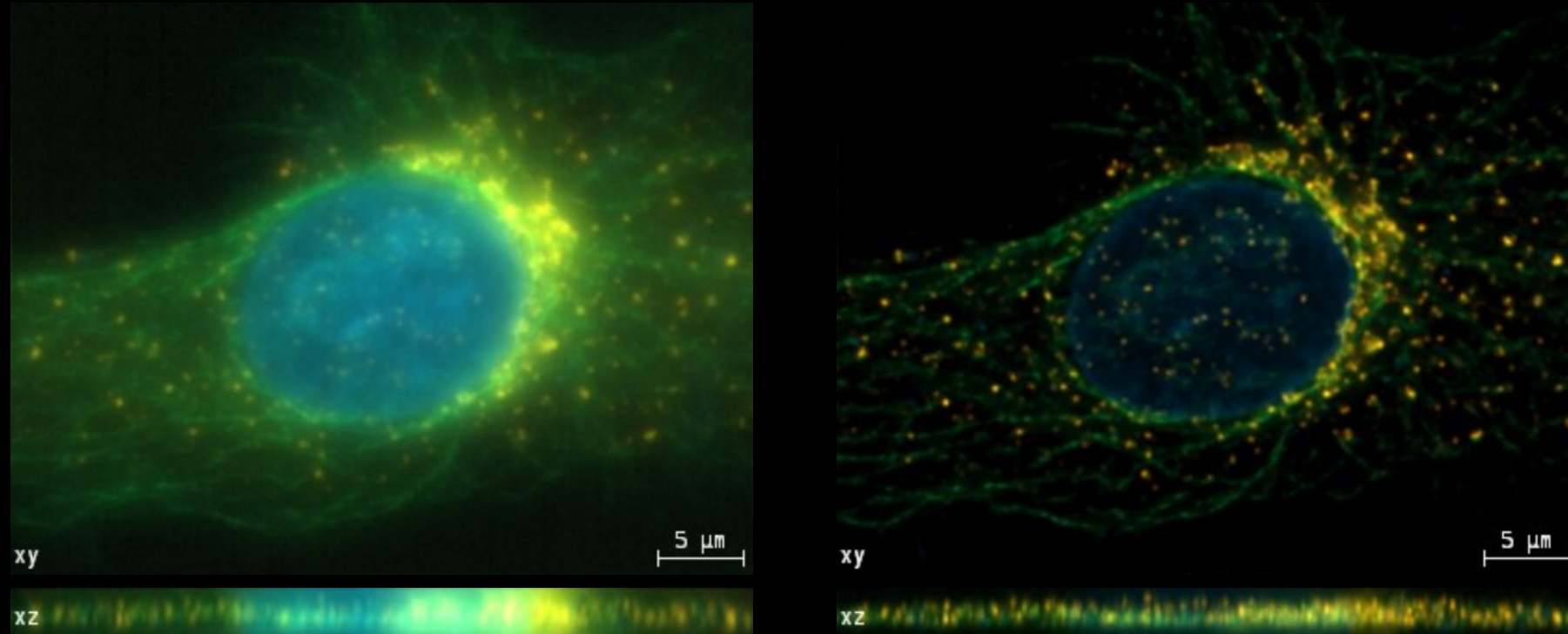
```
$ python analyze_membranes.py --folder E.GS3.56.1_reseeding --result result.csv --max-workers 8
```



# General-purpose projects

	General-purpose projects
Target audience	Many users
Specificity	Generalized
Team composition	Large and diverse team
Problem focus	Broad
Complexity	Moderate to complex
Scalability	High
User input	Initial and iterative (at release)
Project time	Long
Resource allocation	Very high
Field knowledge	Moderate
Code quality/testing	Very high

# General-purpose projects :: Huygens Remote Manager (HRM)

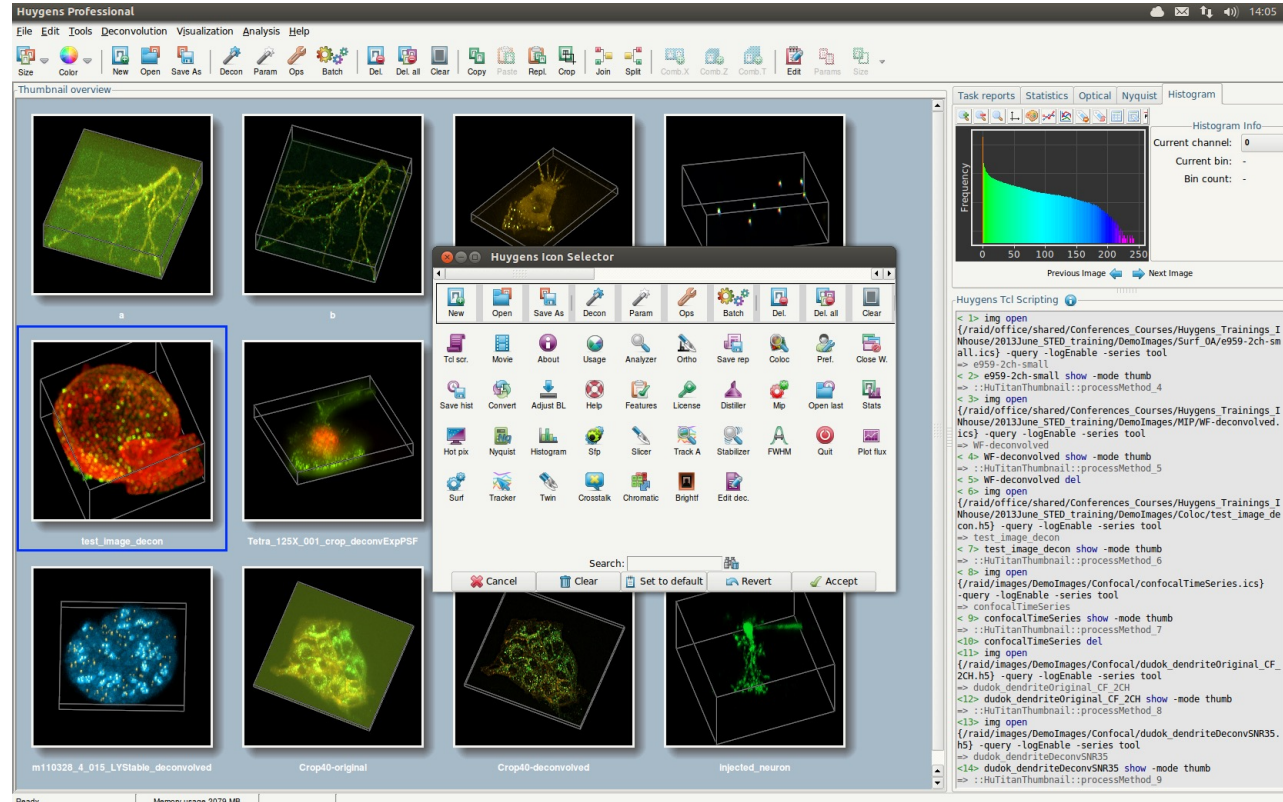


**Deconvolution of an HeLa cell acquired on a widefield microscope.**  
*Image courtesy Dr. Yury Belyaev. EMBL, Heidelberg, Germany.*

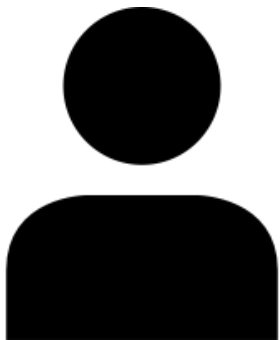
# General-purpose projects :: Huygens Remote Manager (HRM)



Scientific Volume Imaging  
Deconvolution - Visualization - Analysis



Huygens Professional



Single-user desktop  
application

Ponti A., Sevilla Sanchez D., Ehrenfeuchter N., Belyaev Y. G.I.T. Imaging & Microscopy 2:22 – 24. 2015  
Ponti A., Gulati A., Bäcker V. and Schwarb P. Imaging & Microscopy 9(2):57-58. 2007.

# General-purpose projects :: Huygens Remote Manager (HRM)

## Our scope

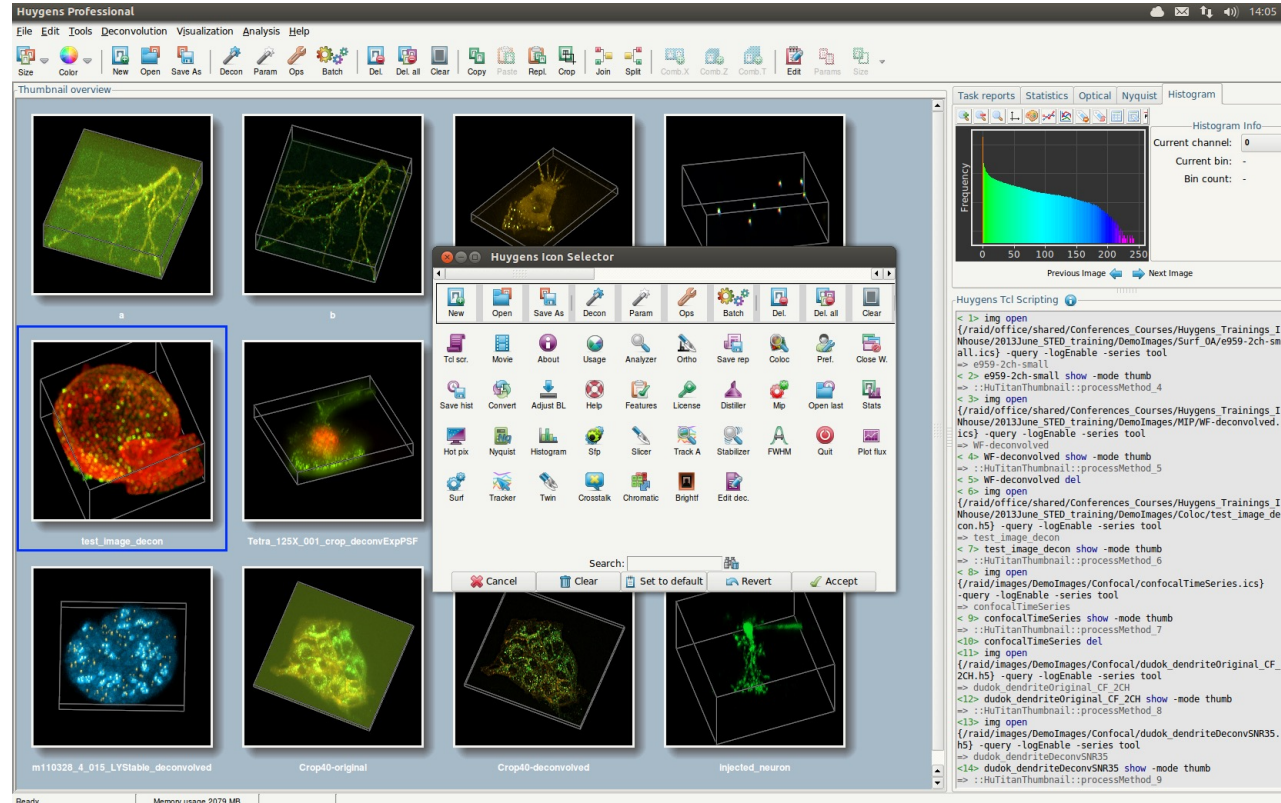
Scientific facility with around  
100 users



1 workstation



Scientific Volume Imaging  
Deconvolution - Visualization - Analysis



Huygens Professional



Ponti A., Sevilla Sanchez D., Ehrenfeuchter N., Belyaev Y. G.I.T. Imaging & Microscopy 2:22 – 24. 2015  
Ponti A., Gulati A., Bäcker V. and Schwarb P. Imaging & Microscopy 9(2):57-58. 2007.

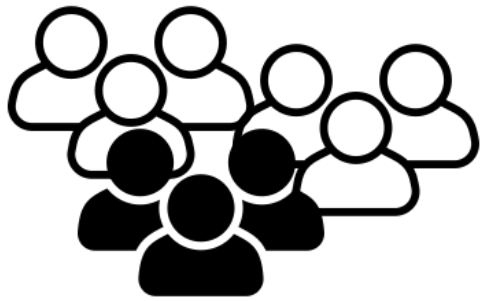
# General-purpose projects :: Huygens Remote Manager (HRM)

## Our scope

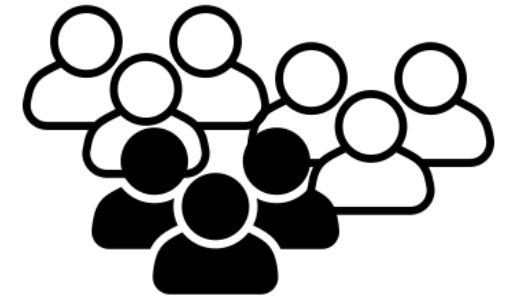
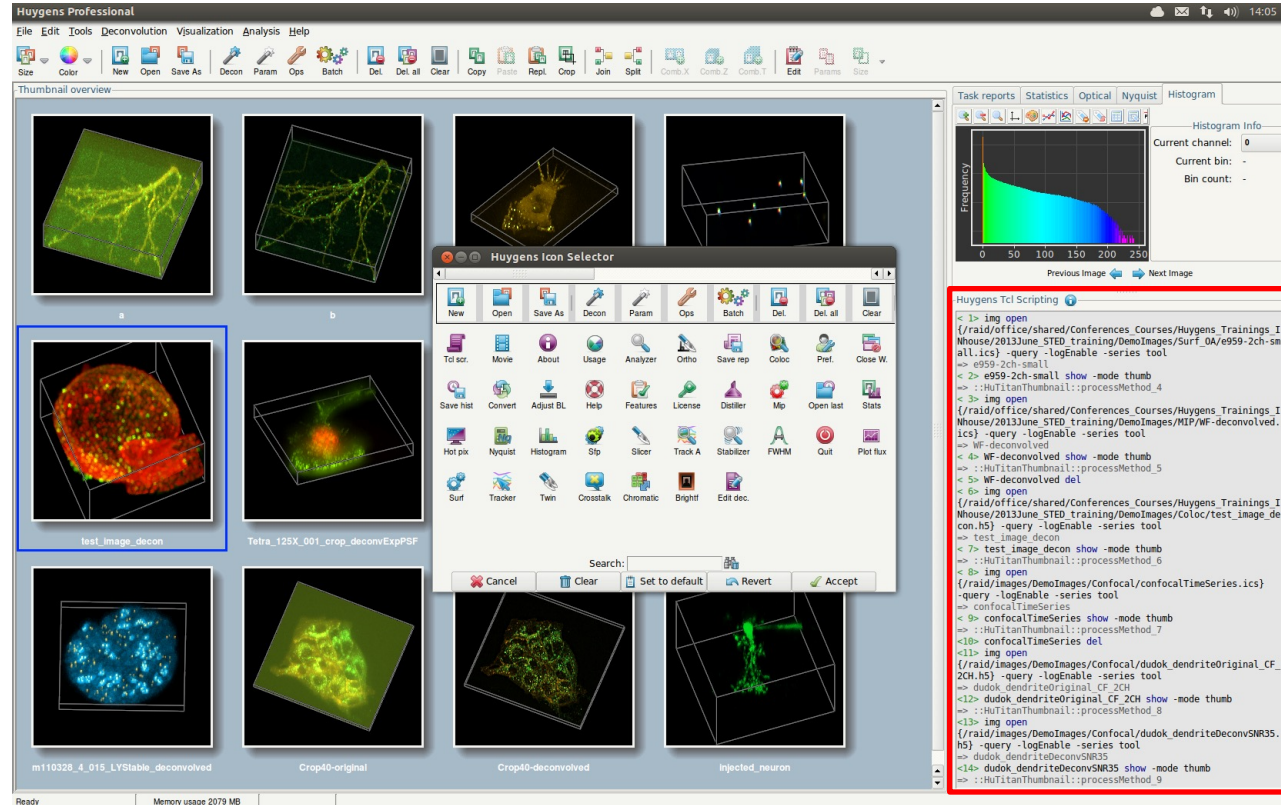
Scientific facility with around  
100 users



1 workstation



Scientific Volume Imaging  
Deconvolution - Visualization - Analysis



Fully scriptable

**Huygens Core**  
*Headless compute engine*

Huygens Professional

Ponti A., Sevilla Sanchez D., Ehrenfeuchter N., Belyaev Y. G.I.T. Imaging & Microscopy 2:22 – 24. 2015  
Ponti A., Gulati A., Bäcker V. and Schwarb P. Imaging & Microscopy 9(2):57-58. 2007.

# General-purpose projects :: Huygens Remote Manager (HRM)

**Huygens Remote Manager**

Help Resources User name Password Log in Reset

The *Huygens Remote Manager* is an easy to use interface to the *Huygens Software* by *Scientific Volume Imaging B.V.* that allows for multi-user, large-scale deconvolution and analysis.

collaboration

**MPI WBI**  
Montpellier RIO Imaging  
National Center for Scientific Research Montpellier

**FMI**  
Friedrich Miescher Institute for Biomedical Research  
Facility for Advanced Imaging and Microscopy  
Friedrich Miescher Institute

**EPFL**  
Bioimaging and Optics platform  
EPF Lausanne

**SVI**  
Scientific Volume Imaging  
Hilversum

**DBSSE**  
Single-Cell Facility  
ETH Zurich

**BIOZENTRUM**  
Universität Basel  
The Center for Molecular Life Sciences  
Imaging Core Facility  
Biozentrum University of Basel

**LIN**  
Leibniz Institute for Neurobiology  
Magdeburg

**cni**  
combinatorial neuroimaging  
Combinatorial Neuroimaging  
Magdeburg

**UNI FR**  
UNIVERSITÉ DE Fribourg  
UNIVERSITÄT Fribourg  
Bioimage | Light Microscopy Facility  
University of Fribourg

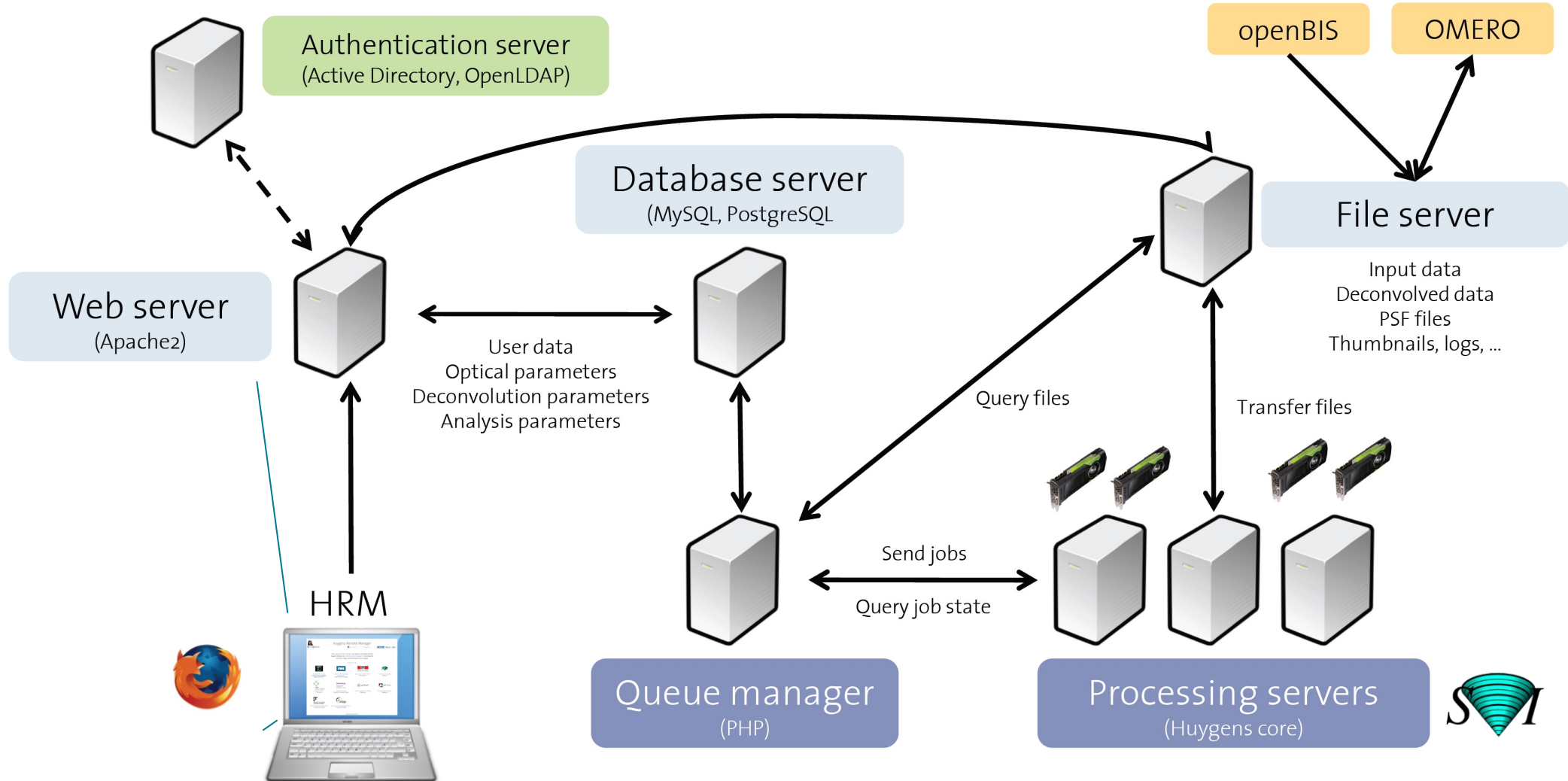
**miap**  
Microscopy and Image Analysis Platform  
University of Freiburg

**MANCHESTER**  
The University of Manchester  
BioImaging Facility  
University of Manchester

Huygens Remote Manager v3.9 Theme: dark light

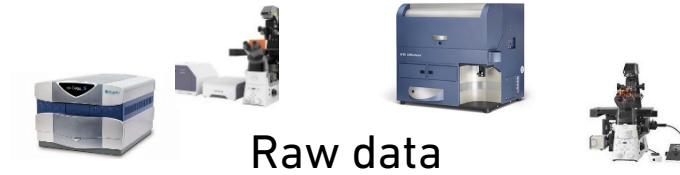
<https://github.com/aarpon/hrm/>

# General-purpose projects :: Huygens Remote Manager (HRM)



# General-purpose projects :: openBIS Importer Toolset (oBIT)

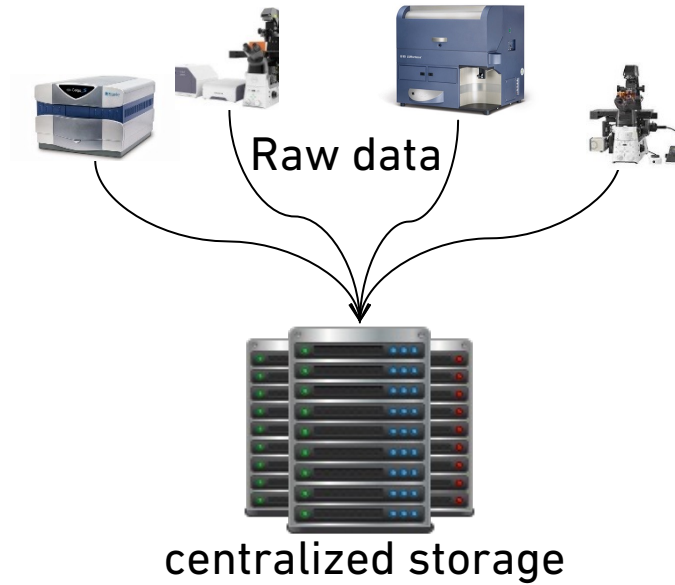
~ 40x at SCF/LAF





# General-purpose projects :: openBIS Importer Toolset (oBIT)

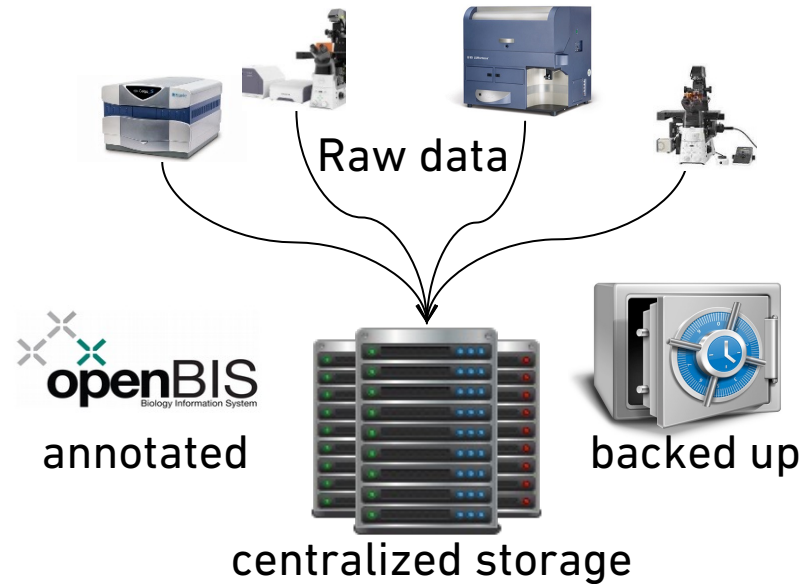
~ 40x at SCF/LAF



# General-purpose projects :: openBIS Importer Toolset (oBIT)

~ 40x at SCF/LAF

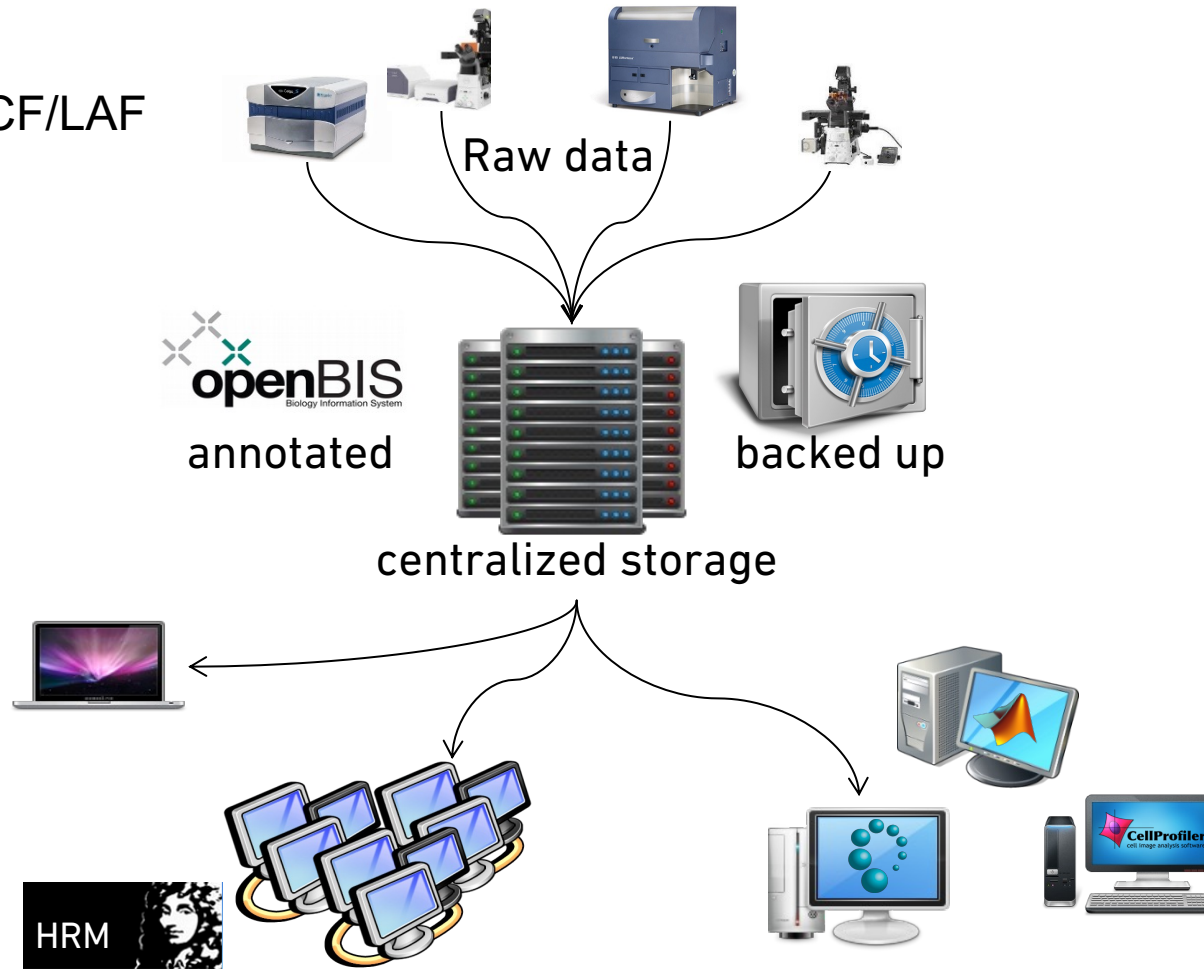
<https://sis.id.ethz.ch/>



# General-purpose projects :: openBIS Importer Toolset (oBIT)

~ 40x at SCF/LAF

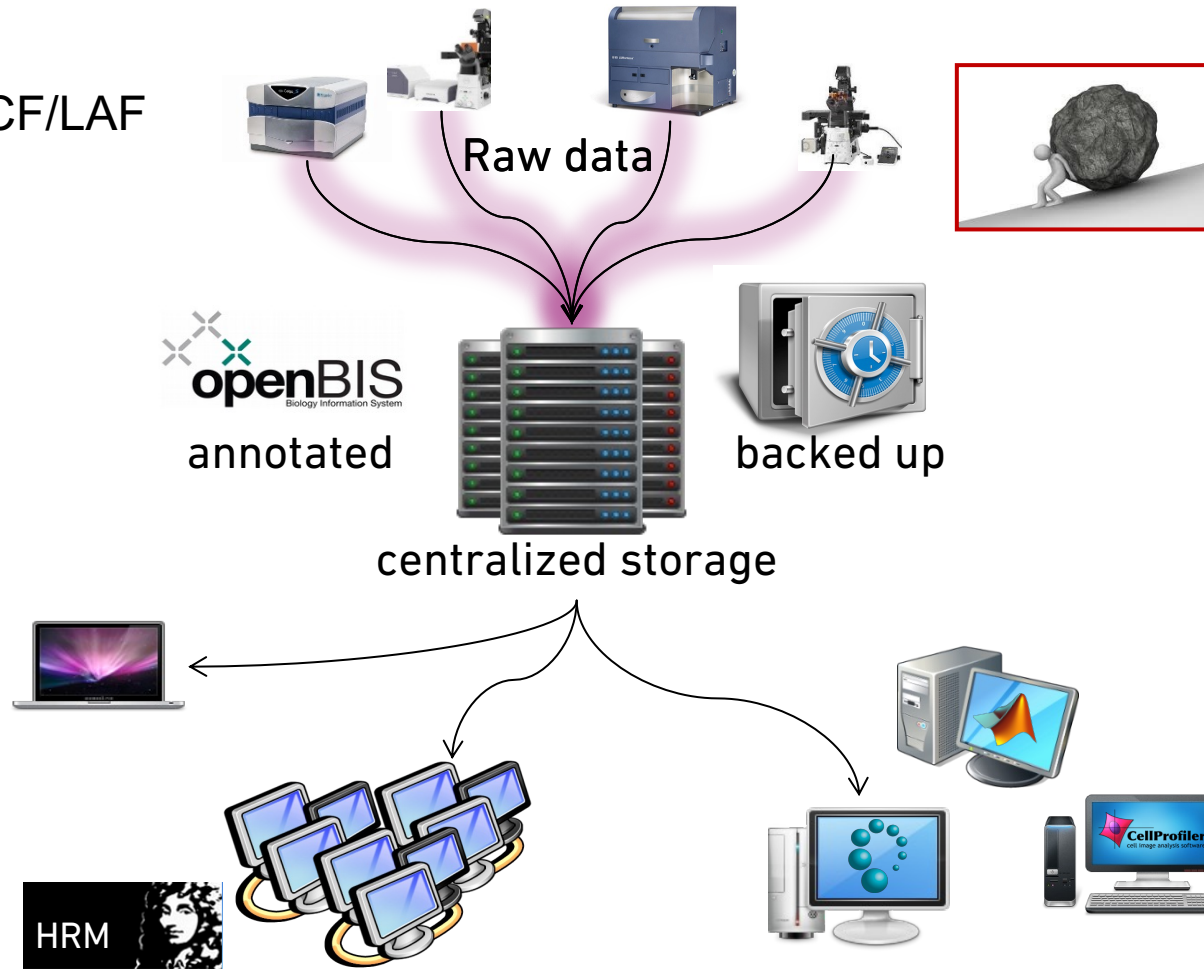
<https://sis.id.ethz.ch/>



# General-purpose projects :: openBIS Importer Toolset (oBIT)

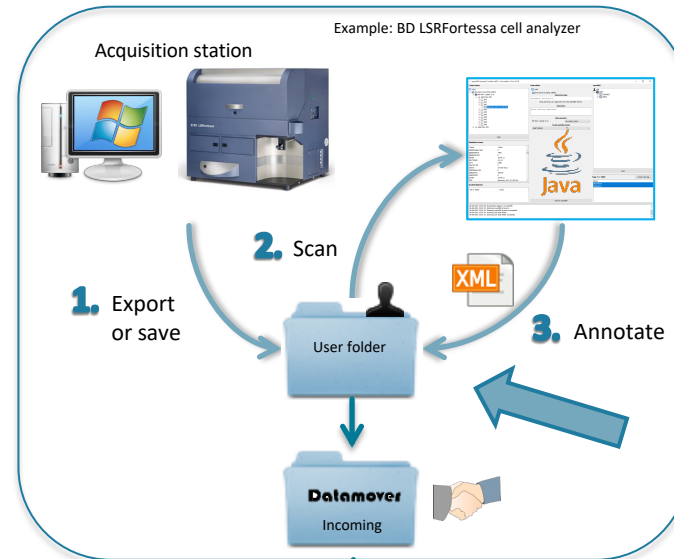
~ 40x at SCF/LAF

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



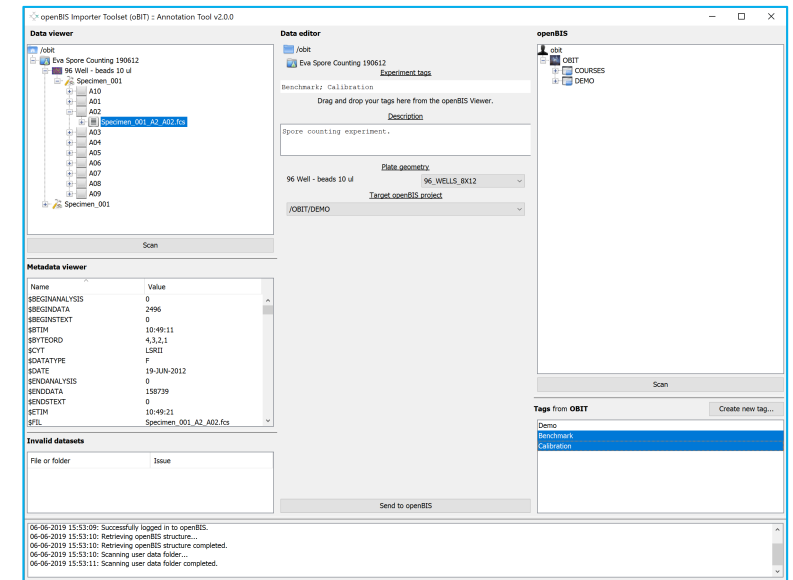
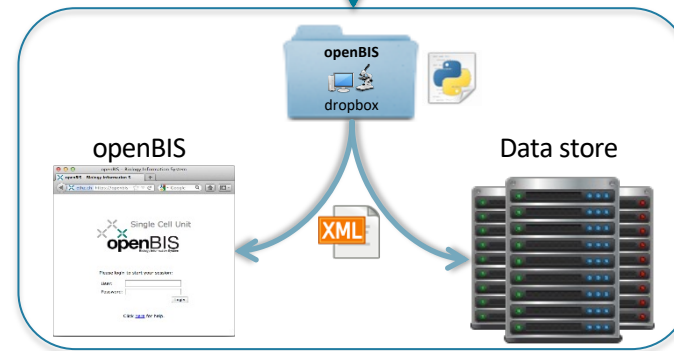
# General-purpose projects :: openBIS Importer Toolset (oBIT)

New openBIS core technologies  
Microscopy  
Flow Cytometry



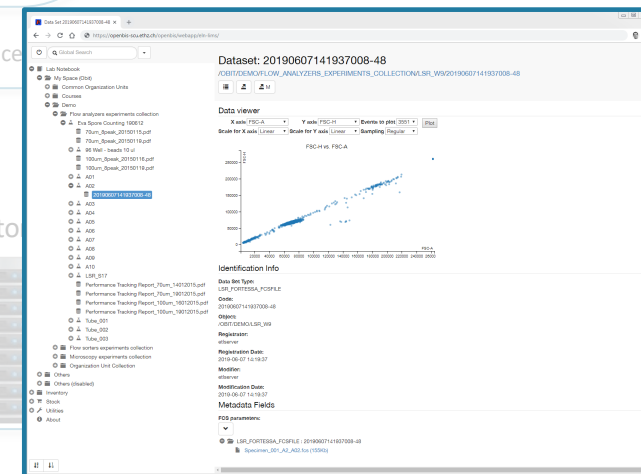
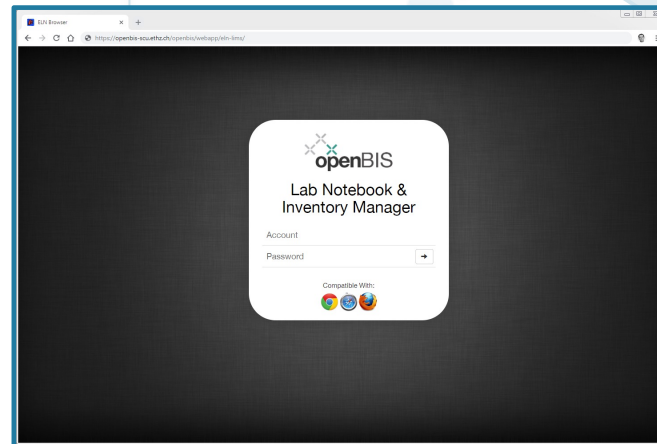
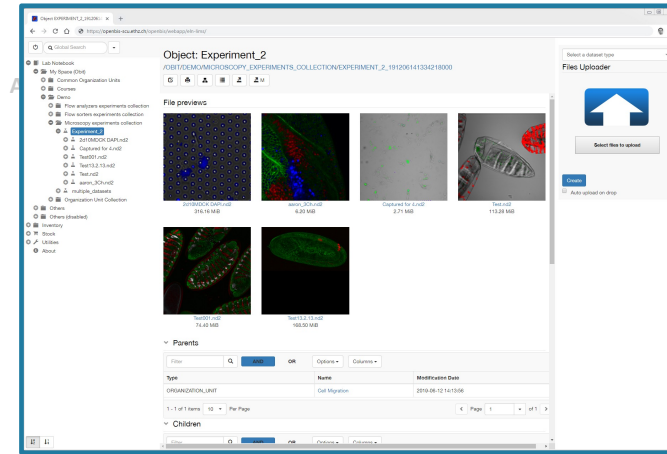
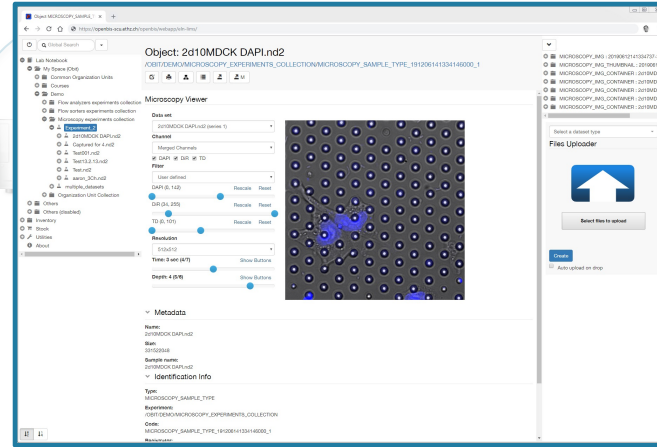
<https://github.com/aarpon/obit>

**Datamover** as a Windows service



# General-purpose projects :: openBIS Importer Toolset (oBIT)

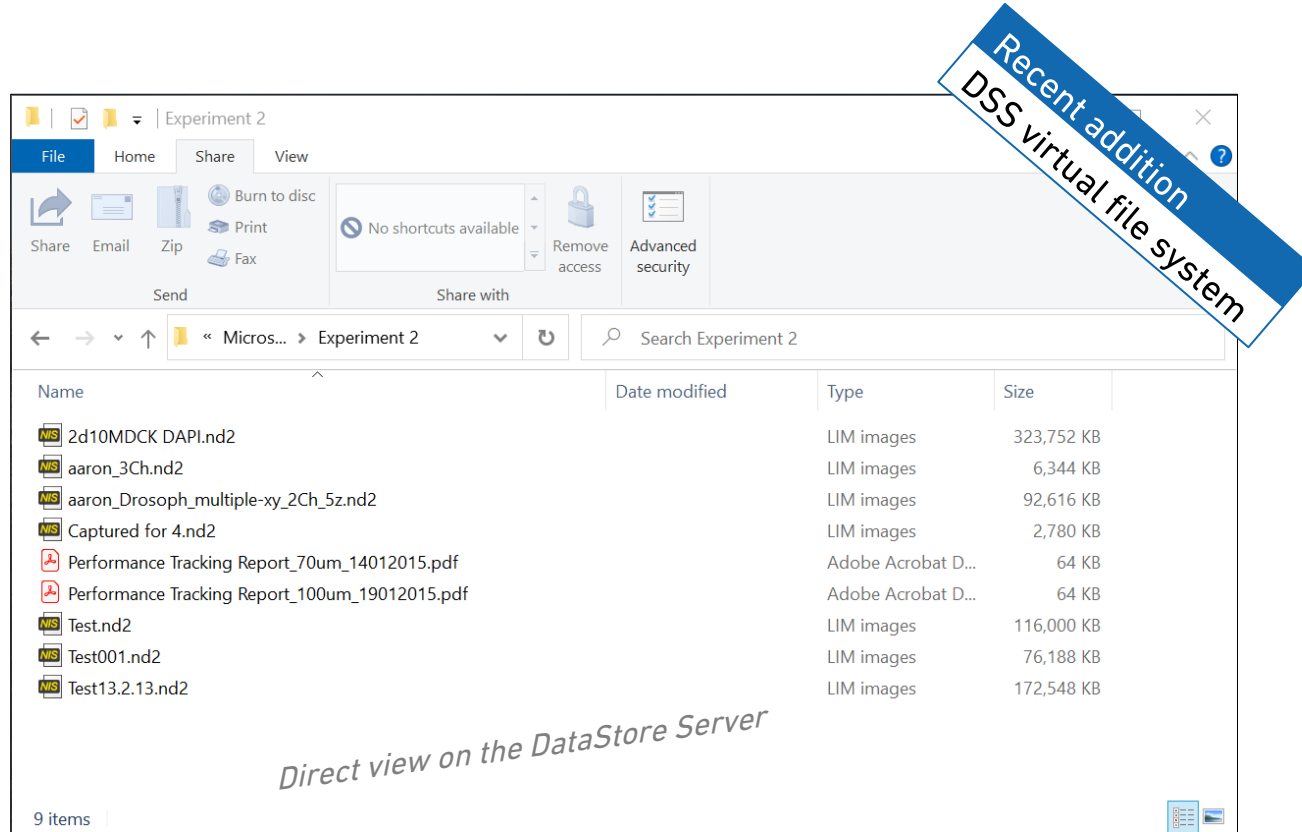
New openBIS core technologies  
Microscopy  
Flow Cytometry



Integration into ELN



# General-purpose projects :: openBIS Importer Toolset (oBIT)



**ETH zürich**  
Scientific IT Services

# Expert-driven projects

	Expert-driven projects
Target audience	Several users
Specificity	Specialized but scalable
Team composition	Me and field experts (with user feedback)
Problem focus	Niche problems
Complexity	Complex
Scalability	Moderate
User input	Initial and iterative (at release)
Project time	Moderate to long
Resource allocation	High
Field knowledge	Very high (experts)
Code quality/testing	High



# Expert-driven projects

	Expert-driven projects
Target audience	Several users
Specificity	Specialized but scalable
Team composition	Specialized
Problem focus	Well-defined problems
Complexity	High
Scalability	Moderate
User input	Initial and iterative (at release)
Project time	Moderate to long
Resource allocation	High
Field knowledge	Very high (experts)
Code quality/testing	High

**Three example projects**

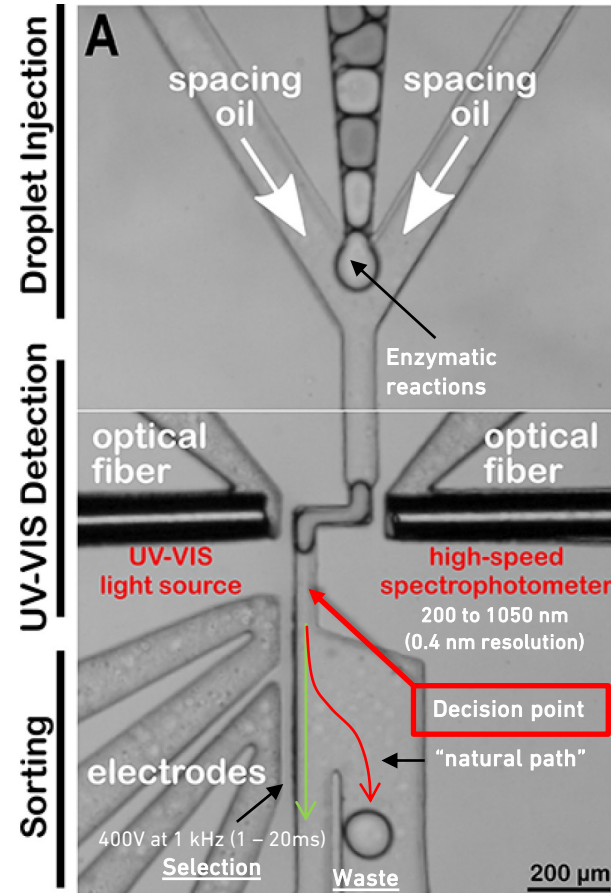
SpectraSorter  
pyPOCQuant  
pyMINFLUX

## Expert-driven projects :: SpectraSorter

Duncombe T. A., Ponti A., Dittrich P. S. SoftwareX, Volume 19, 2022, 101160 DOI:10.1016/j.softx.2022.101160  
 Duncombe T. A., Ponti A., Seebeck F. P., Dittrich P. S. 2021. DOI: 10.1021/acs.analchem.1c02822

## Goal

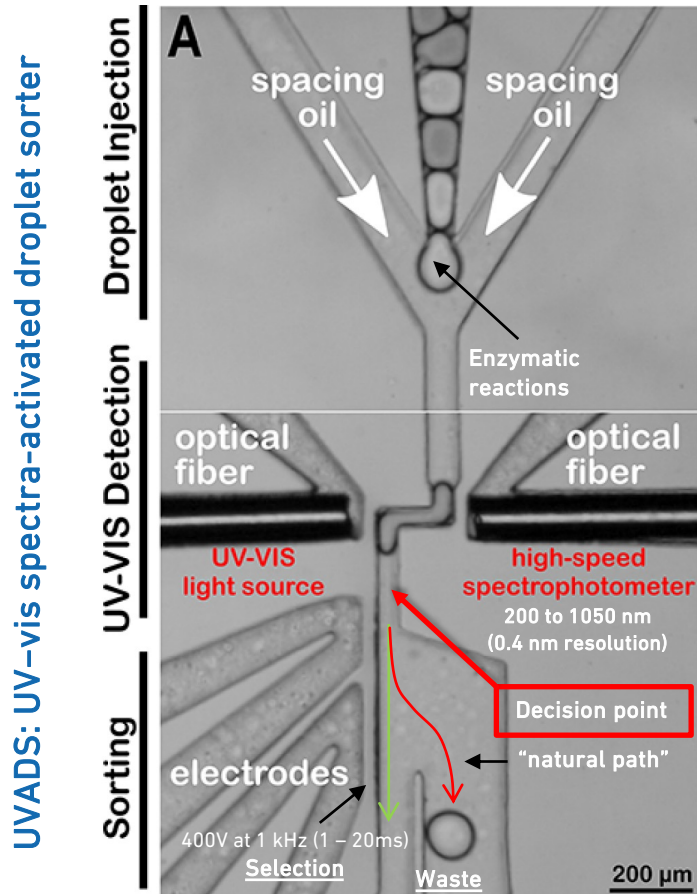
Design a **microfluidic platform** for the **high-throughput analysis** of (enzymatic) **reactions** inside small **droplets**.



Expert: **Todd Duncombe**, Petra Dittrich Group, D-BSSE ETHZ

# Expert-driven projects :: SpectraSorter

Duncombe T. A., Ponti A., Dittrich P. S. SoftwareX, Volume 19, 2022, 101160 DOI:10.1016/j.softx.2022.101160  
 Duncombe T. A., Ponti A., Seebeck F. P., Dittrich P. S. 2021. DOI: 10.1021/acs.analchem.1c02822



In **droplet microfluidic platforms**, droplets containing cells or reagents flow through capillaries. Inside each droplet, a reaction takes place, and the output is measured *by some means*. These platforms aim at achieving high-throughput and/or massively parallel analytics.

Previous approaches were very low throughput (1-10 droplets / second) and used fluorescence microscopy to measure single-wavelength absorbance in the visible spectrum.

Recent **label-free UV-vis spectroscopy** interrogates **molecular structures** directly by **chemical absorbance** of incident light over a large spectrum (200 to 1050 nm)

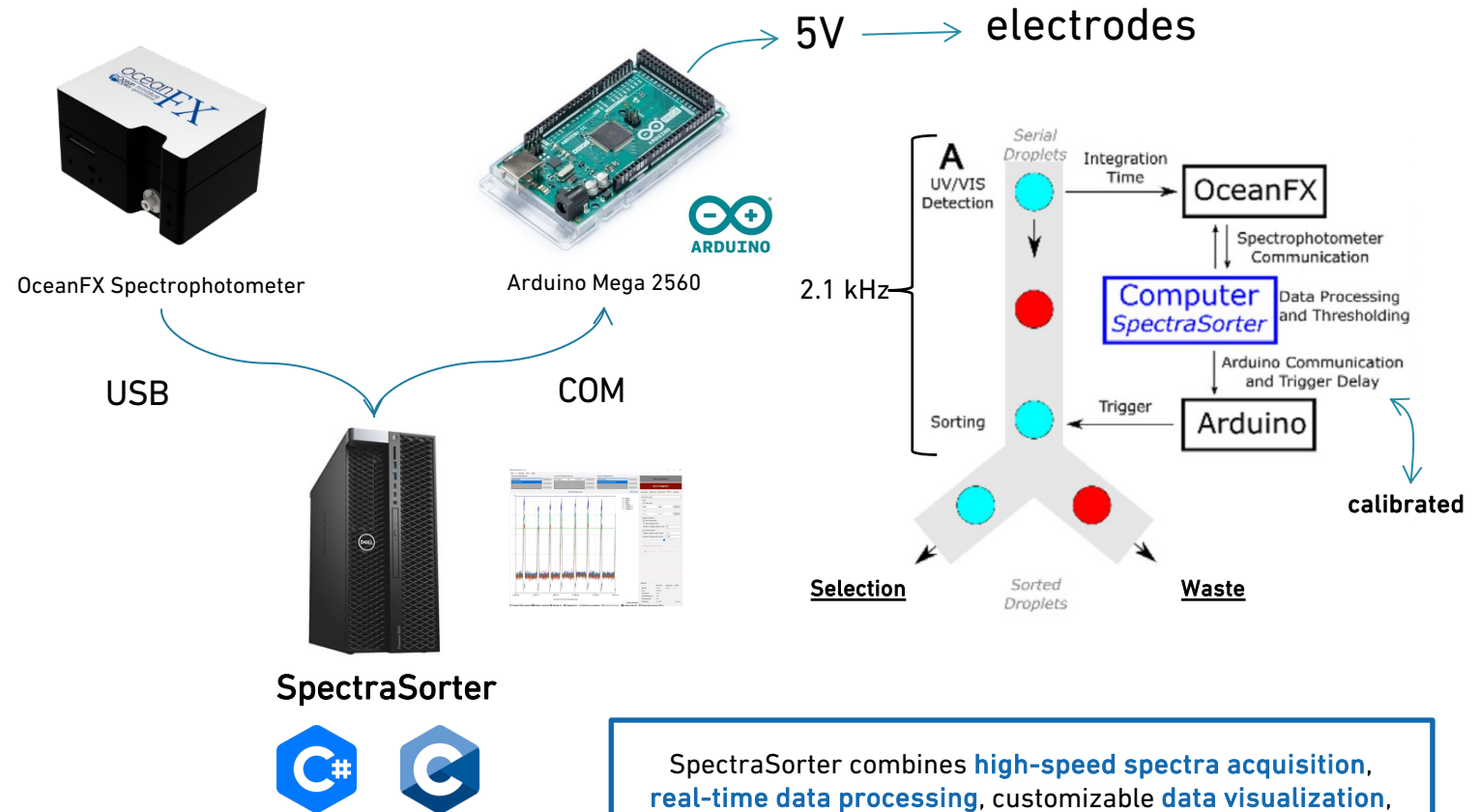
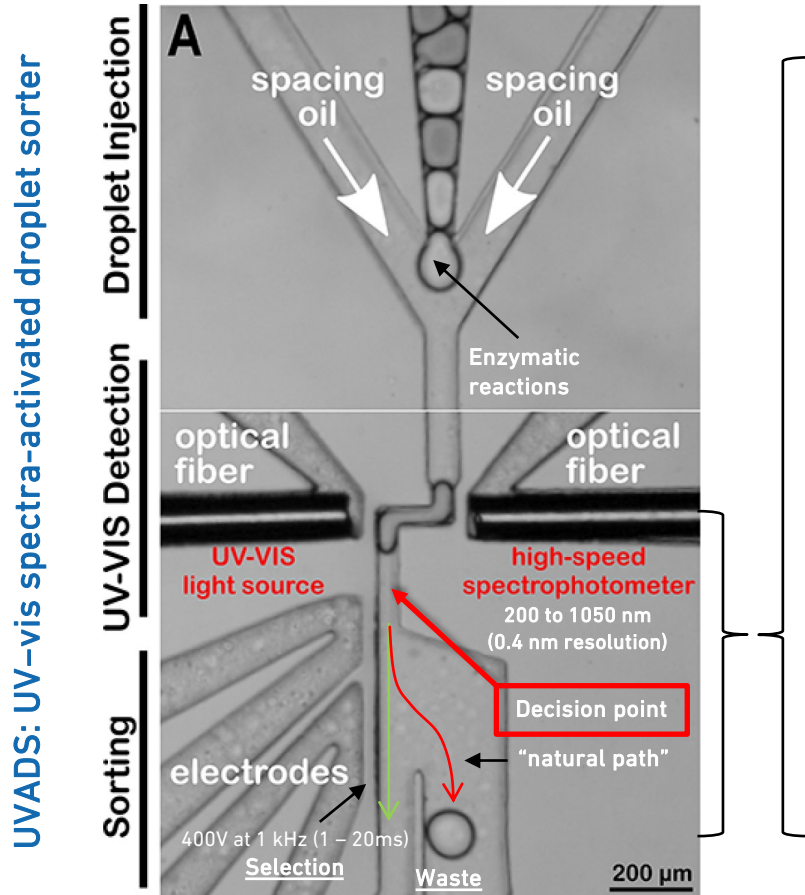
Our **microfluidic platform (UVADS)** can perform screening assays of enzymatic activity in **droplets** of bacterial micro-colonies by directly measuring the conversion to product by **UV-vis spectroscopy** and **sort** selected droplets using **electrodes**.

High-throughput label-free chemical identification in droplets and **on-demand** collection (via electrodes)

# Expert-driven projects :: SpectraSorter

Duncombe T. A., Ponti A., Dittrich P. S. SoftwareX, Volume 19, 2022, 101160 DOI:10.1016/j.softx.2022.101160  
 Duncombe T. A., Ponti A., Seebeck F. P., Dittrich P. S. 2021. DOI: 10.1021/acs.analchem.1c02822

<https://github.com/SpectraSorter/SpectraSorter>

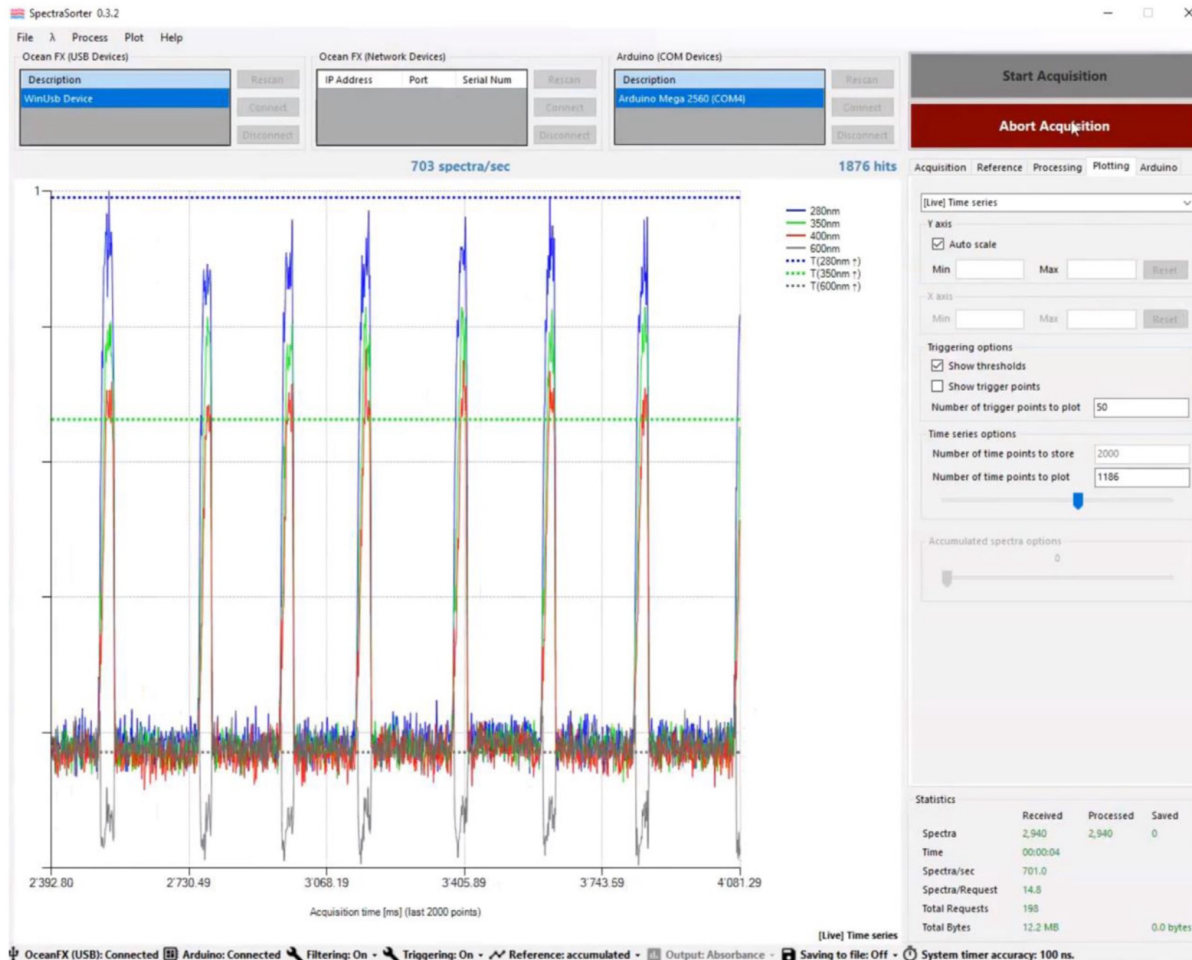


SpectraSorter combines **high-speed spectra acquisition**, **real-time data processing**, **customizable data visualization**, **precise control of external triggering** using any **multi-spectral** feature, **configuration management**, and more.

High-throughput label-free chemical identification in droplets and **on-demand** collection (via electrodes)

# Expert-driven projects :: SpectraSorter

Duncombe T. A., Ponti A., Dittrich P. S. SoftwareX, Volume 19, 2022, 101160 DOI:10.1016/j.softx.2022.101160  
 Duncombe T. A., Ponti A., Seebeck F. P., Dittrich P. S. 2021. DOI: 10.1021/acs.analchem.1c02822

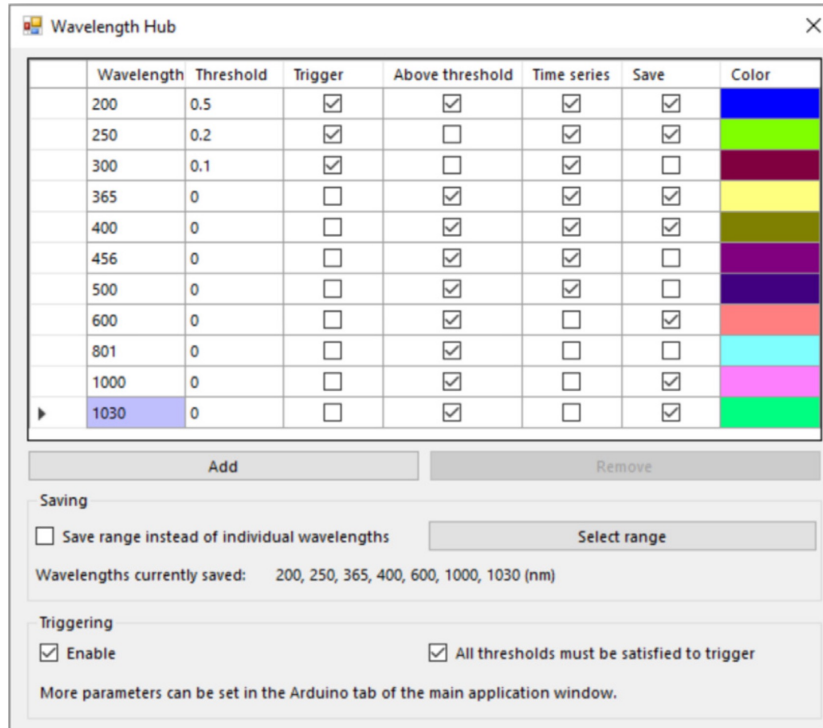


## SpectraSorter runs **four parallel queues**:

- **Acquisition** queue: collects up to 4500 spectra/s from the spectrometer
- **Compute** queue: performs all operations to decide if an event should trigger the Arduino microcontroller (low-pass filtering, transformation for absorbance or transmission, dark- and reference-correction, testing against all user-defined thresholds, triggering via Arduino)
- **Plotting** queue: plots the last spectrum at very low rate (10 Hz) for visual feedback
- **Saving** queue: writes to file the processed wavelengths (either a selection, a range, or the full spectrum)

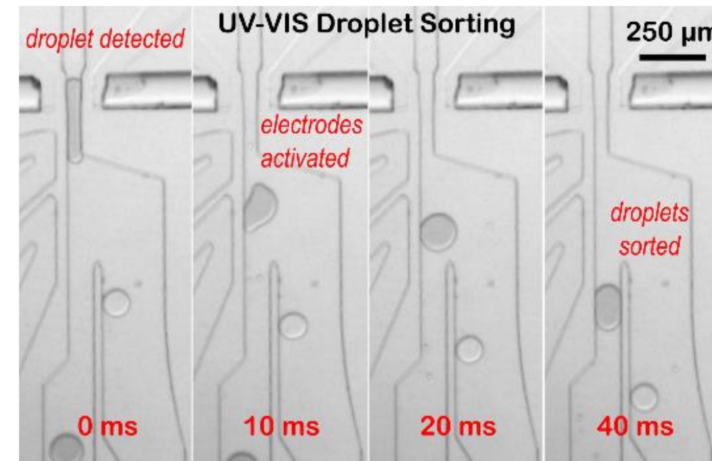


# Expert-driven projects :: SpectraSorter



The **Wavelength Hub** allows selection of any number of wavelengths that:

- act as a **threshold** for a **triggering** event (and in which way)
- will be displayed in the plotter as **time series** and/or as **full spectra**
- will be **saved** to disk



# Expert-driven projects :: SpectraSorter

## Expert contributions

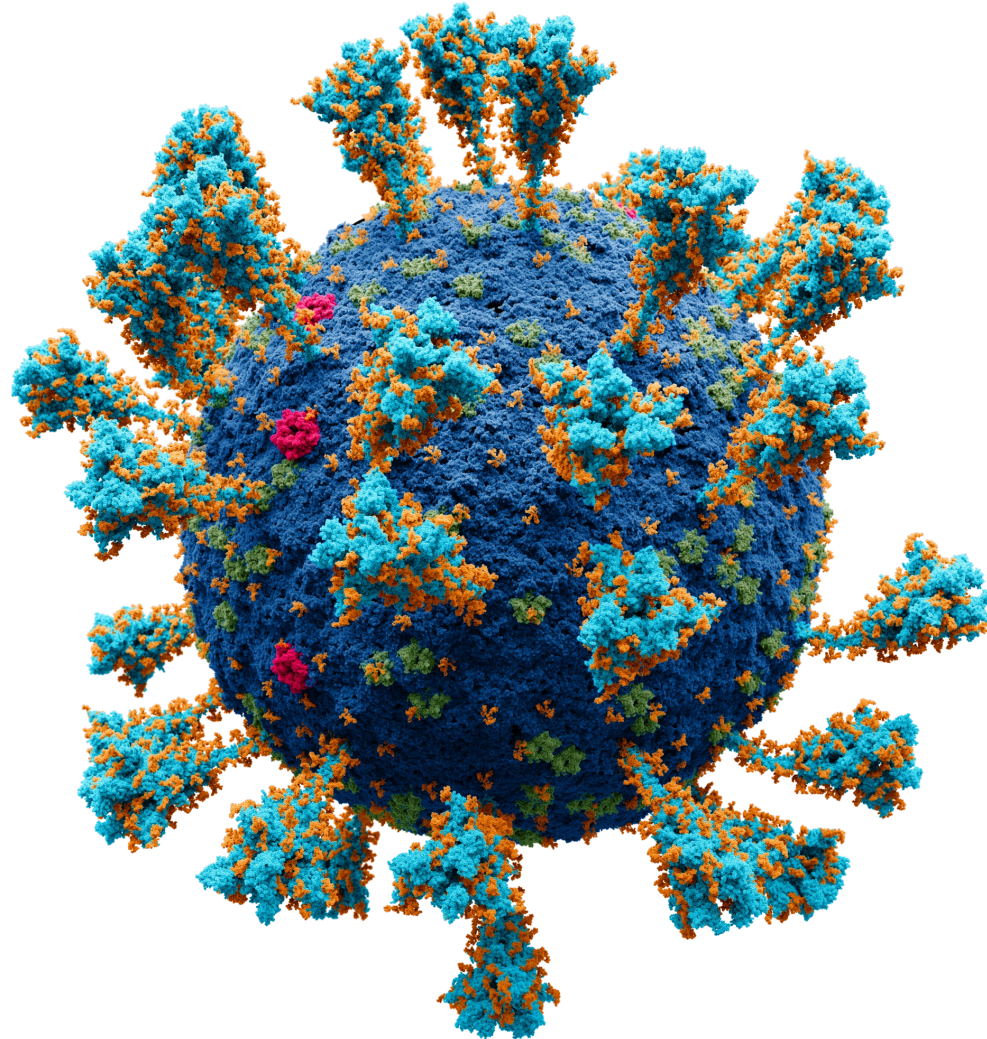
**Todd Duncombe**: development of the UVADS methodology and the microfluidics chip

**Aaron Ponti**: software development

## Users

Petra Dittrich group

# Expert-driven projects :: pyPOCQuant



<https://en.wikipedia.org/wiki/SARS-CoV-2>



# Expert-driven projects :: pyPOCQuant

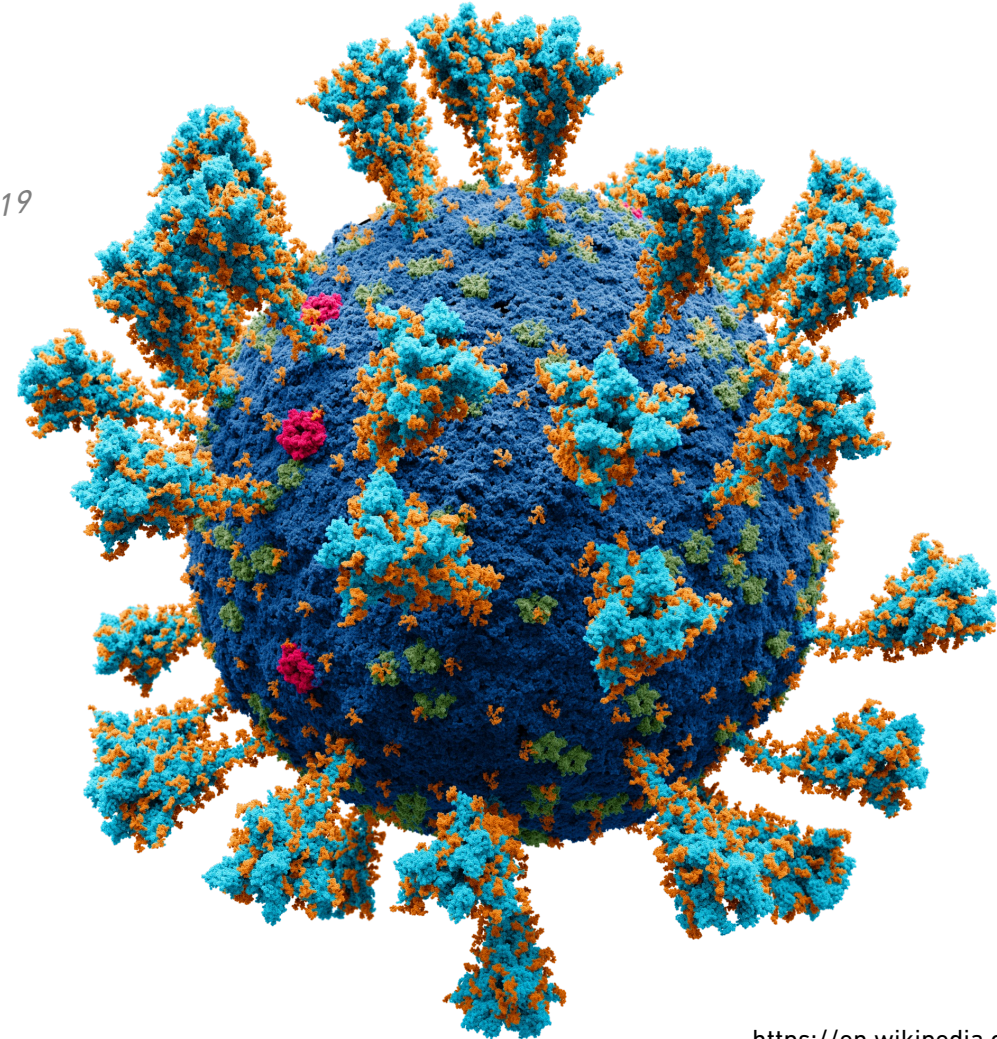
On request by  
Baselland Test Center (via Biolytix AG)



## Requirements

1. **Quantify** the levels of **IgG** and **IgM** antibodies in patients' blood samples **over time** to study the body's immune response to SARS-CoV-2
2. **Associate patient metadata** to each analysis and store the results in a database

*First wave of COVID-19  
Spring 2020*



## Experts:

**Fabian Rudolf**, Bundesamt für Gesundheit, CH

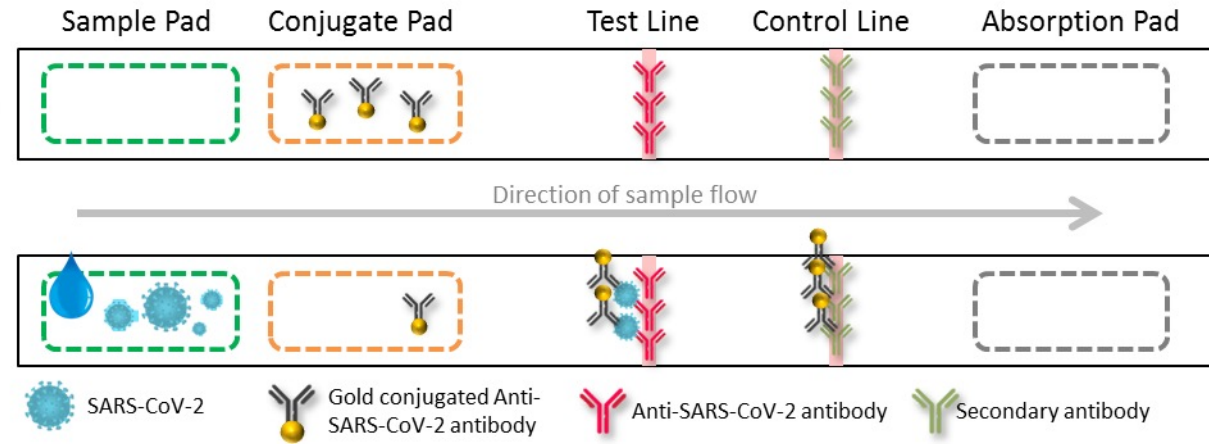
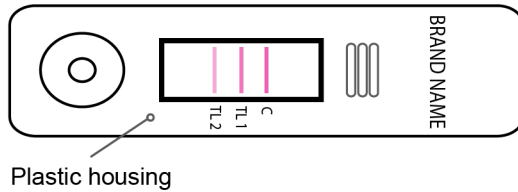
**Andreas Cuny**, Joerg Stelling group, D-BSSE ETHZ

<https://en.wikipedia.org/wiki/SARS-CoV-2>

# Expert-driven projects :: pyPOCQuant

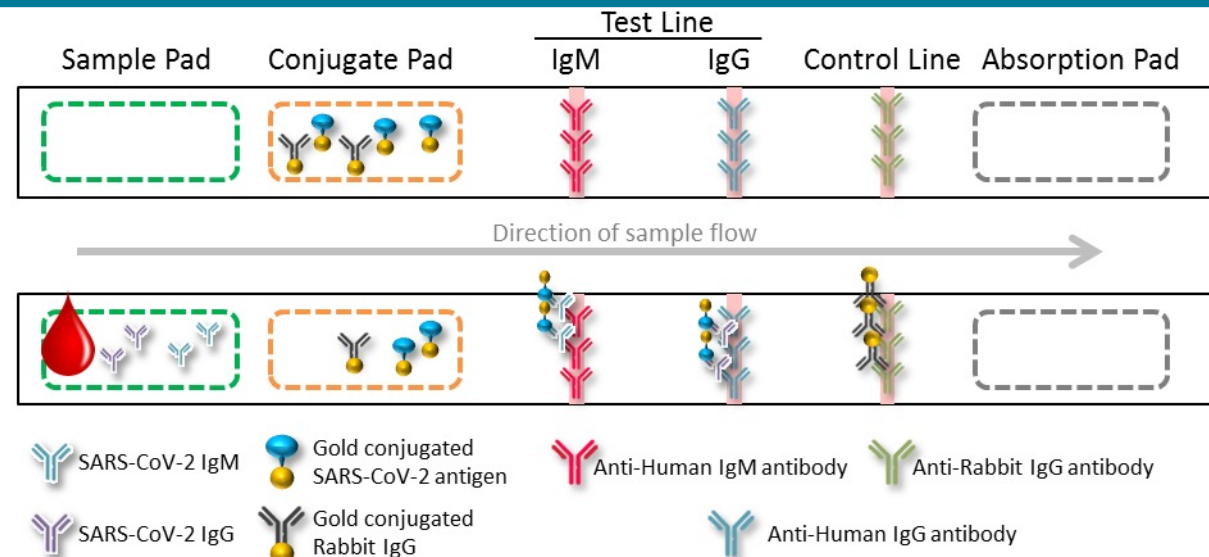
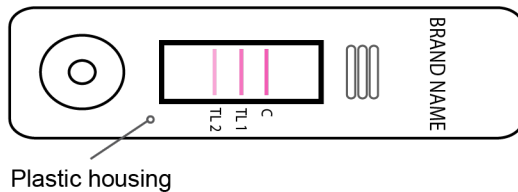
Image source: <https://www.acebiolab.com/EN/news/44>

## Antigen (Ag) test



## Lateral-flow point-of-care tests (POCTs)

## Antibody (Ab) test



Lateral flow Point-Of-Care Tests (POCTs) are a valuable tool for rapidly detecting pathogens and the associated immune response in humans and animals.

# Expert-driven projects :: pyPOCQuant

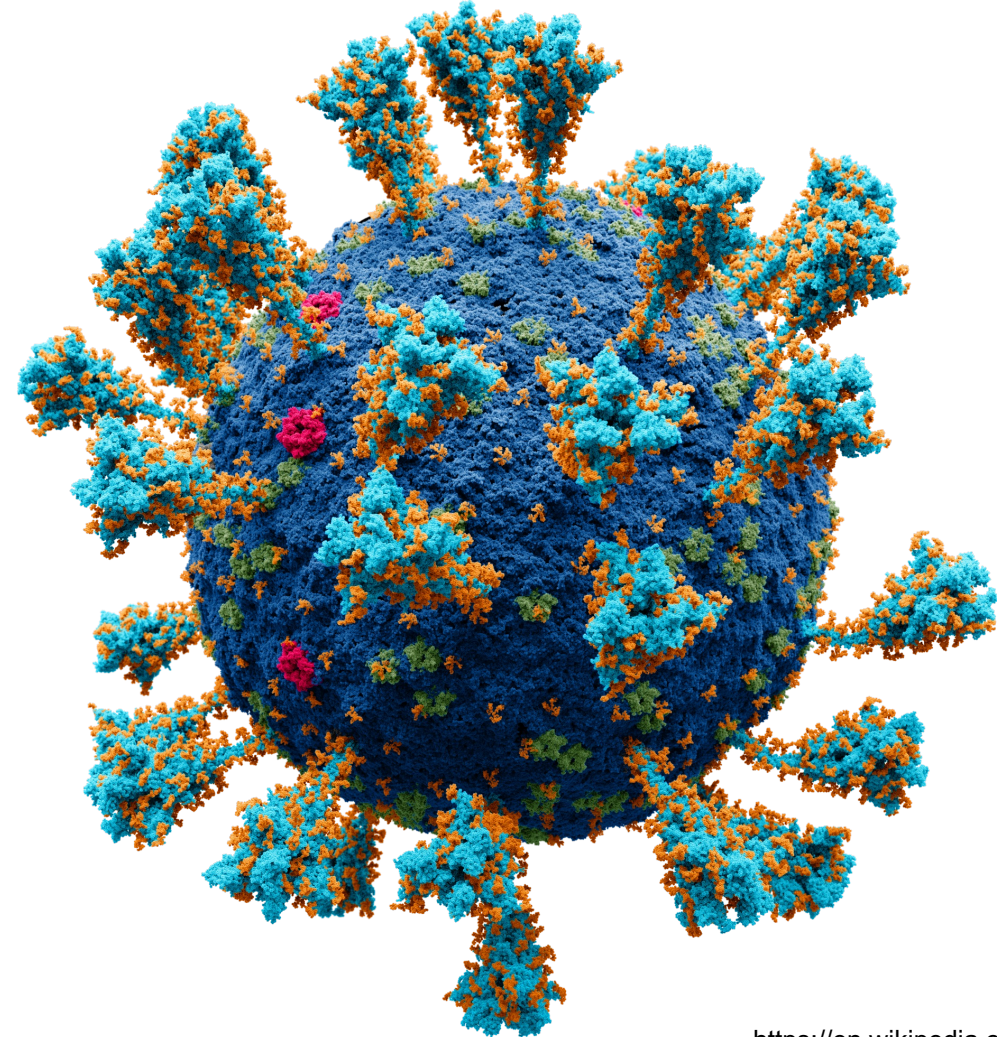
Sudden, high demand for **robust** and **quantitative** analysis of POCTs from **different vendors** and from **large numbers** of images

*We don't want to waste any test!*

*Which ones are quantitative?*

**No freely available pipelines for robust, large-scale batch processing**

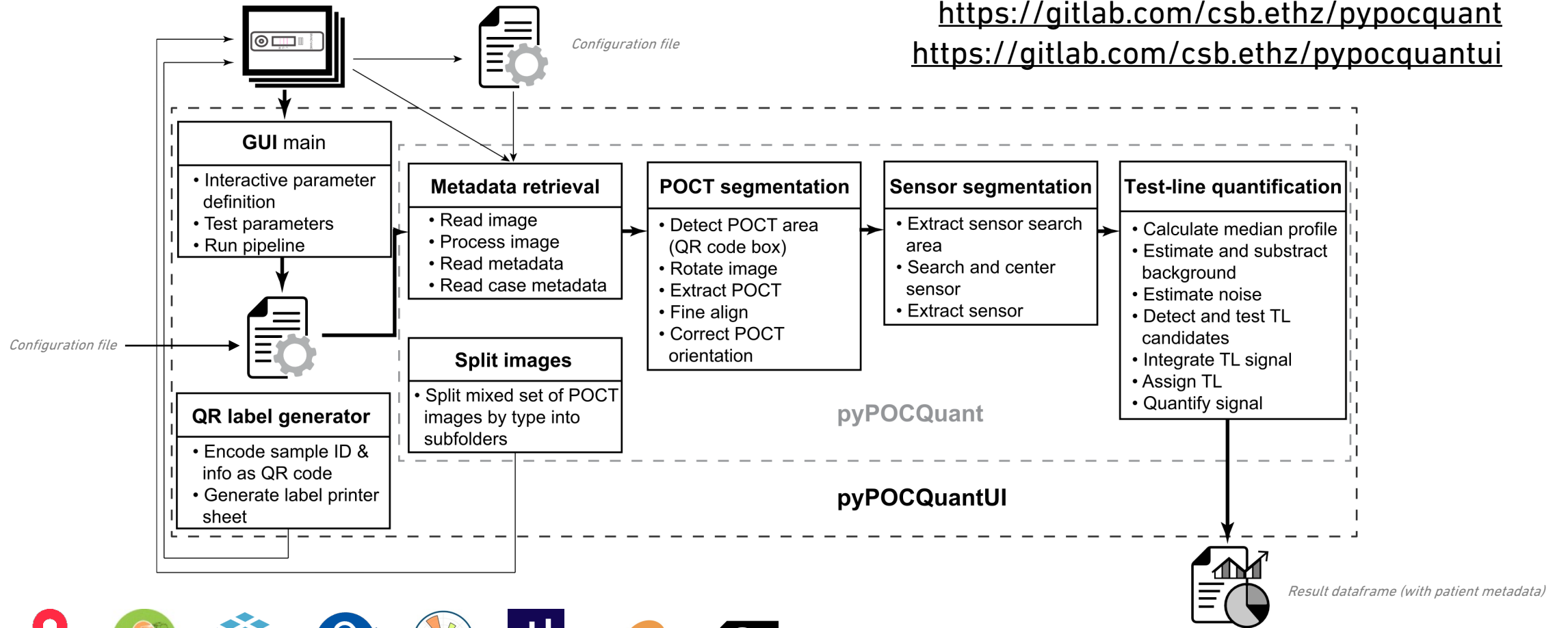
*There were commercial readers, but they were slow, expensive, proprietary, and only compatible with certain types of tests.*



<https://en.wikipedia.org/wiki/SARS-CoV-2>

# Expert-driven projects :: pyPOCQuant

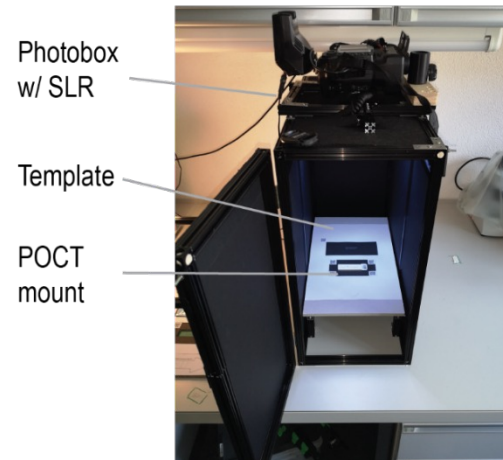
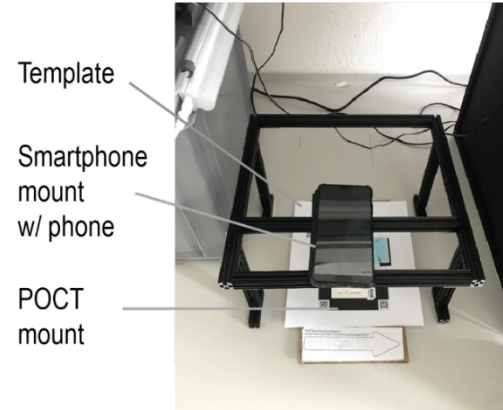
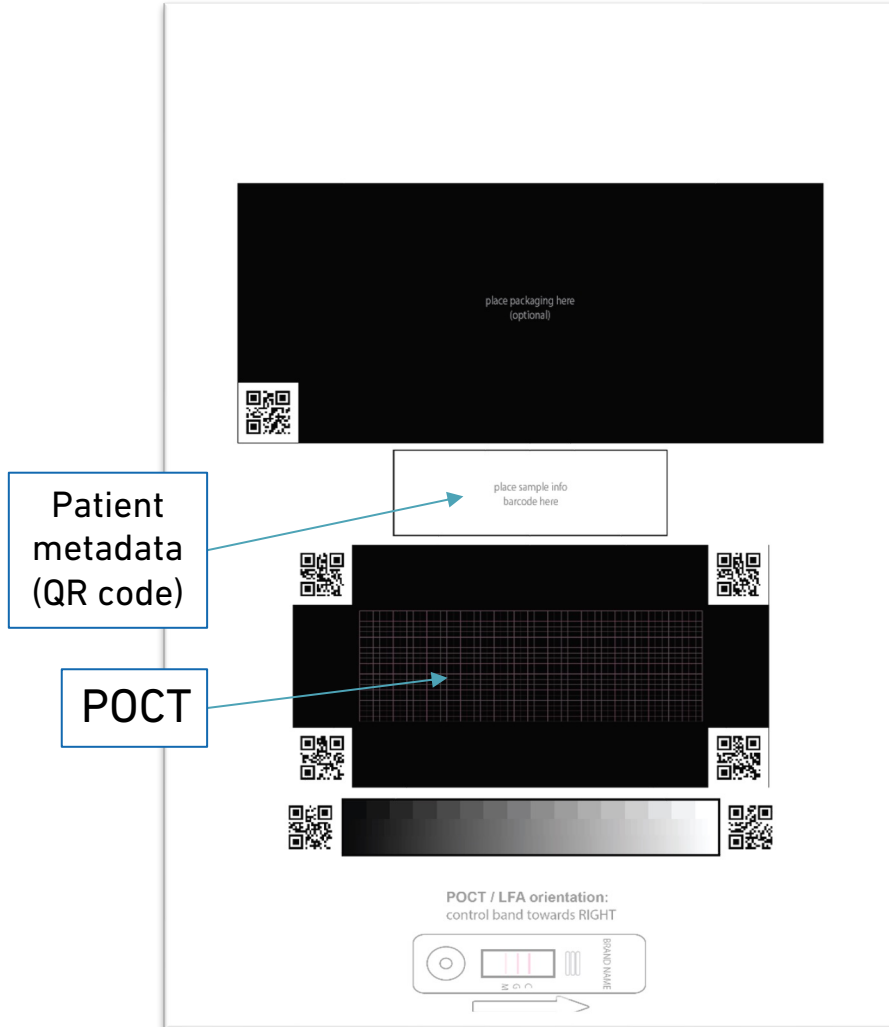
<https://gitlab.com/csb.ethz/pypocquant>  
<https://gitlab.com/csb.ethz/pypocquantui>



# Expert-driven projects :: pyPOCQuant

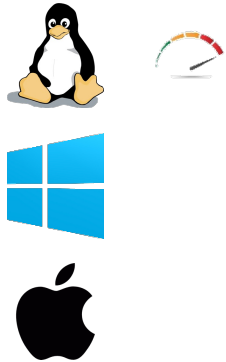
- We needed to be able to analyse batches of **hundreds to thousands of tests** with very low failure rate
- However, several parameters strongly affect the robustness of our quantitative analysis:
  - **illumination variability** (given by acquisitions performed at different times of the day, weather conditions, shadows, reflections)
  - **scaling** and **orientation** of the images (due to freehand shots with cameras or smartphones)
  - **intensity corrections** by the camera (intensity stretching, white balancing, and compression) and its file formats (*e.g.*, RAW vs. JPG)
  - **different sizes, colors, and shapes** of POCTs from different vendors
  - **random positioning** and **orientation** of the POCT in the field of view
  - often **difficult localization** of the detector window

# Expert-driven projects :: pyPOCQuant



Work by [Andreas Cuny](#)

# Expert-driven projects :: pyPOCQuant



Interactively create and save a configuration for batch processing

The screenshot shows the pyPOCQuant application window. On the left, there are file selection fields (1, 2) and a file list (3). Below that is a parameter configuration table (10) with sections for Runtime parameters, Basic parameters, and Sensor band names. A red arrow points to the 'Number of sensor bands' field (8). At the bottom, there are 'Test parameters' (9) and a 'Run' button (11). The main area shows two images: the top one (6) has a red test line (7) drawn on it, and the bottom one (4) shows the extracted POCT result. A status bar at the bottom indicates 'Extracting POCT from image finished successfully' (5). A log window at the very bottom shows timestamps and messages.

User defines window and test line positions (and number)

Automatic extraction of POCT

# Expert-driven projects :: pyPOCQuant

User defines window and test line positions (and number)



Inter  
con  
bat



# Expert-driven projects :: pyPOCQuant

## Run from console

```
python pyPOCQuant.py -f examples/images -o examples/images/results -s examples/config.conf -w 4
```

## Run from scripts or notebooks

```
from pypocquant.lib.pipeline import run_pipeline
from pypocquant.lib.settings import default_settings

# Get the default settings
settings = default_settings()

# Change settings manually as needed
settings["sensor_band_names"] = ('igm', 'igg', 'ctl')

# Alternatively, Load existing settings file
# from pypocquant.lib.settings import load_settings
# settings = load_settings('full/path/to/settings/file.conf')

# Set final argument
input_folder_path = 'full/path/to/input/folder'
results_folder_path = 'full/path/to/results/folder'
max_workers = 8

# Run the pipeline
run_pipeline(
    input_folder_path,
    results_folder_path,
    **settings,
    max_workers=max_workers
)
```

## Run from user interface

The screenshot shows the pyPOCQuant user interface with several numbered annotations:

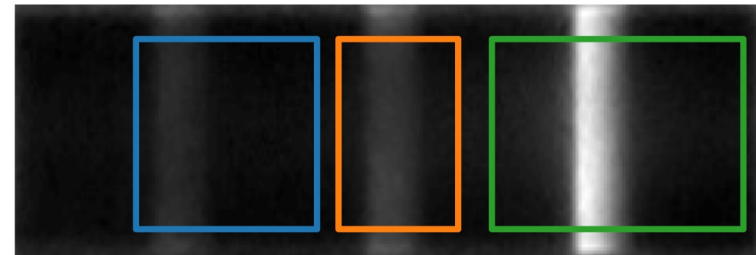
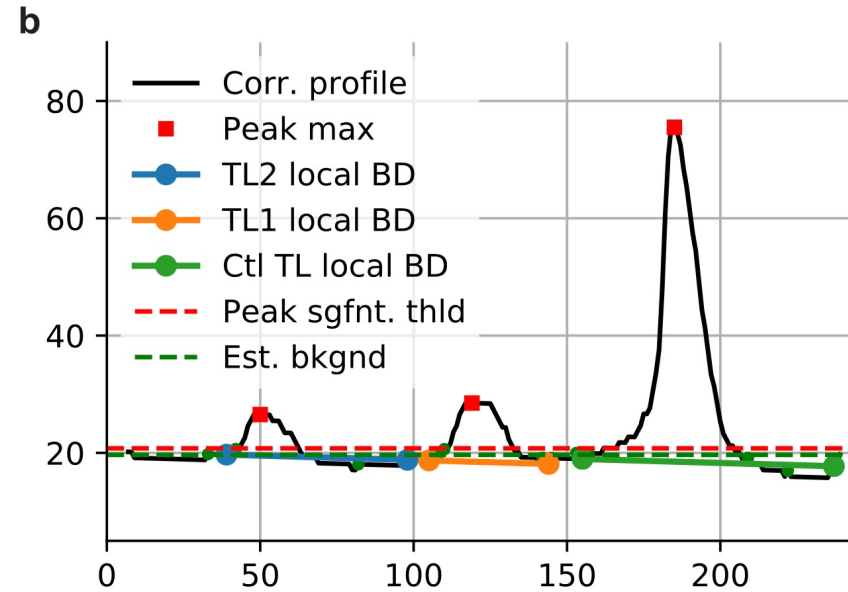
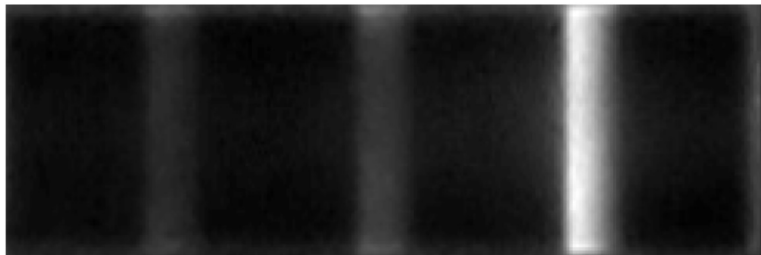
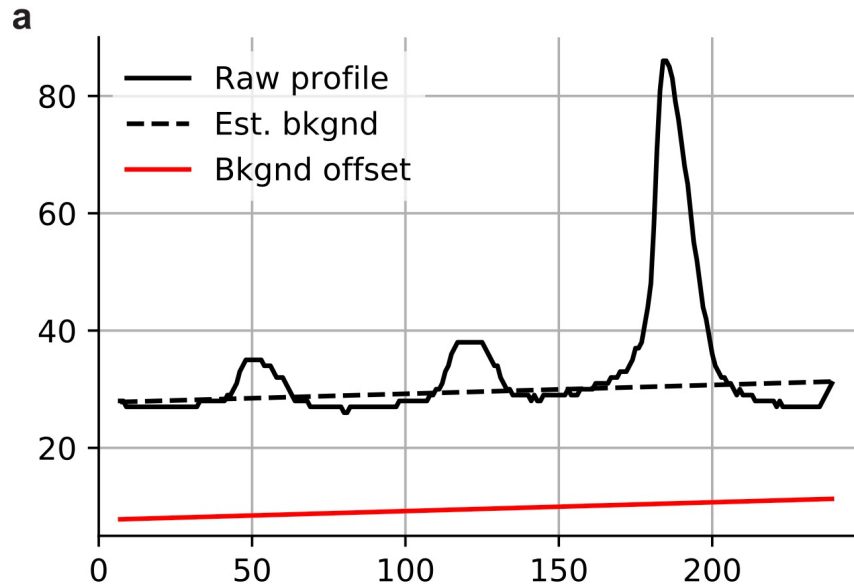
- 1: File menu
- 2: Select the output folder field
- 3: File list showing IMG\_8489.JPG and Img0619.jpg
- 4: Image of a COVID-19 test strip
- 5: Run button
- 6: Input folder field
- 7: Red arrow pointing to a sensor band on the test strip image
- 8: Band expected relative location parameter
- 9: Test parameters section
- 10: Number of cores parameter
- 11: Run button

Parameter	Value
<b>Runtime parameters</b>	
Number of cores (max=96)	2
QC	<input checked="" type="checkbox"/>
Verbose	<input checked="" type="checkbox"/>
<b>Basic parameters</b>	
Number of sensor bands	3
Control band index	-1
<b>Sensor band names</b>	
0	igm
1	igg
2	ctl
<b>Band expected relative location</b>	
0	0.25

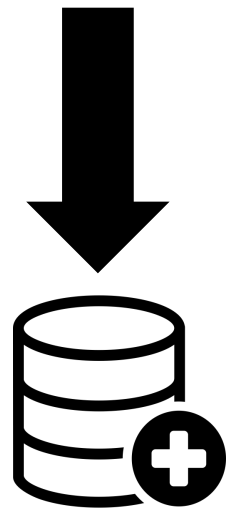
Log window output:

```
2021-01-05 15:14:30 - ui.mainwindow - Extracting POCT from image finished successfully.
2021-01-05 15:16:15 - ui.mainwindow - Drawing sensor 0
2021-01-05 15:16:17 - ui.mainwindow - Drawing sensor 1
2021-01-05 15:16:18 - ui.mainwindow - Drawing sensor 2
```

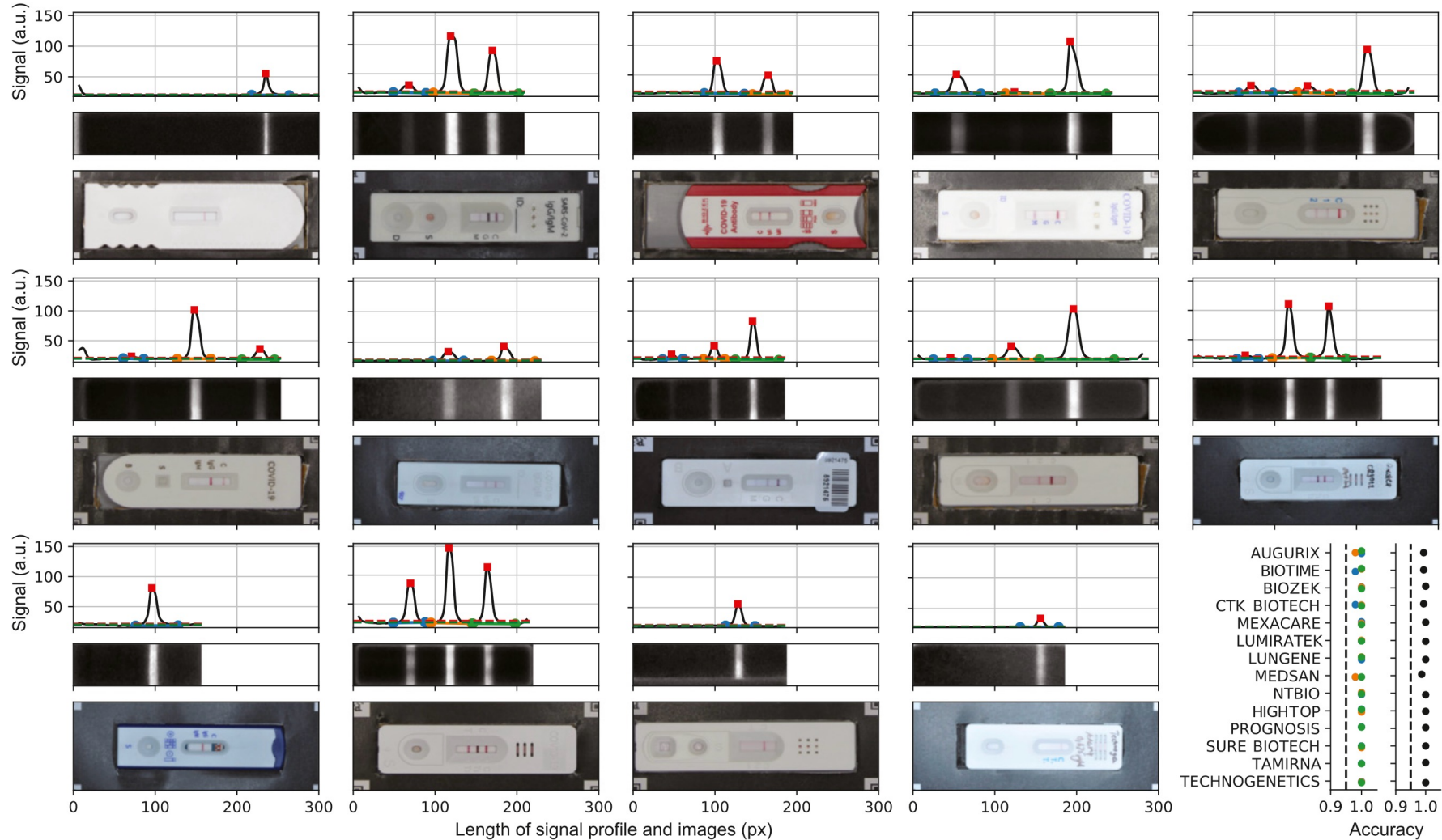
# Expert-driven projects :: pyPOCQuant



Analysis results  
 Patient metadata  
 Experiment metadata



# Expert-driven projects :: pyPOCQuant



And more!

# Expert-driven projects :: pyPOCQuant

Cuny A. P., Rudolf F., Ponti A. SoftwareX 15:2021, 100710. DOI: 10.1016/j.softx.2021.100710

Work by **Andreas Cuny**

## Expert contributions

**Fabian Rudolf:** development of the testing methodology, field work, and project management

**Andreas Cuny:** development of hardware components of pyPOCQuant, application development

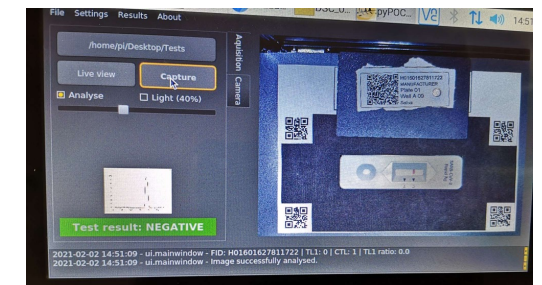
**Aaron Ponti:** computer vision algorithm development, application development

## Users

- Baselland Test Center (via Biolytix AG)
- Swiss Tropical and Health Institute (TPH), Basel
- Fachhochschule Nordwestschweiz (FHNW), Muttenz
- Swiss Armed Forces ← **Large-scale testing before Rekrutenschule Summer 2020**
- Canton Grisons and Swiss Federal Office of Public Health (Kantonaler Führungsstab Graubünden)
- Purdue University, Indiana, USA
- Test centers in Argentina and Greece



New hardware prototype (with touch screen and Raspberry Pi 4 Model B) for the Swiss Armed Forces



Adapted user interface (and template) for the new hardware

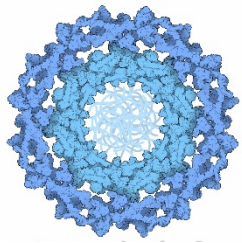
# Expert-driven projects :: pyMINFLUX



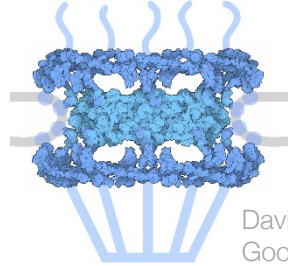
Expert: [Javier Casares Arias](#), Single Cell Facility, D-BSSE ETHZ

# Expert-driven projects :: pyMINFLUX

QC sample: Nuclear pore complex imaging



Top view



Side view

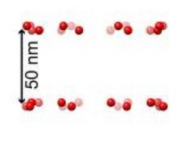
David  
Goodsell

c EM density

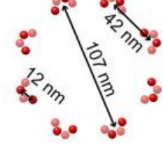


Red  
C-termini of  
Nup96

d side view



e top view

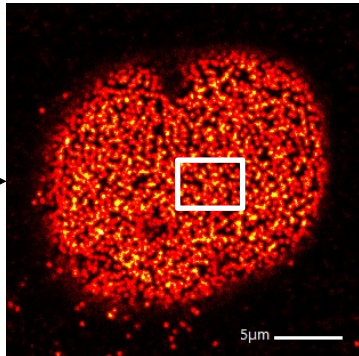


NPCs have  
8-fold  
symmetry

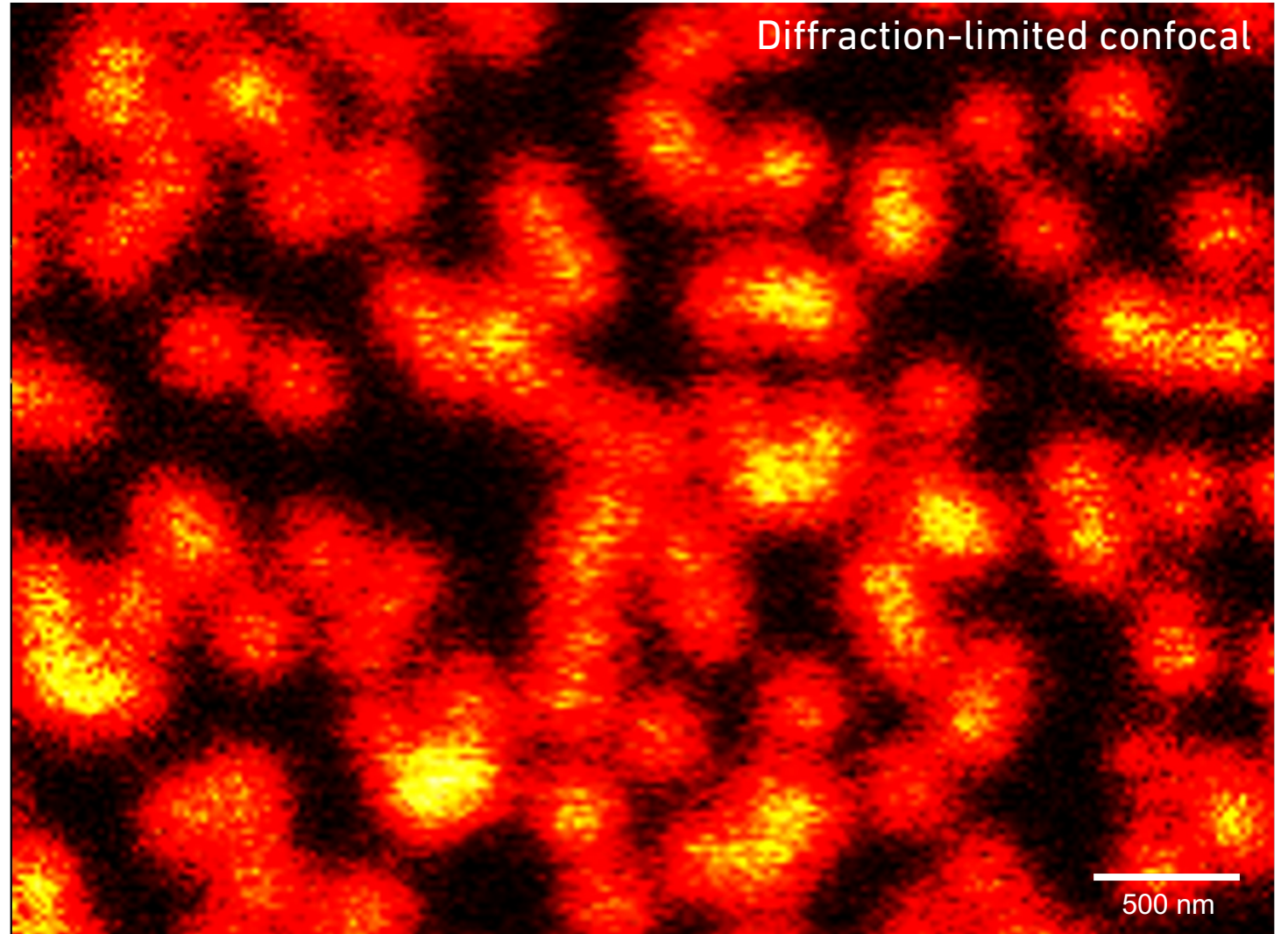
doi: [10.1038/s41592-019-0574-9](https://doi.org/10.1038/s41592-019-0574-9)

U2OS/Nup96-SNAP + BG-AF647

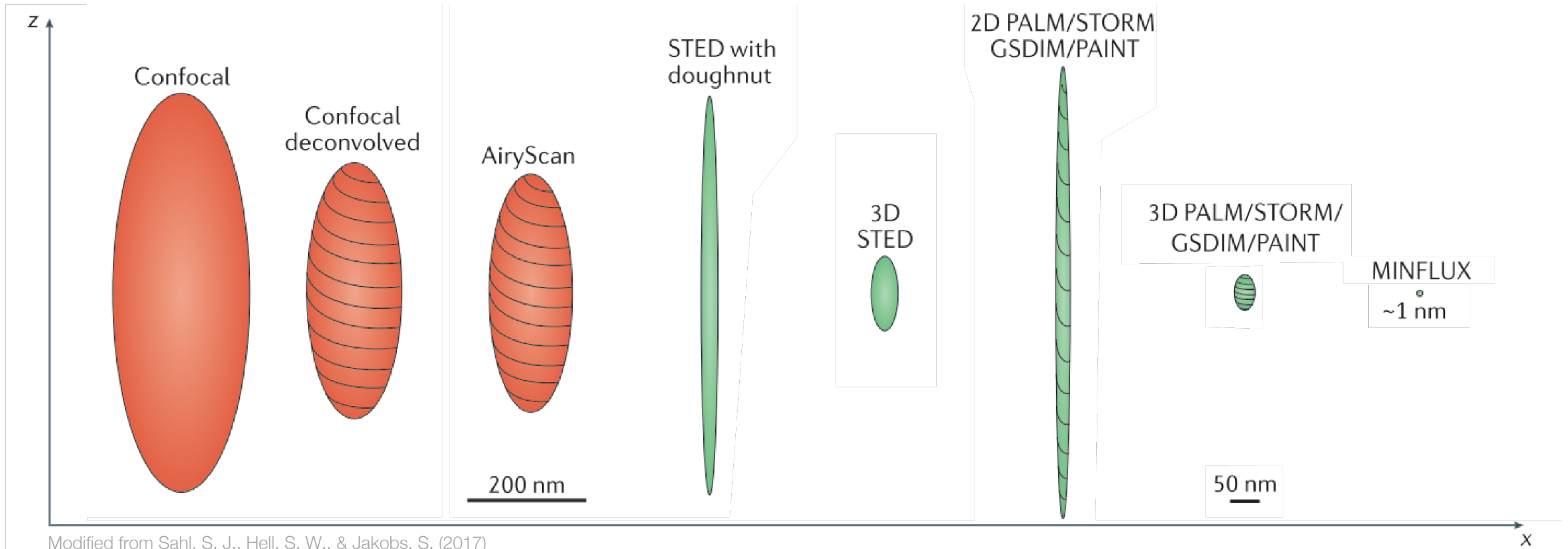
Cell  
nucleus



Diffraction-limited confocal



# Expert-driven projects :: pyMINFLUX

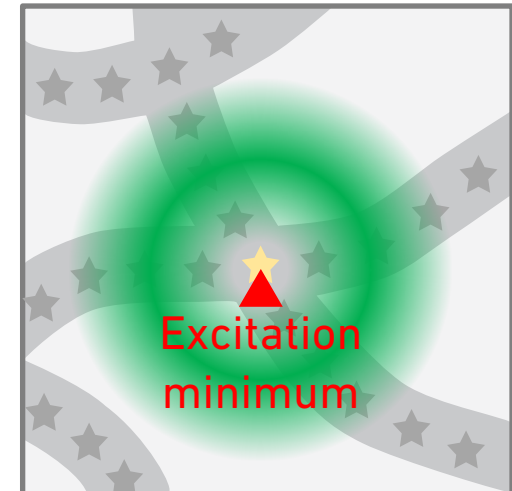


XY	200 nm	150 nm	120 nm	30 nm	50 nm	20 nm	20 nm	1 nm
Z	600 nm	400 nm	350 nm	600 nm	150 nm	600 nm	100 nm	1 nm

# Expert-driven projects :: pyMINFLUX

When the minimum at the center of the **excitation donut** and the fluorophore overlap there is no excitation, and thus no emission

Beam position = Fluorophore position

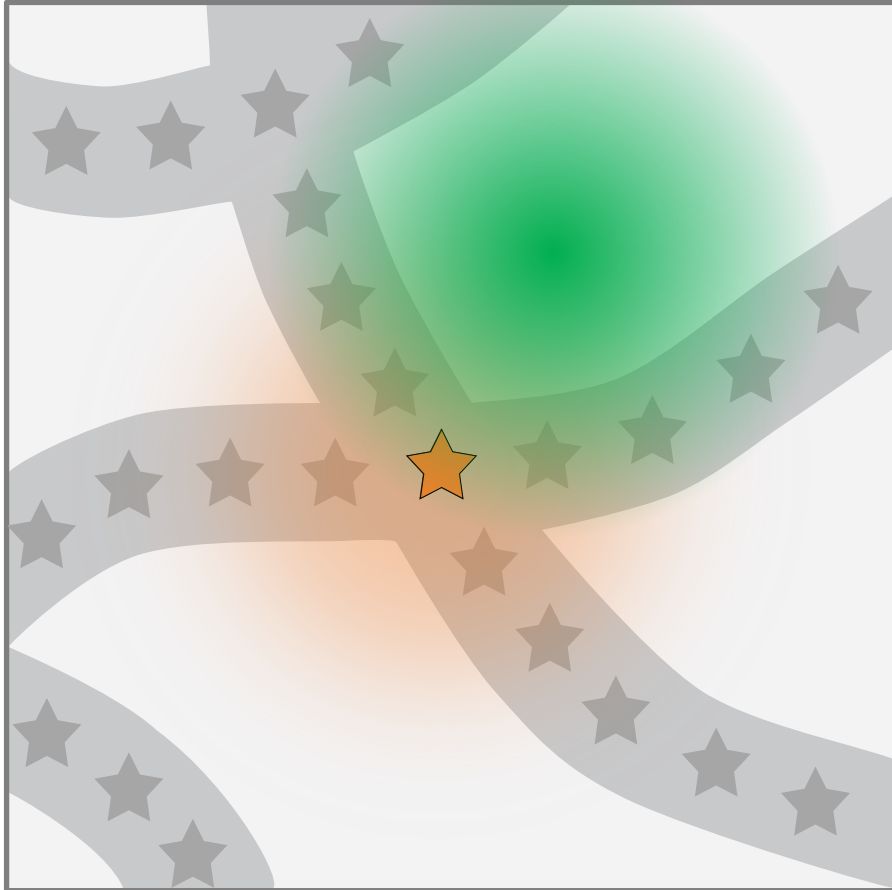




# Expert-driven projects :: pyMINFLUX

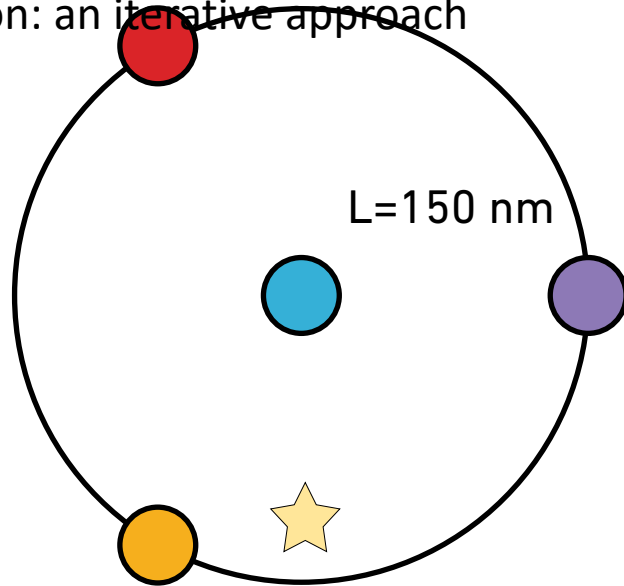
Localization: an iterative approach

Estimate fluorophore position with spot-shaped beam



# Expert-driven projects :: pyMINFLUX

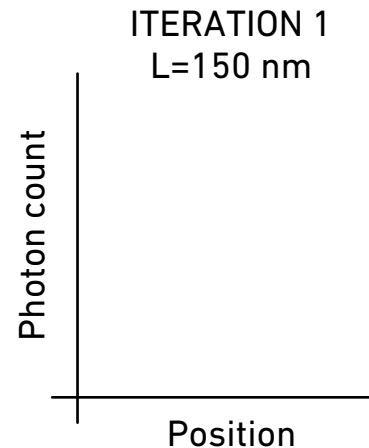
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

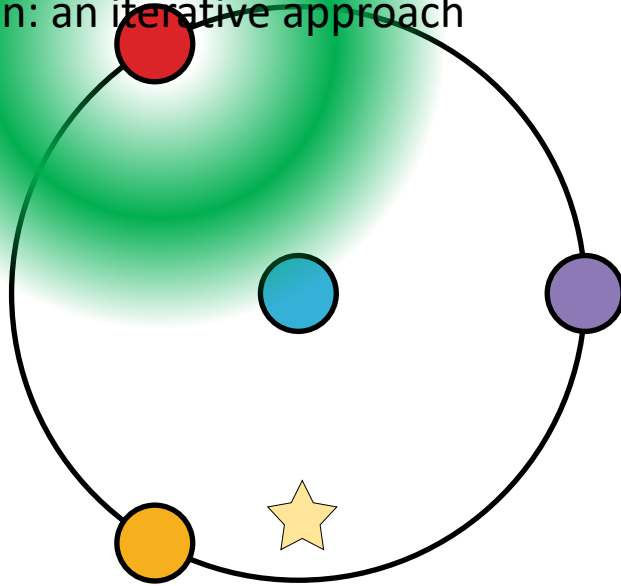
MINFLUX iteration:

1. Define scanning region according to previous estimation



# Expert-driven projects :: pyMINFLUX

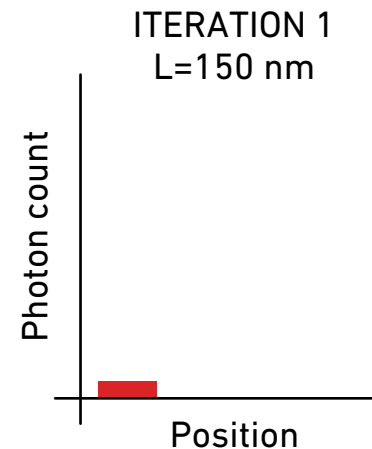
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

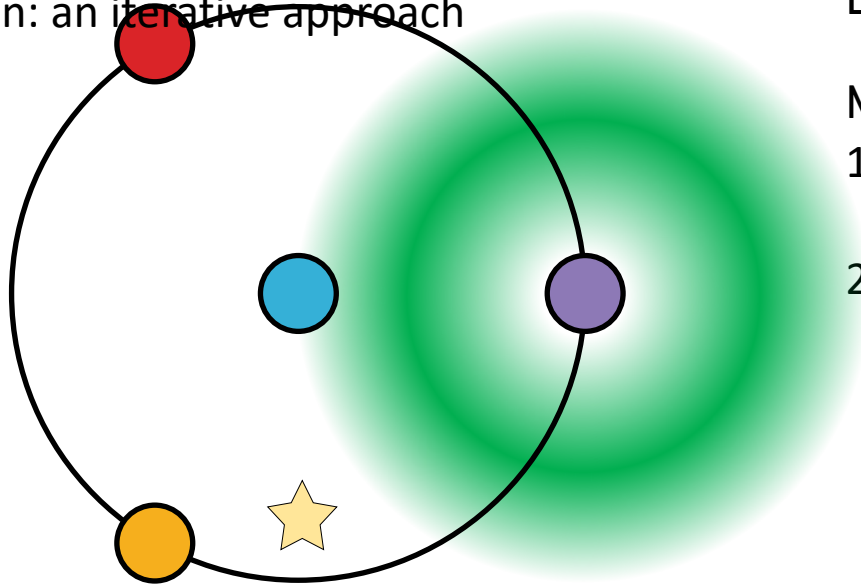
MINFLUX iteration:

1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern



# Expert-driven projects :: pyMINFLUX

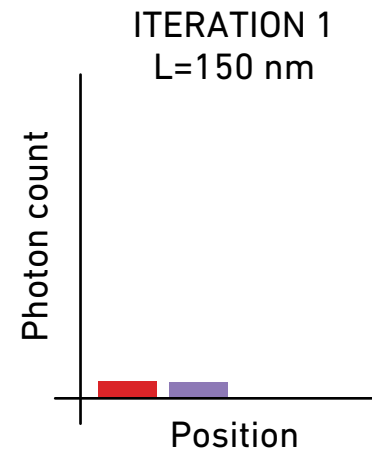
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

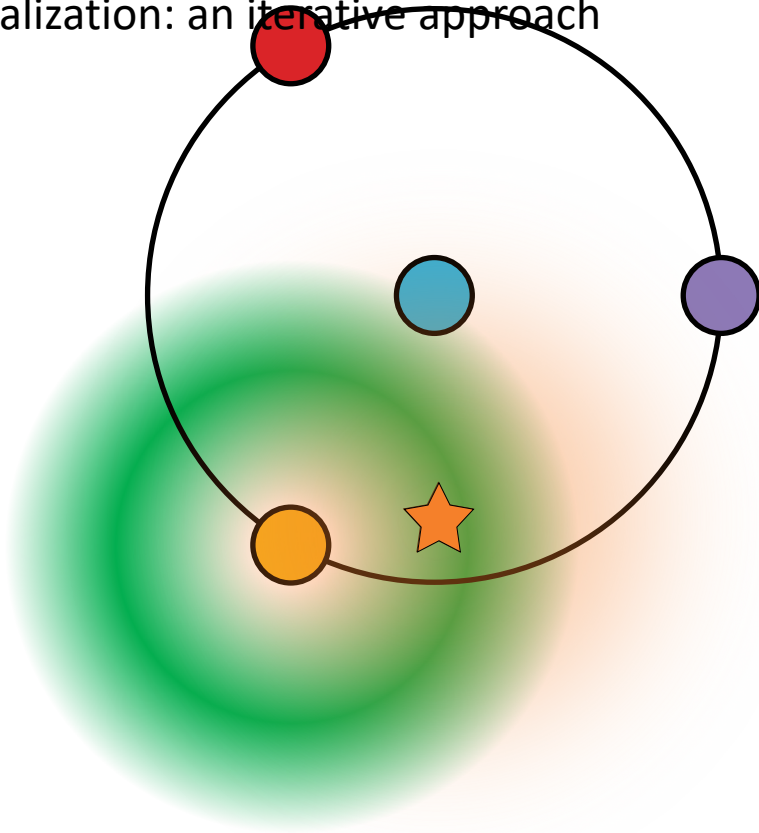
MINFLUX iteration:

1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern



# Expert-driven projects :: pyMINFLUX

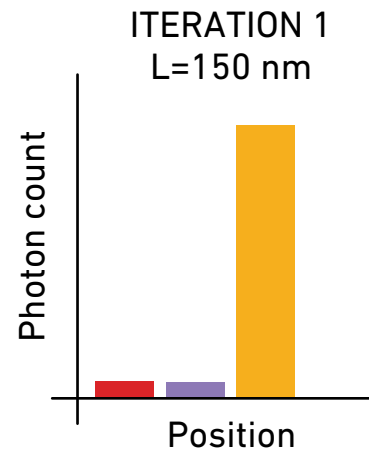
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

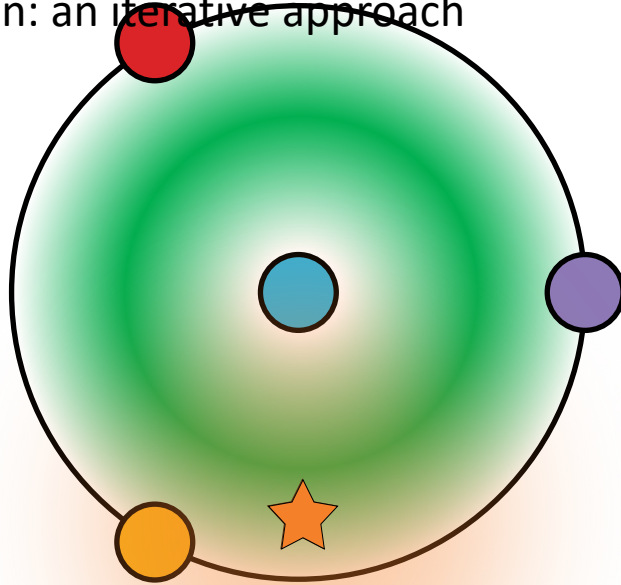
MINFLUX iteration:

1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern



# Expert-driven projects :: pyMINFLUX

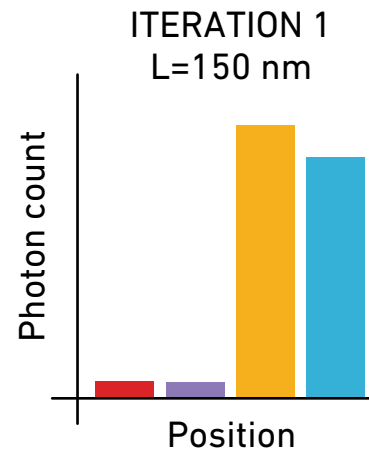
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

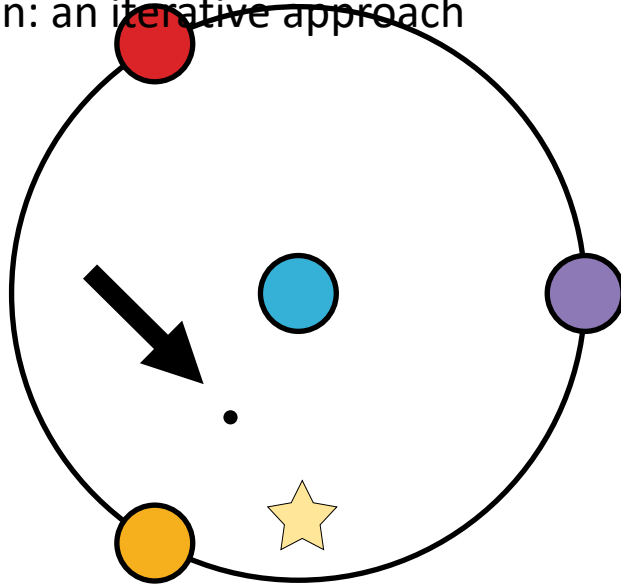
MINFLUX iteration:

1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern



# Expert-driven projects :: pyMINFLUX

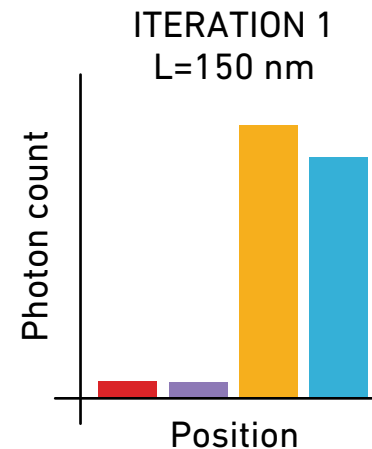
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

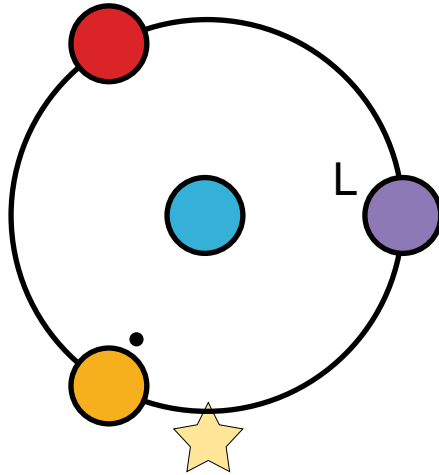
MINFLUX iteration:

1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern
3. Analyse photon counts and refine fluorophore position estimation



# Expert-driven projects :: pyMINFLUX

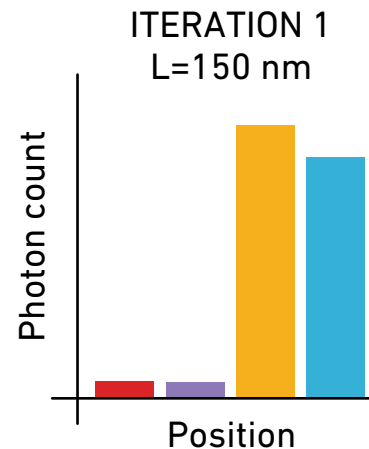
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

MINFLUX iteration:

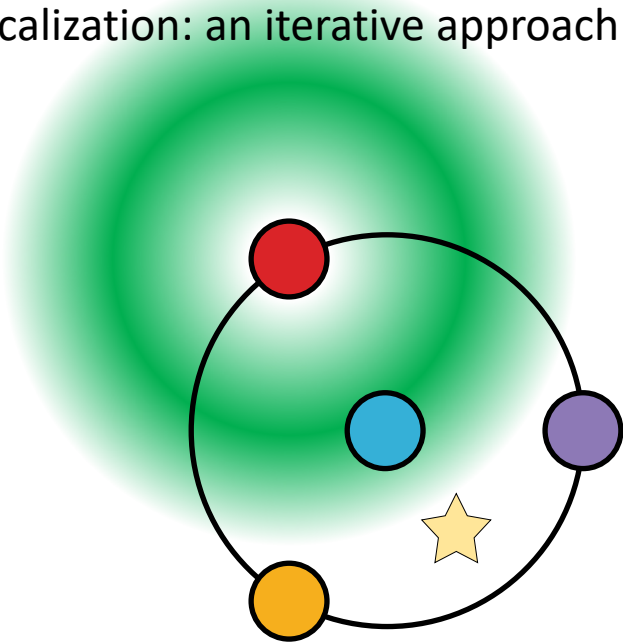
1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern
3. Analyse photon counts and refine fluorophore position estimation
4. Decrease "L"





# Expert-driven projects :: pyMINFLUX

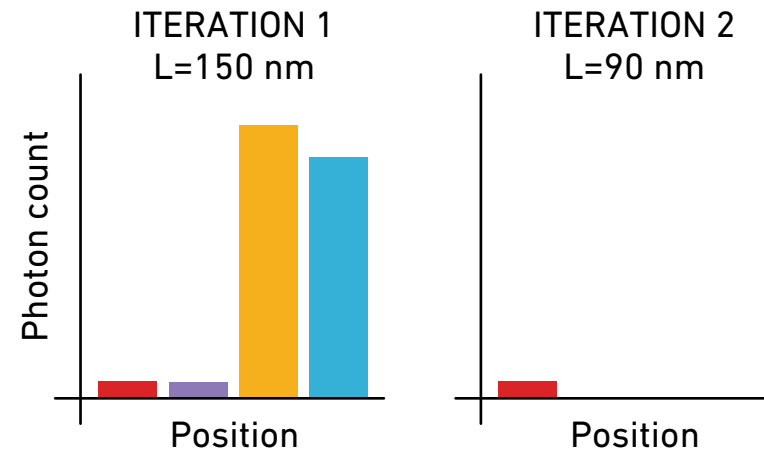
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

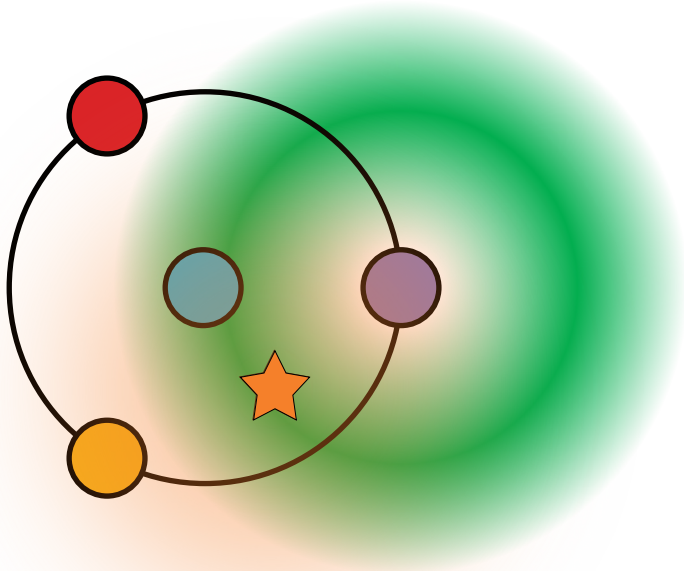
MINFLUX iteration:

1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern
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# Expert-driven projects :: pyMINFLUX

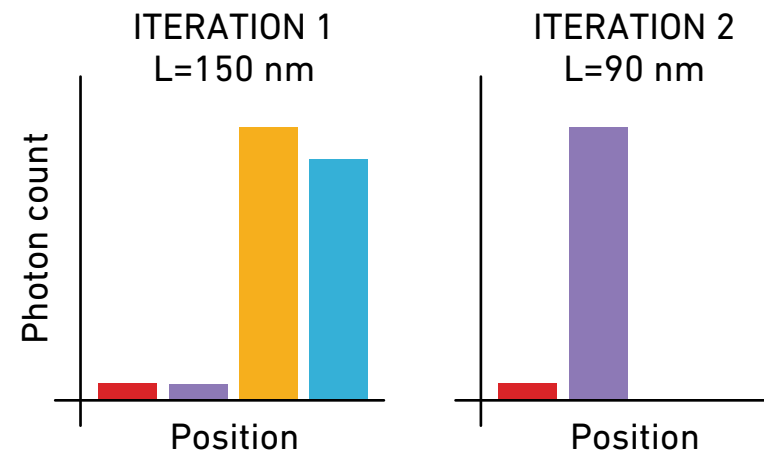
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

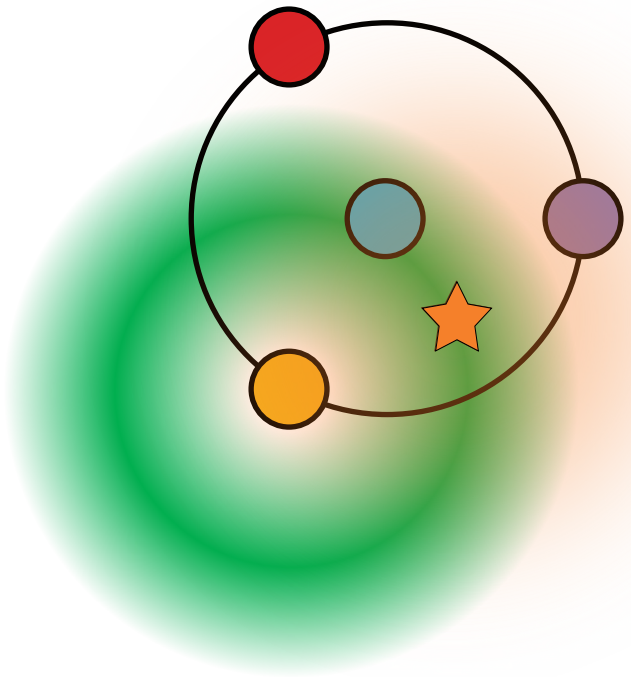
MINFLUX iteration:

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# Expert-driven projects :: pyMINFLUX

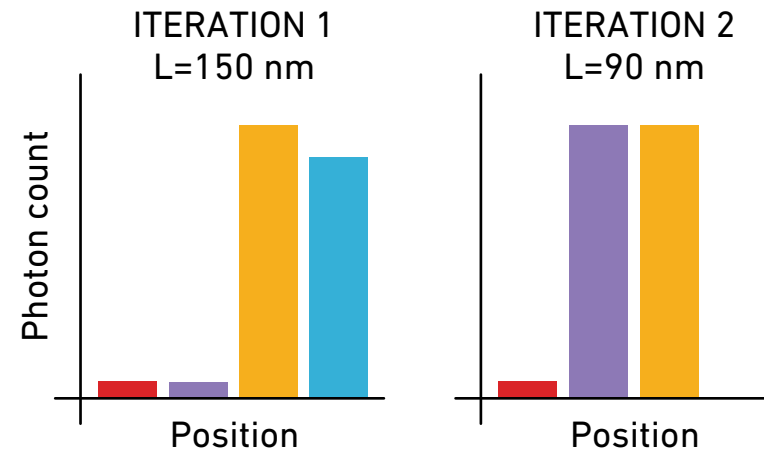
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

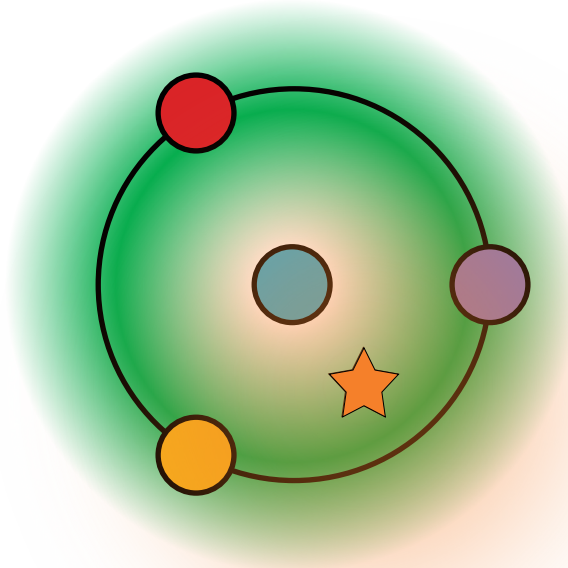
MINFLUX iteration:

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# Expert-driven projects :: pyMINFLUX

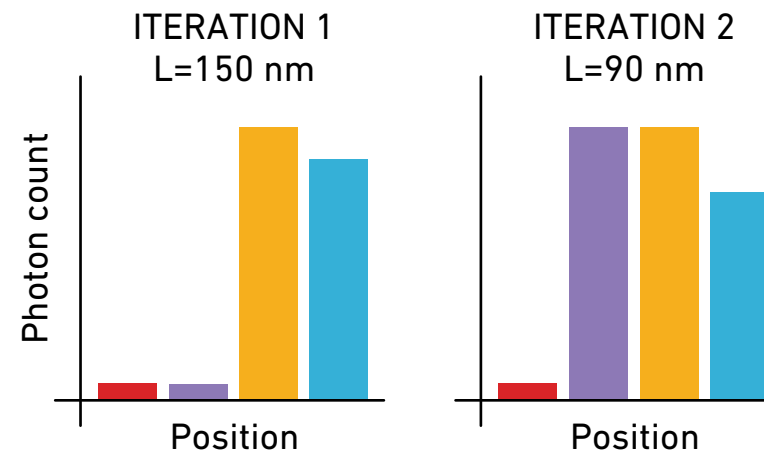
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

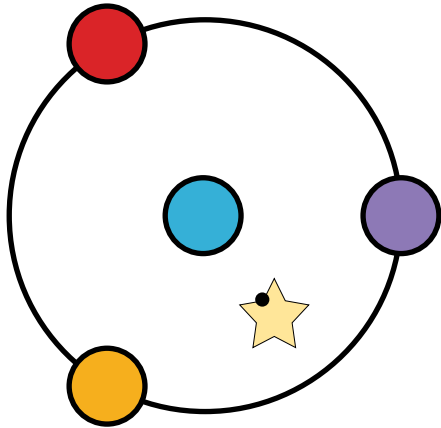
MINFLUX iteration:

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# Expert-driven projects :: pyMINFLUX

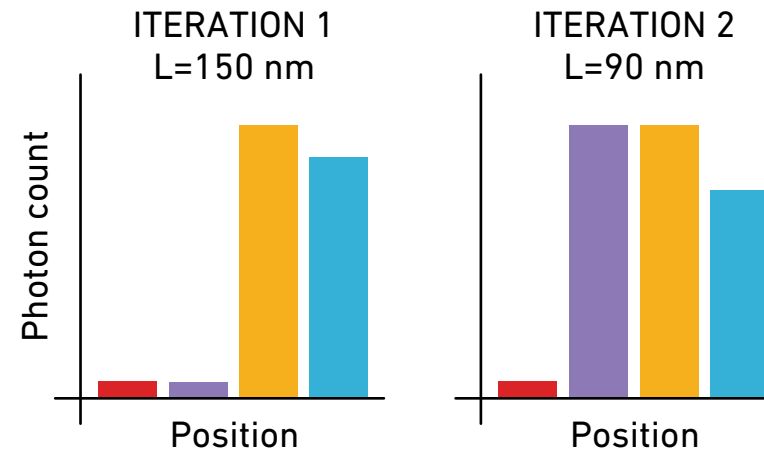
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

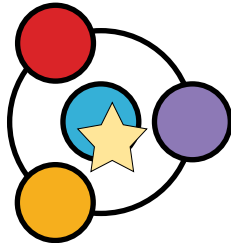
MINFLUX iteration:

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# Expert-driven projects :: pyMINFLUX

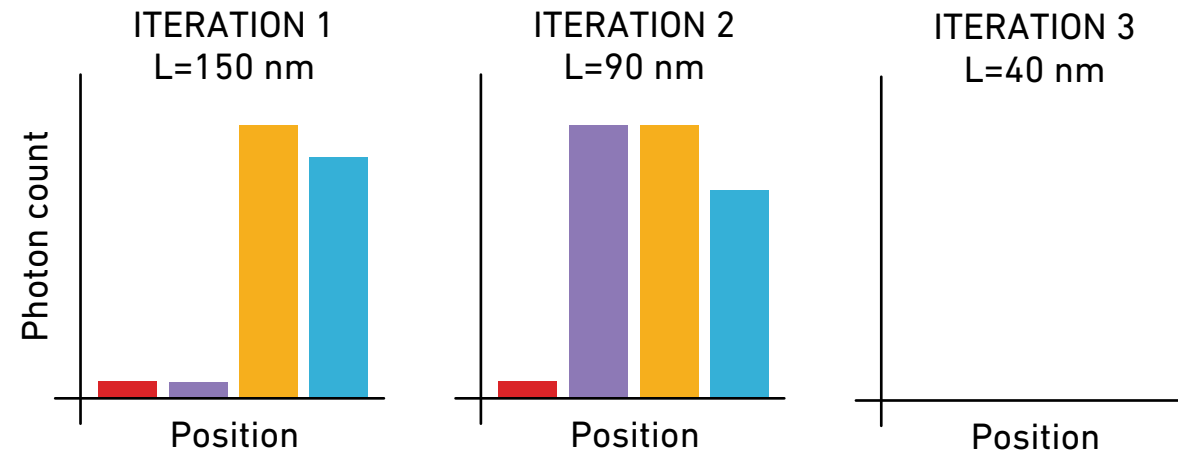
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

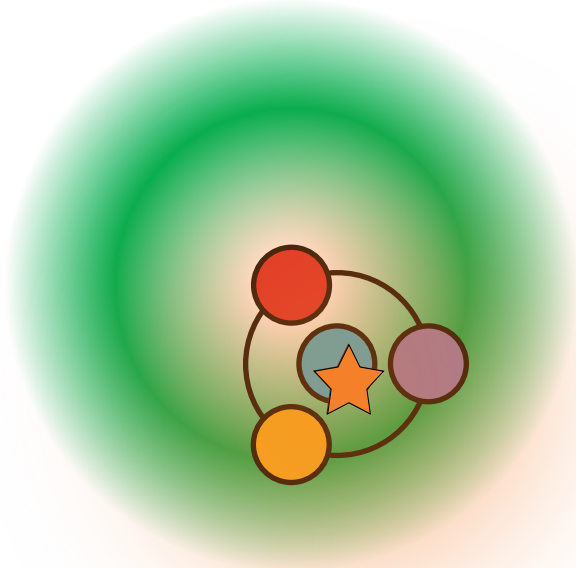
MINFLUX iteration:

1. Define scanning region according to previous estimation
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# Expert-driven projects :: pyMINFLUX

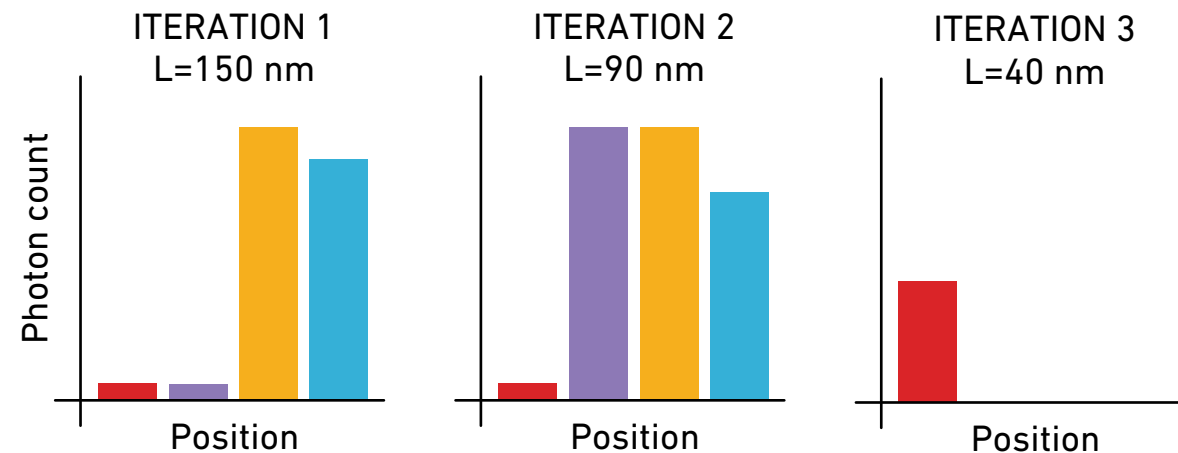
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

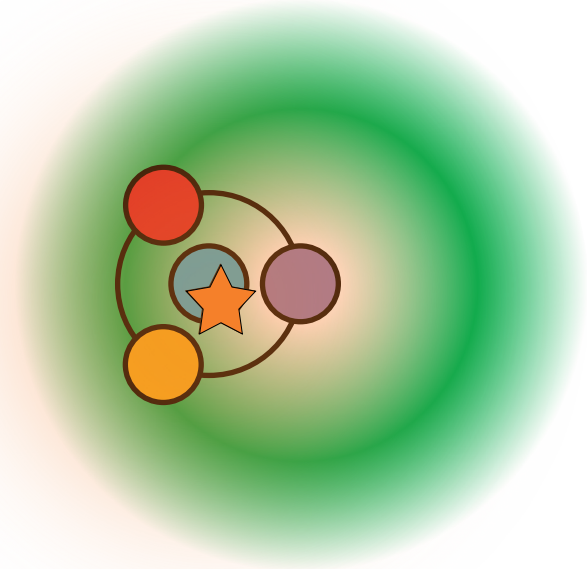
MINFLUX iteration:

1. Define scanning region according to previous estimation
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3. Analyse photon counts and refine fluorophore position estimation
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# Expert-driven projects :: pyMINFLUX

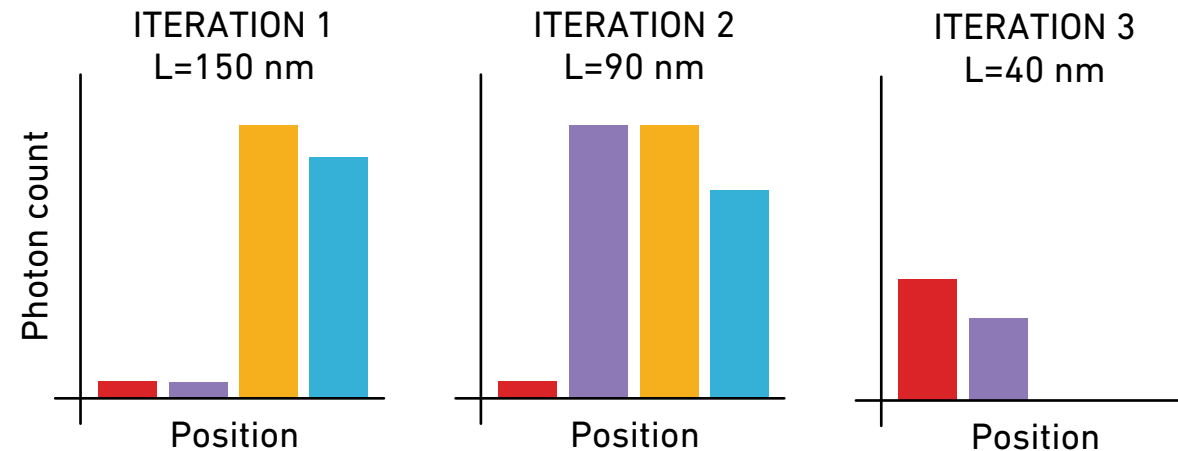
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

MINFLUX iteration:

1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern
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4. Decrease "L"





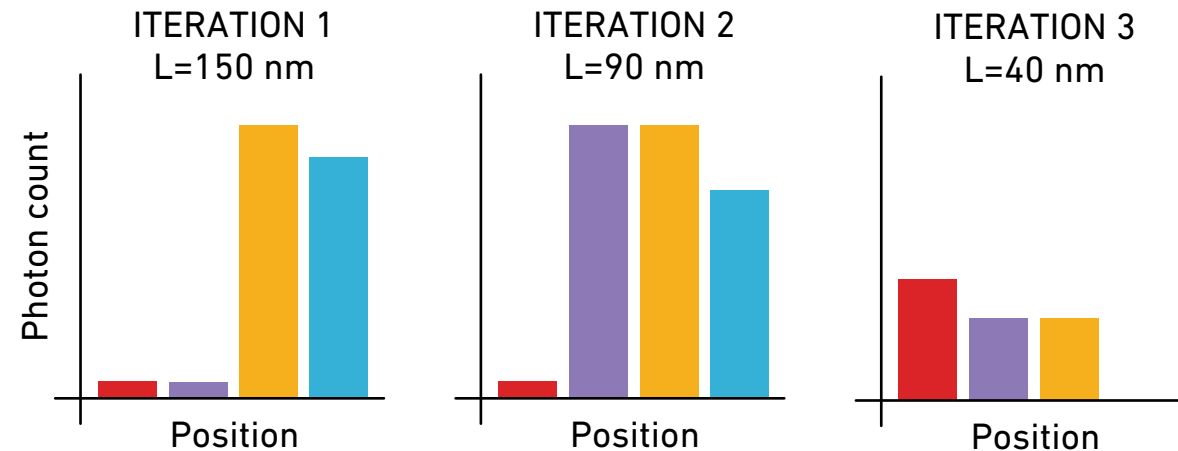
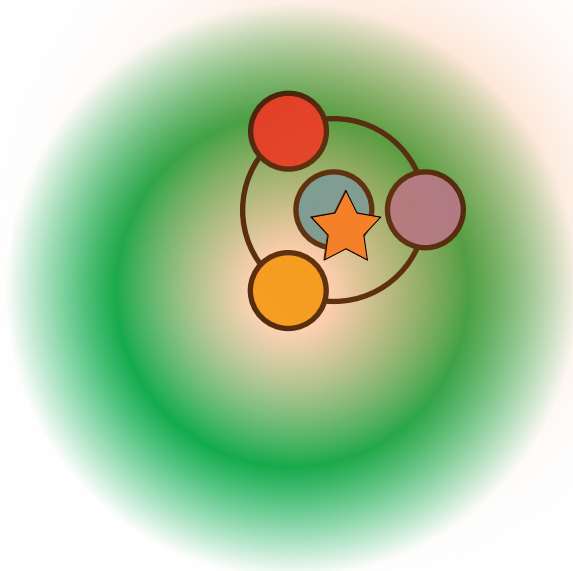
# Expert-driven projects :: pyMINFLUX

Localization: an iterative approach

Estimate fluorophore position with spot-shaped beam

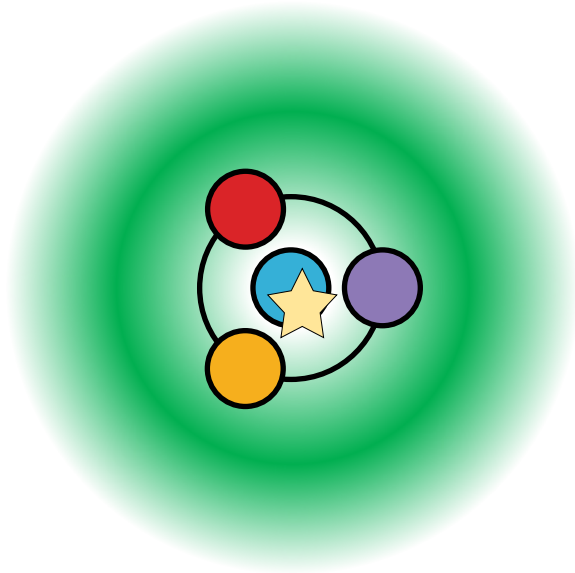
MINFLUX iteration:

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4. Decrease "L"



# Expert-driven projects :: pyMINFLUX

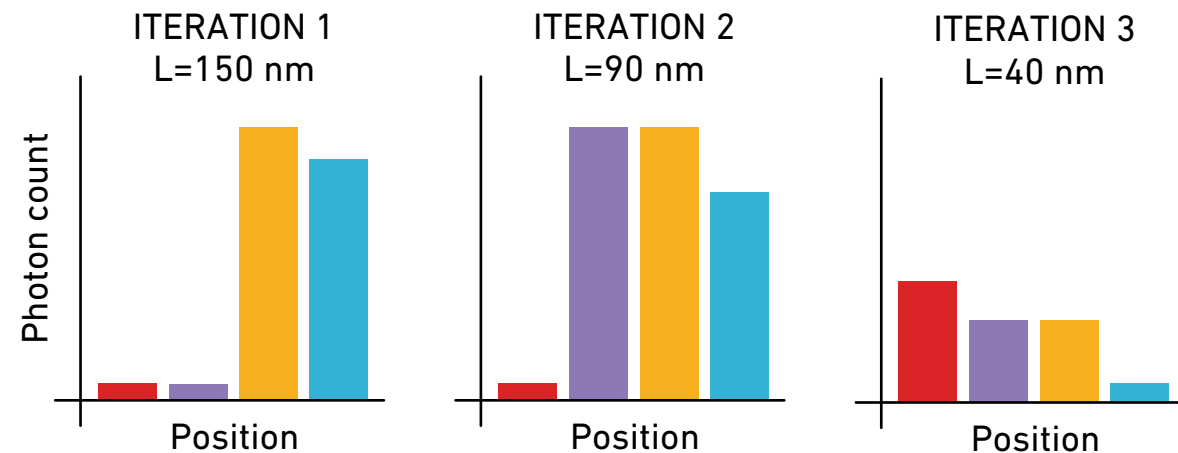
Localization: an iterative approach



Estimate fluorophore position with spot-shaped beam

MINFLUX iteration:

1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern
3. Analyse photon counts and refine fluorophore position estimation
4. Decrease "L"



# Expert-driven projects :: pyMINFLUX

Localization: an iterative approach

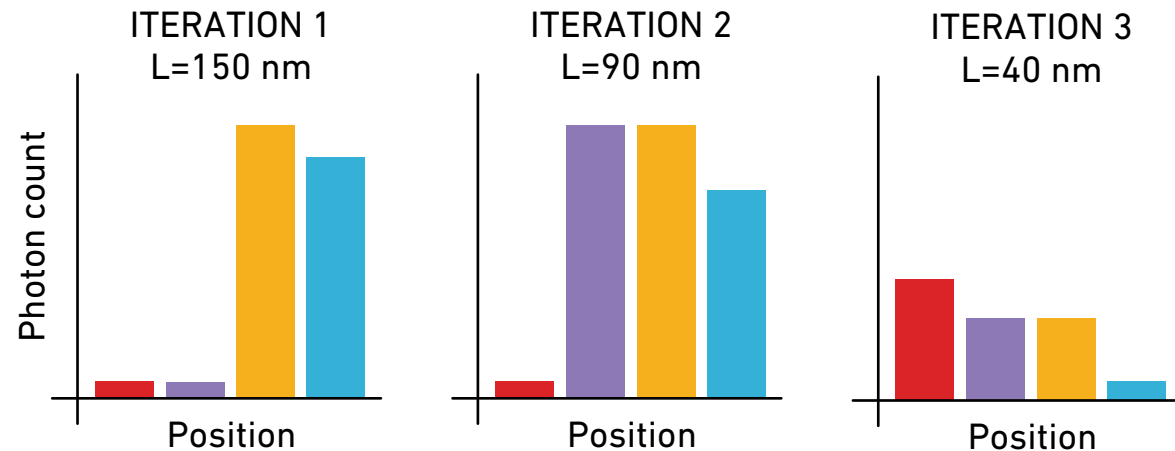
Estimate fluorophore position with spot-shaped beam

MINFLUX iteration:

1. Define scanning region according to previous estimation
2. Register emitted photons from every position in the pattern
3. Analyse photon counts and refine fluorophore position estimation
4. Decrease "L"



FLUOROPHORE  
LOCALIZATION



# Expert-driven projects :: pyMINFLUX

Operation modalities

Localization



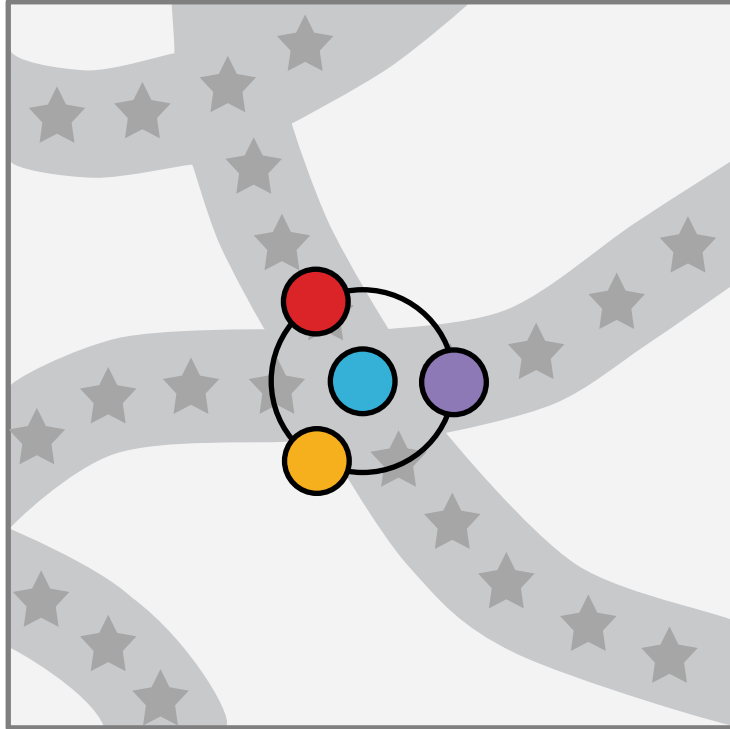
*Important: only very few fluorophores must be emitting at any given time!*

1. Fluorophores are localized 1 by 1

# Expert-driven projects :: pyMINFLUX

Operation modalities

Localization

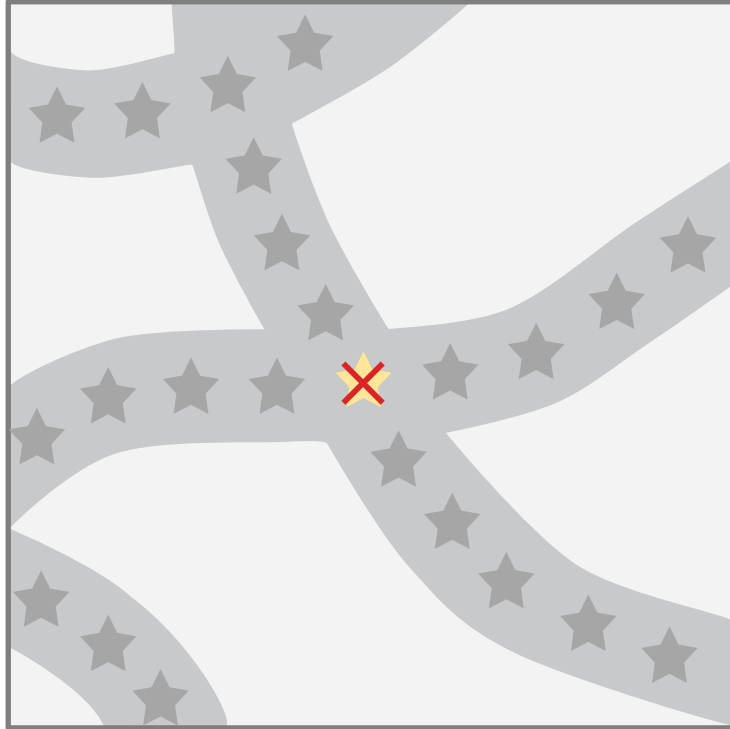


1. Fluorophores are localized 1 by 1

# Expert-driven projects :: pyMINFLUX

Operation modalities

Localization

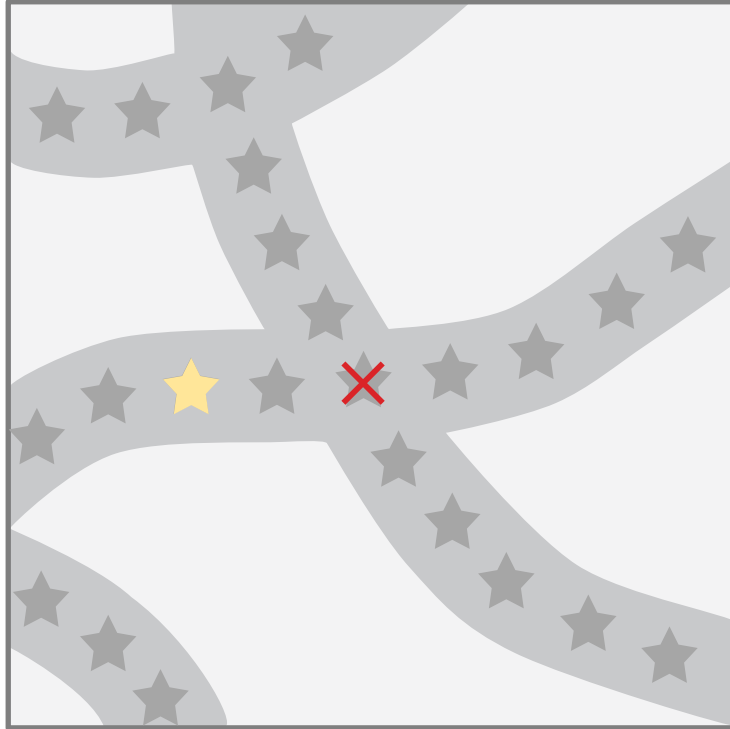


1. Fluorophores are localized 1 by 1

# Expert-driven projects :: pyMINFLUX

Operation modalities

Localization

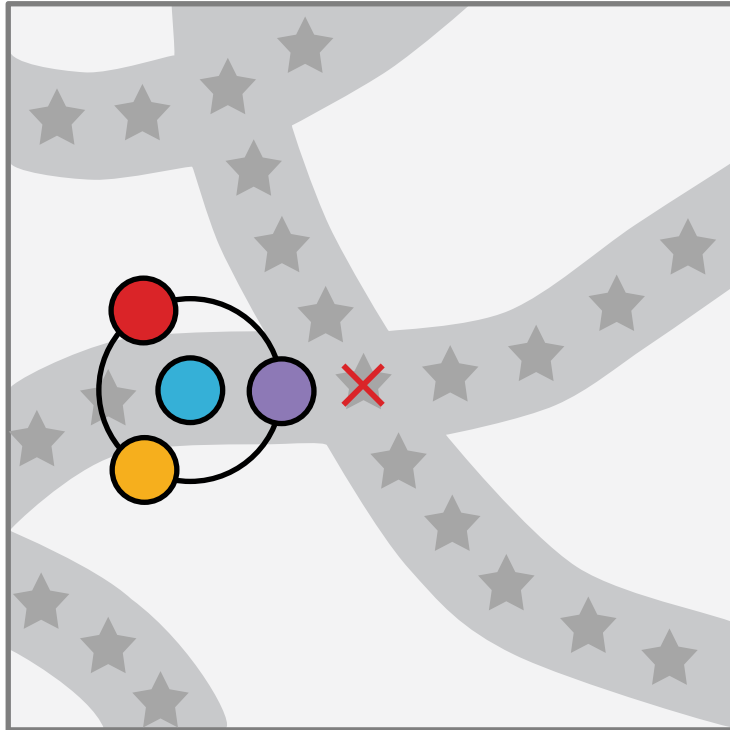


1. Fluorophores are localized 1 by 1

# Expert-driven projects :: pyMINFLUX

Operation modalities

Localization



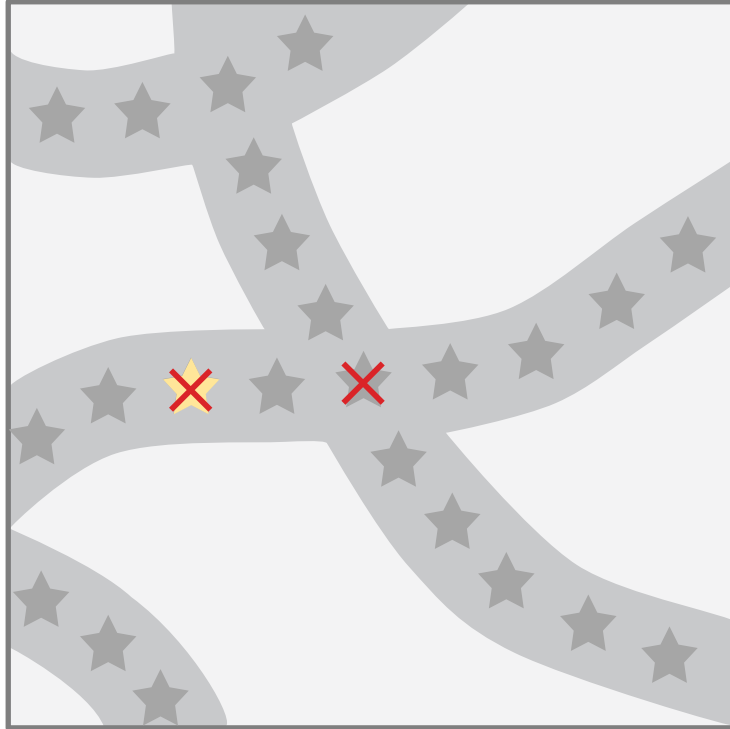
1. Fluorophores are localized 1 by 1



# Expert-driven projects :: pyMINFLUX

Operation modalities

Localization

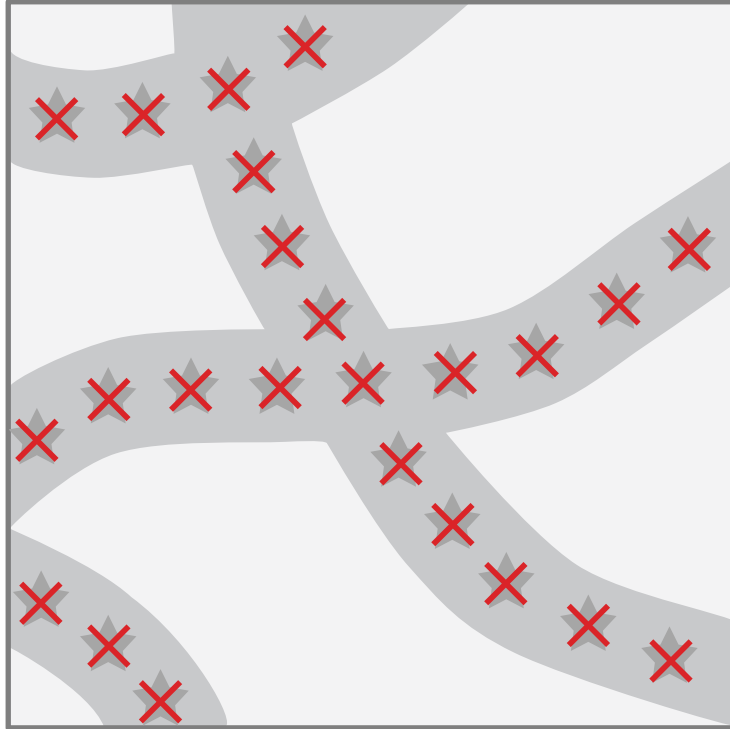


1. Fluorophores are localized 1 by 1

# Expert-driven projects :: pyMINFLUX

Operation modalities

Localization

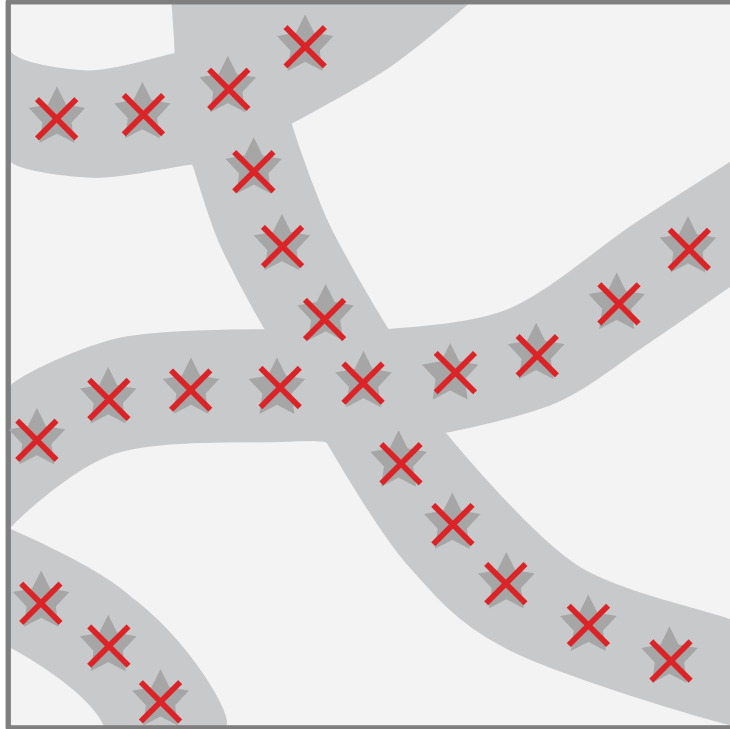


1. Fluorophores are localized 1 by 1

# Expert-driven projects :: pyMINFLUX

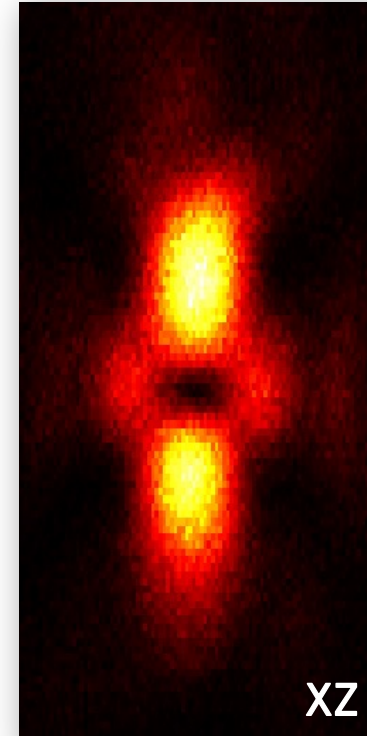
Operation modalities

Localization



3 nm - 2D

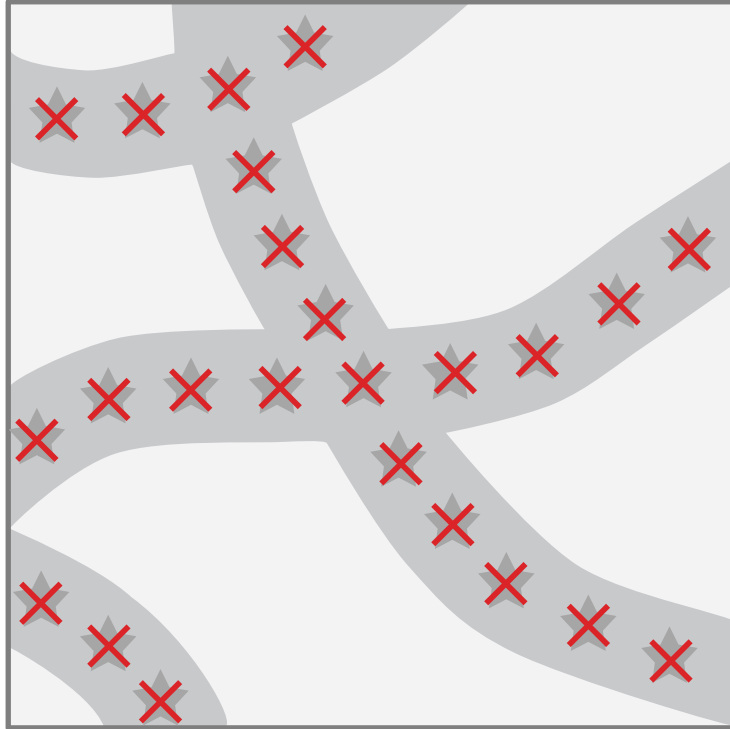
5 nm - 3D



# Expert-driven projects :: pyMINFLUX

Operation modalities

Localization



1. Fluorophores are localized 1 by 1
2. Coordinates turned into an image

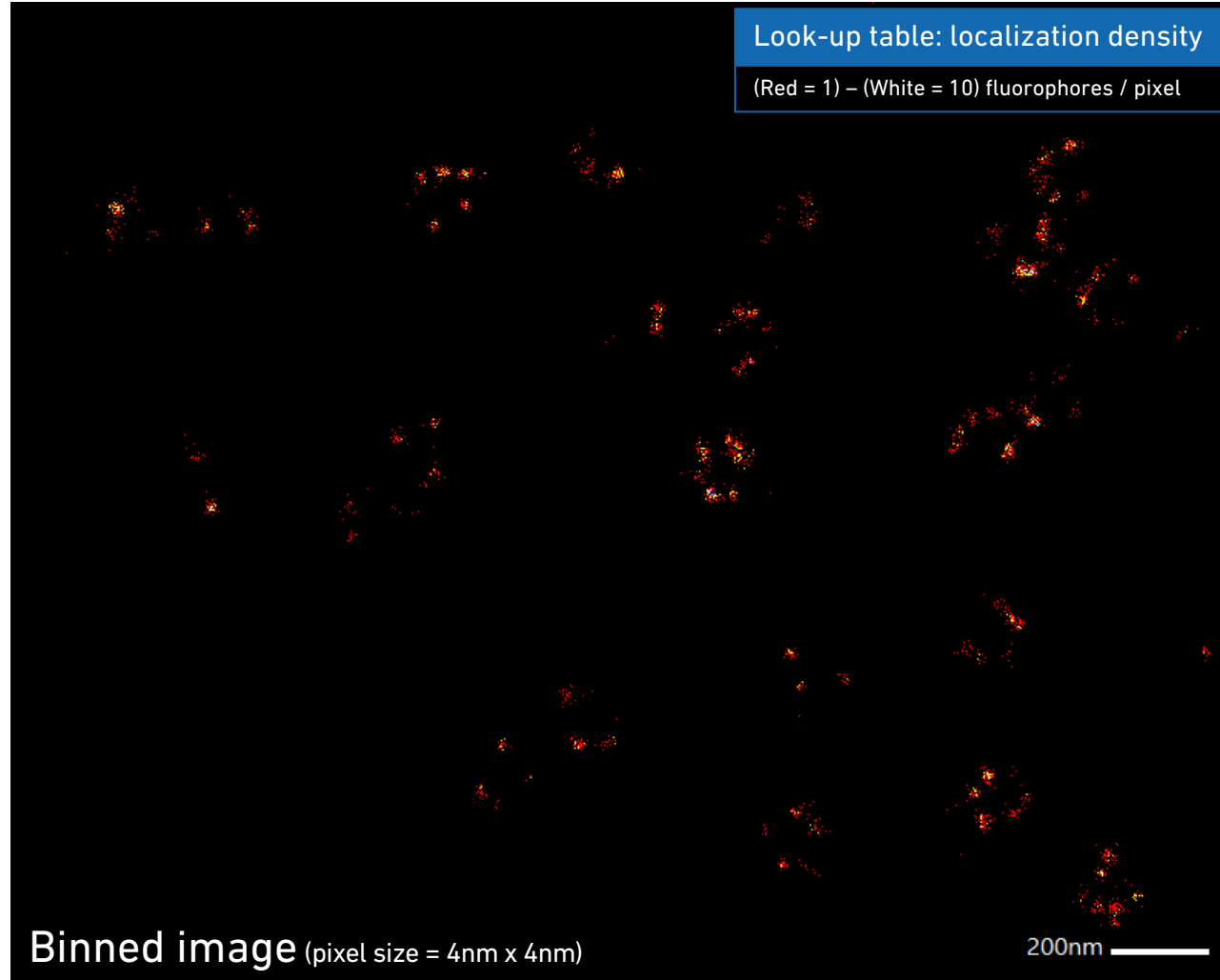
# Expert-driven projects :: pyMINFLUX

At this stage, we hadn't bought the microscope yet...

What we had:

- No localization data
- Low-res raster images

Testing phase



# Expert-driven projects :: pyMINFLUX

At this stage, we *still* hadn't  
bought the microscope

What we had:

- Undocumented .npz files

```
python
>>> import numpy as np
>>>
>>> data = np.load("2D_FA-IIF-AF467.npz")
>>> data
array([[[[0, 88459982, [ 9.20032180e-06, -2.84379534e-06, 0.00000000e+00], [ 9.20032180e-06, -2.84379534e-06, 0.00000000e+00], 191, 0, 47754.77547755,
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```

```
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  "itr",
  [
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    ("sta", "<i4""),
    ("cfr", "<f8""),
    ("dcr", "<f8""),
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]
```

timestamp?  
trace id?  
valid?

Testing phase



# Expert-driven projects :: pyMINFLUX

Now, we bought the microscope



Acquisition  
Data export  
Developed for STED  
Pixel-based  
MINFLUX data visualization  
requires "binning"  
Licensed  
Viewer version available  
Windows-only

+



3D rendering software  
Built-in Inspector bridge  
Tools + Plugins  
Open-source  
Cross-platform

## Missing features

Click-and-inspect  
Localization precision  
Fluorophore unmixing  
Filtering options  
Metrics comparison

Not facility-friendly

Fully open raw data format  
NumPy, MATLAB or JSON

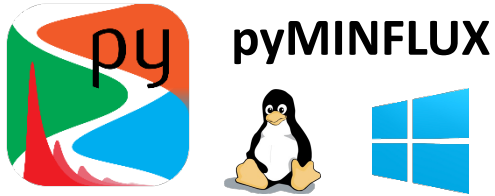
Documentation from Abberior



Production

# Expert-driven projects :: pyMINFLUX

<https://github.com/bsse-scf/pyMINFLUX>



Tailored for MINFLUX data

Open-access and cross-platform  
GUI-Independent API

Visualization, filtering and analysis tools

Selection tool

Data viewer

Customizable scatter plot

2-Color MINFLUX unmixing

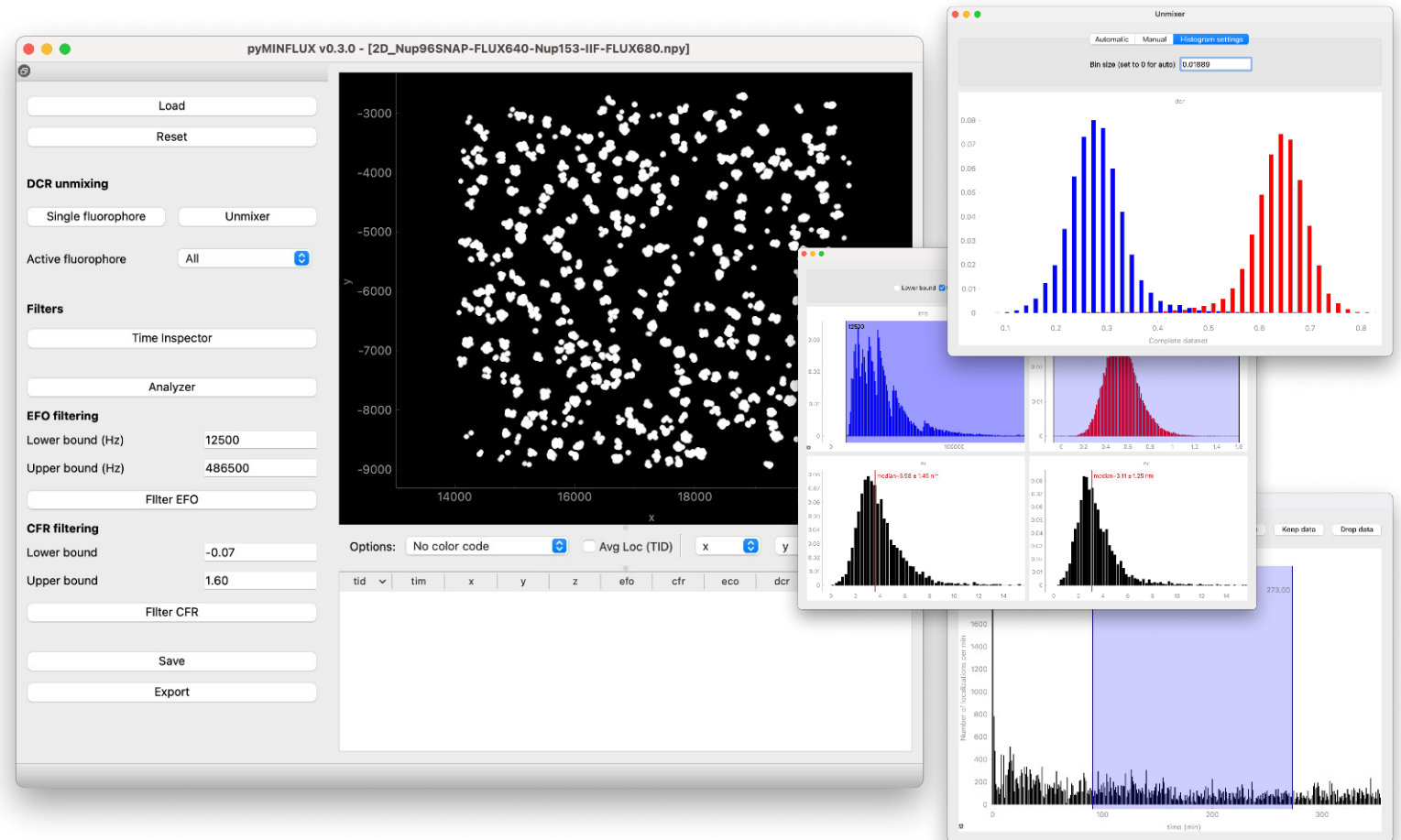
Localization precision calculation

Manual and automatic filtering

Fourier-ring correlation analysis

Integration with Paraview

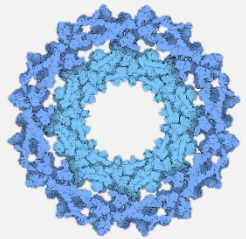
Custom .pmx format (metadata, processing parameters, ...)



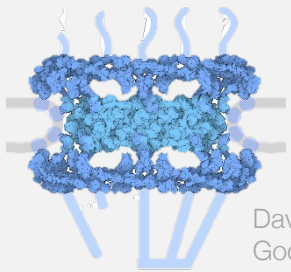


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view

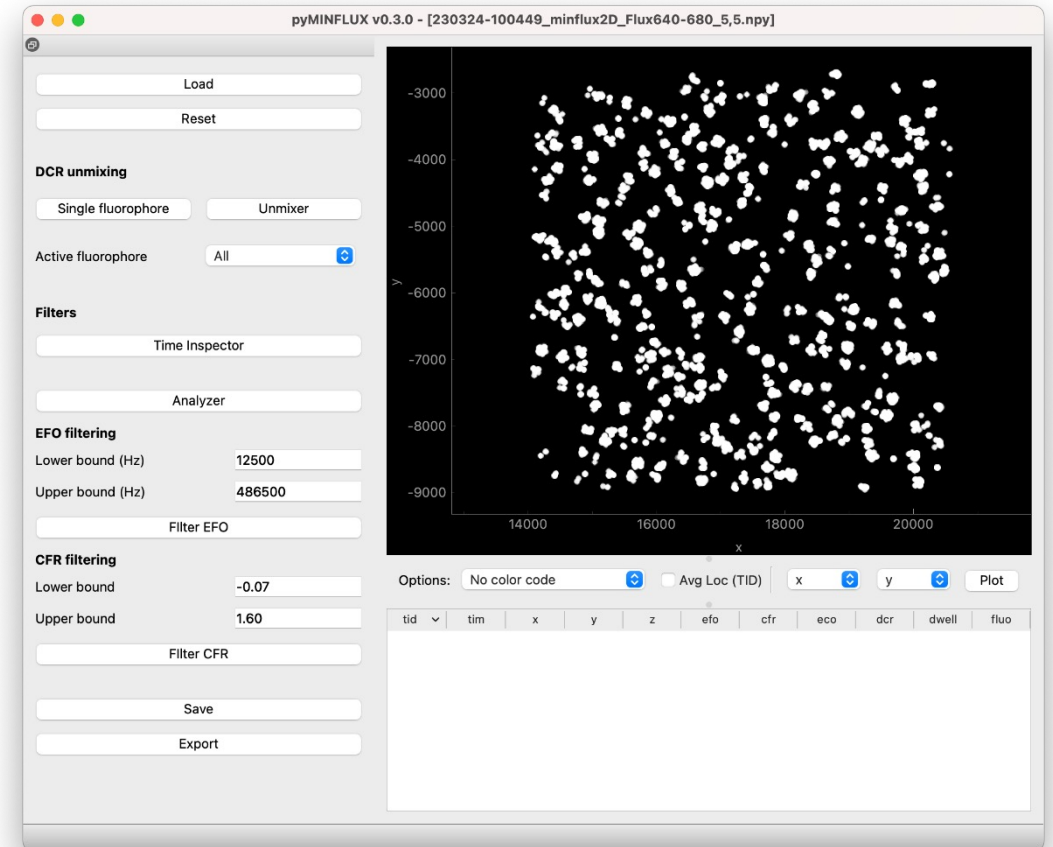
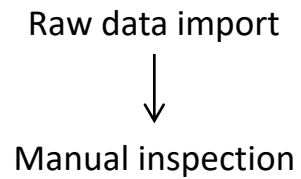


Side view

David  
Goodsell

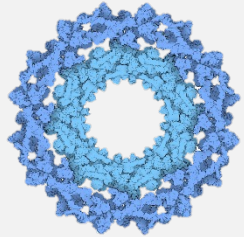
U2OS Nup96-SNAP cells  
BG-Abberior Flux 640  
Ms-Nup153 + 2<sup>ary</sup> Flux 680  
GLOX-MEA (10-15 mM)

pyMINFLUX workflow

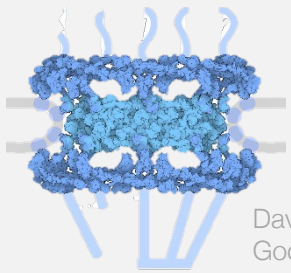


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view



Side view

David  
Goodsell

U2OS Nup96-SNAP cells  
BG-Abberior Flux 640  
Ms-Nup153 + 2<sup>ary</sup> Flux 680  
GLOX-MEA (10-15 mM)

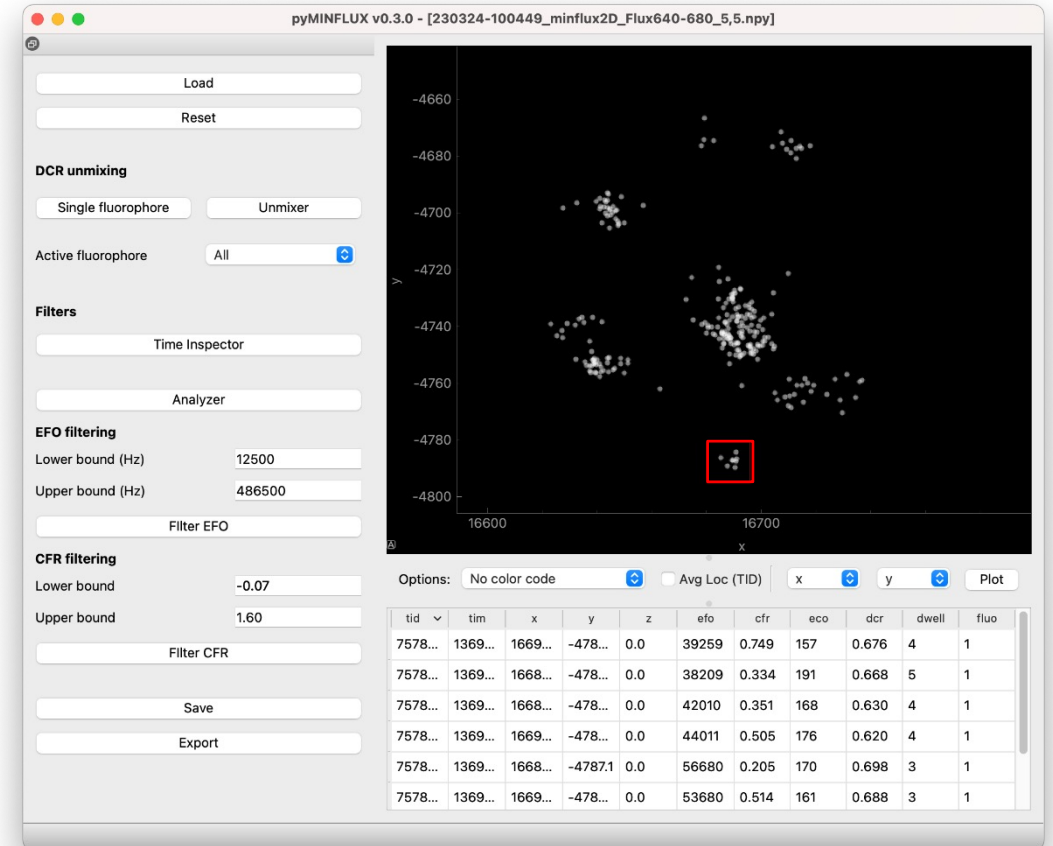
pyMINFLUX workflow

Raw data import



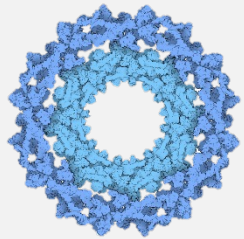
Manual inspection

Dataset navigation

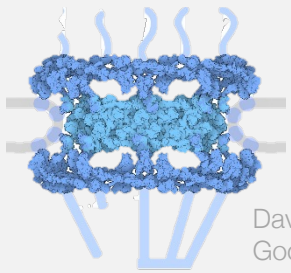


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view



Side view

David  
Goodsell

U2OS Nup96-SNAP cells  
BG-Abberior Flux 640  
Ms-Nup153 + 2<sup>ary</sup> Flux 680  
GLOX-MEA (10-15 mM)

pyMINFLUX workflow

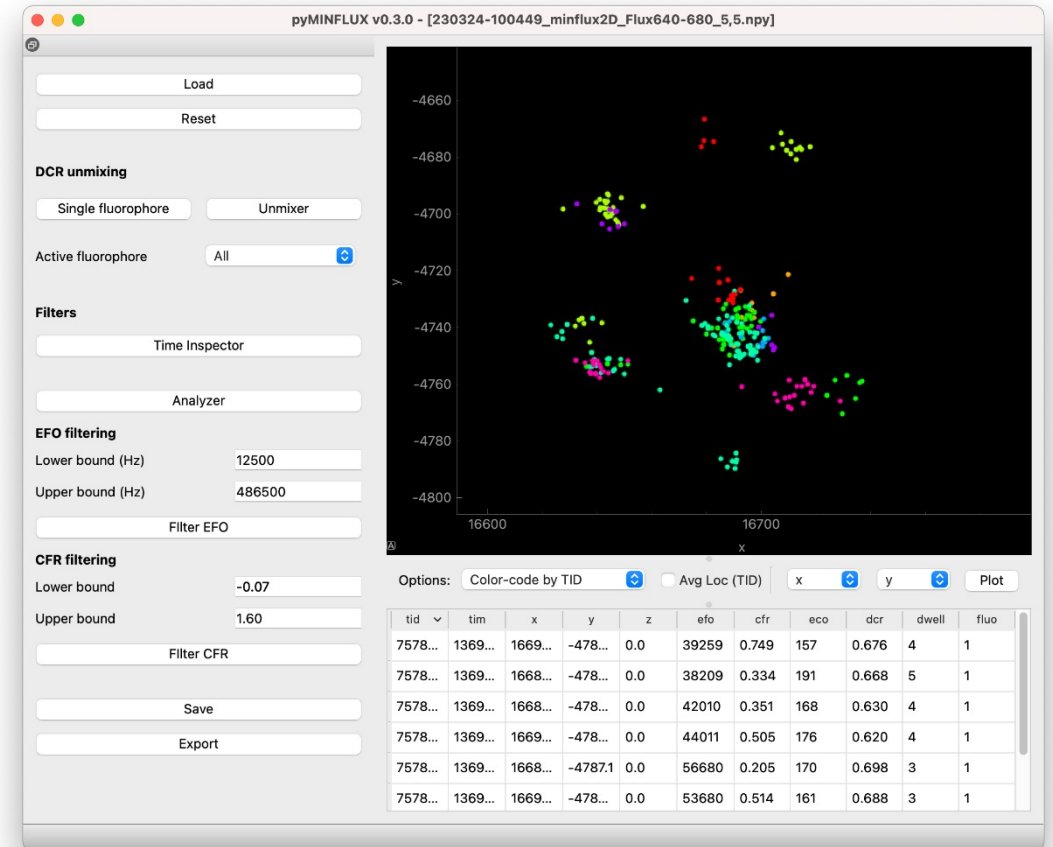
Raw data import



Manual inspection

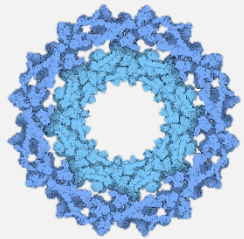
Dataset navigation

Trace coloring

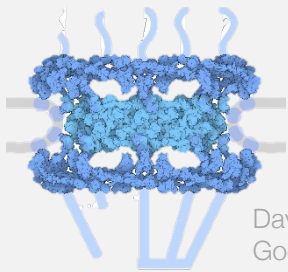


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view

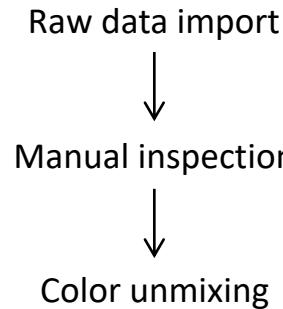


Side view

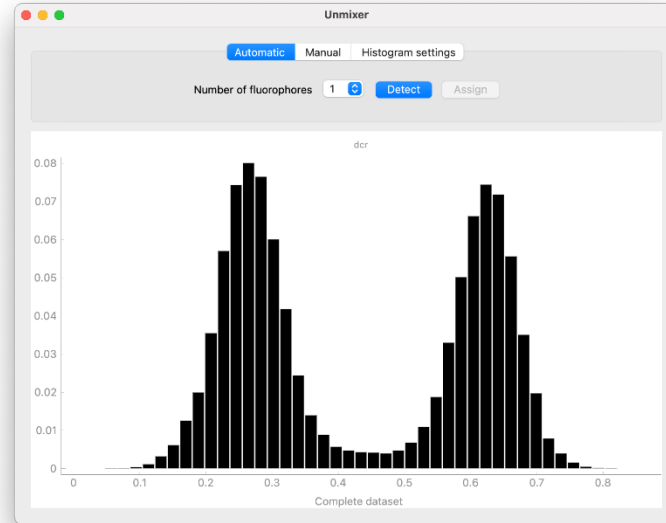
David Goodsell

U2OS Nup96-SNAP cells  
BG-Abberior Flux 640  
Ms-Nup153 + 2<sup>ary</sup> Flux 680  
GLOX-MEA (10-15 mM)

pyMINFLUX workflow



$$dcr = \frac{\text{Photons}_{\text{Detector1}}}{\text{Total}}$$



**EFO filtering**

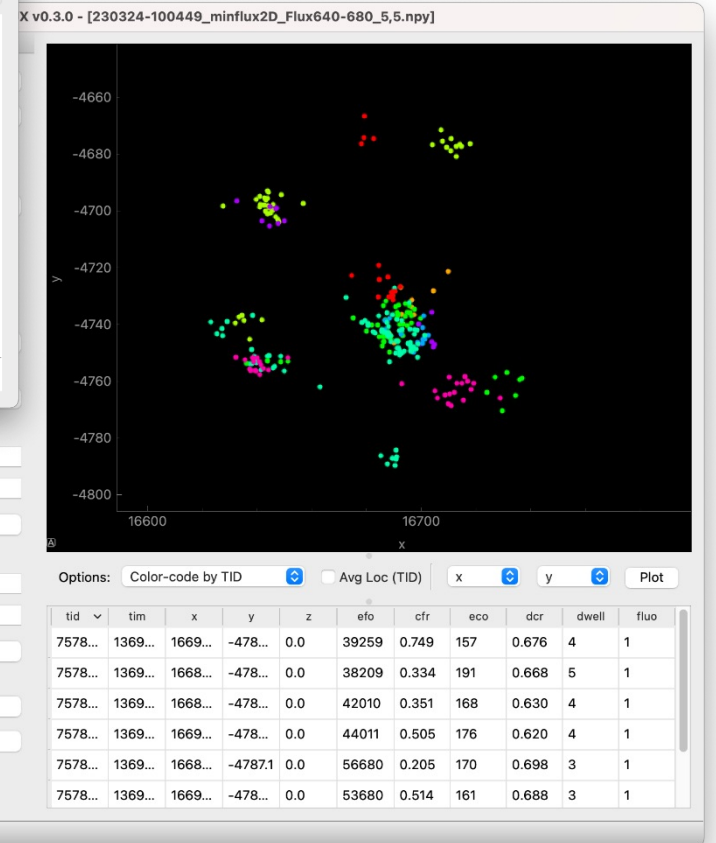
Lower bound (Hz)

Upper bound (Hz)

**CFR filtering**

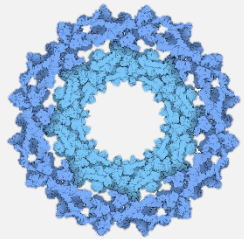
Lower bound

Upper bound

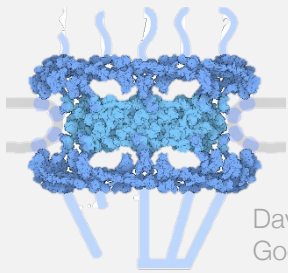


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view

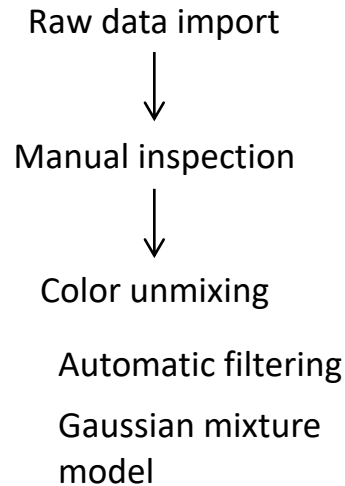


Side view

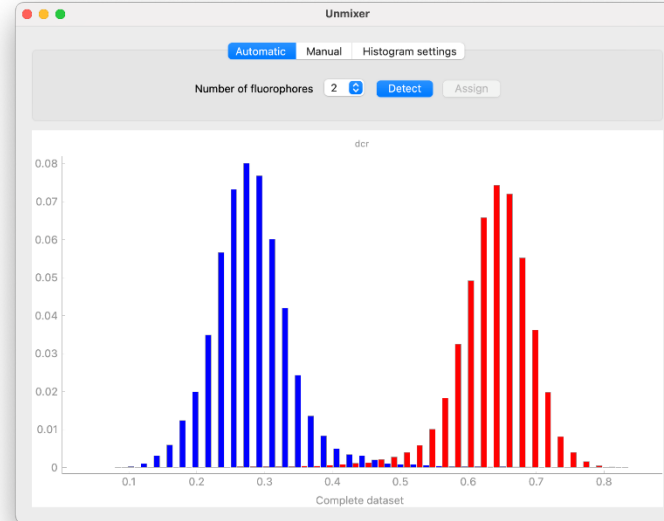
David  
Goodsell

U2OS Nup96-SNAP cells  
BG-Abberior Flux 640  
Ms-Nup153 + 2<sup>ary</sup> Flux 680  
GLOX-MEA (10-15 mM)

pyMINFLUX workflow



$$dcr = \frac{\text{Photons}_{\text{Detector1}}}{\text{Total}}$$



**EFO filtering**

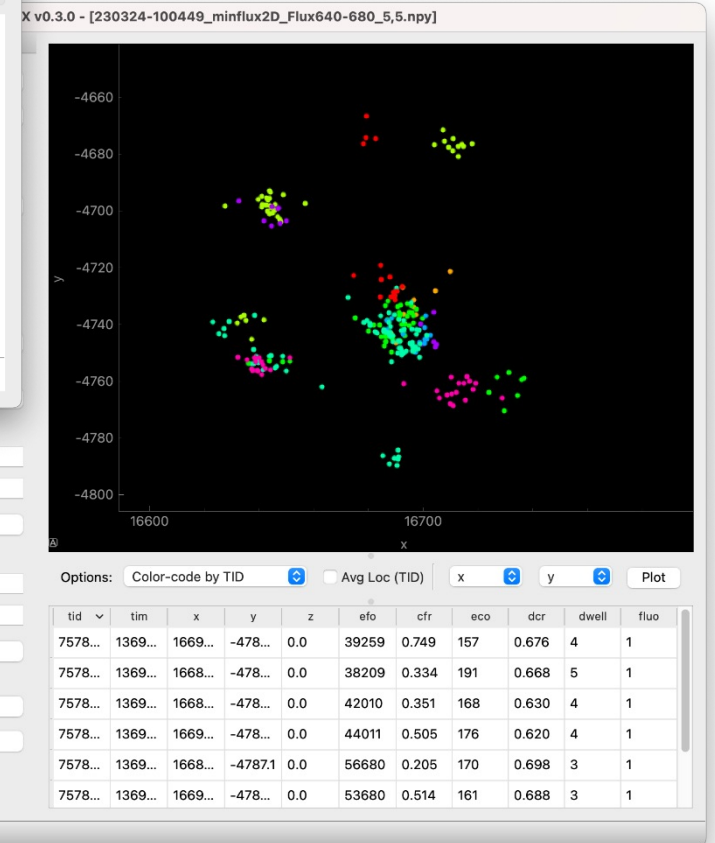
Lower bound (Hz)

Upper bound (Hz)

**CFR filtering**

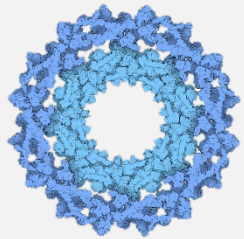
Lower bound

Upper bound

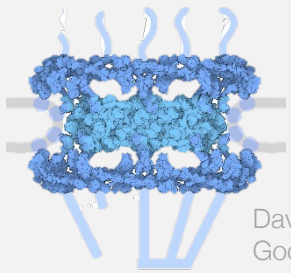


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view

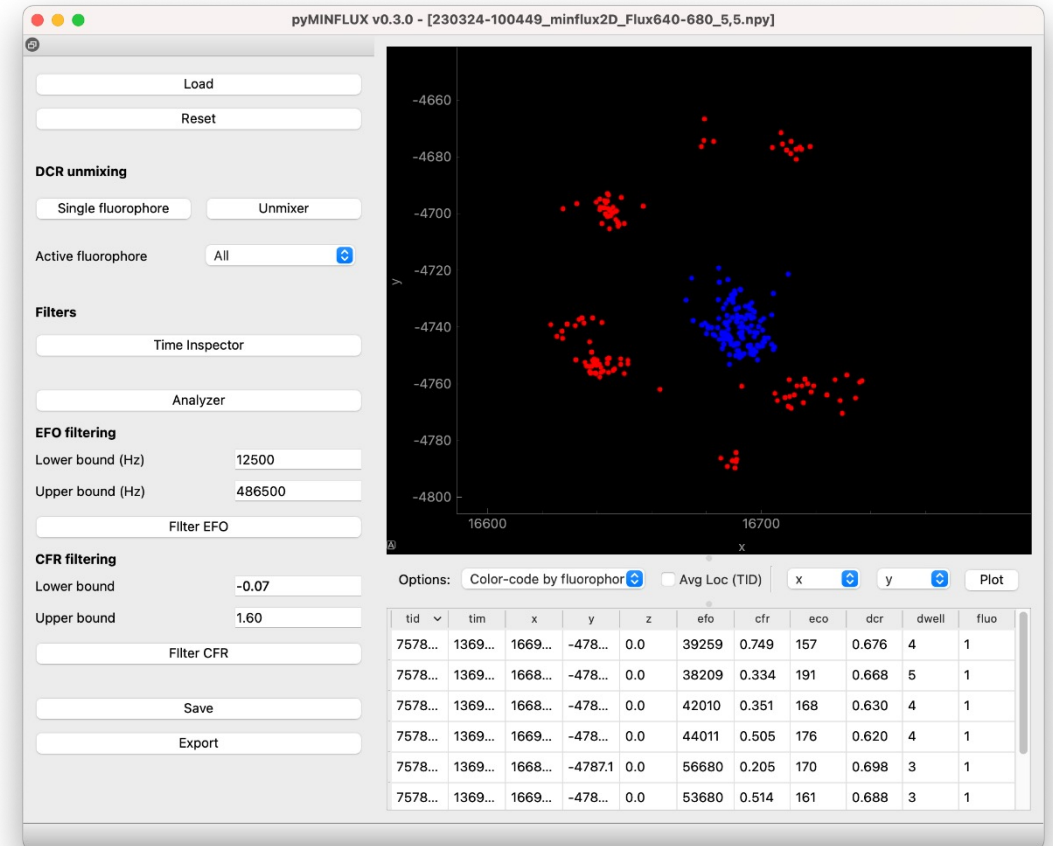
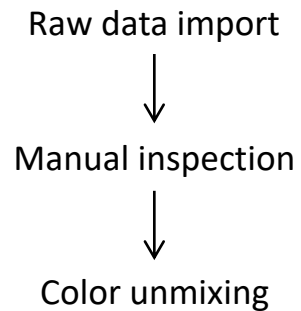


Side view

David  
Goodsell

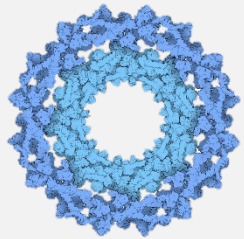
U2OS Nup96-SNAP cells  
BG-Abberior Flux 640  
Ms-Nup153 + 2<sup>ary</sup> Flux 680  
GLOX-MEA (10-15 mM)

pyMINFLUX workflow

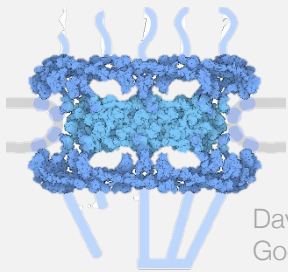


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view

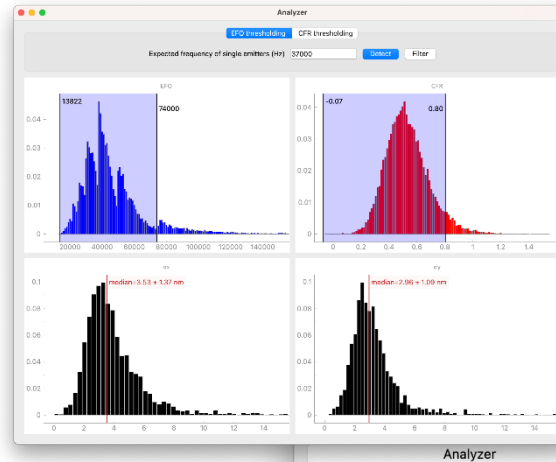
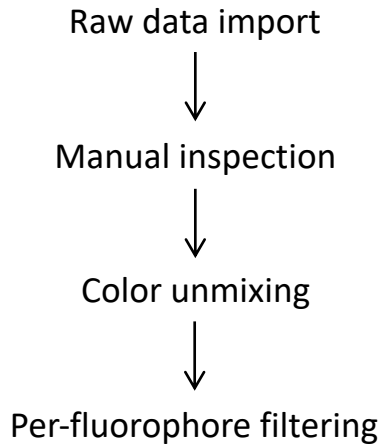


Side view

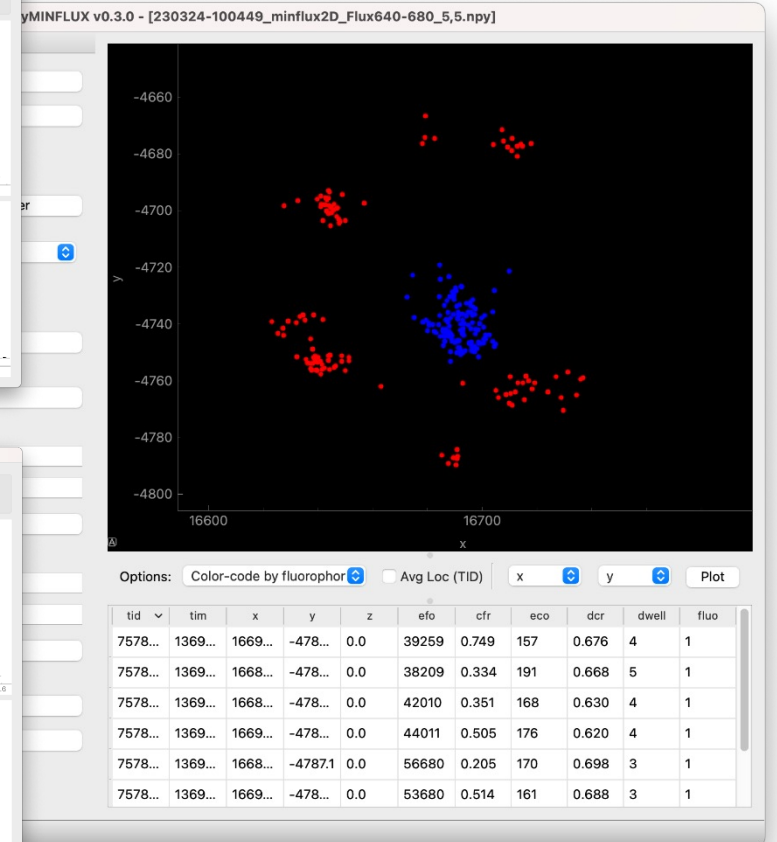
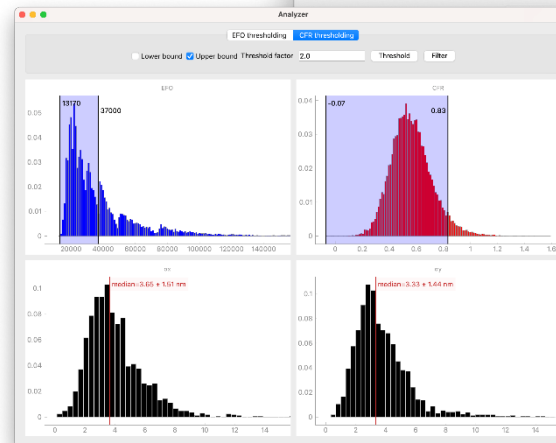
David Goodsell

U2OS Nup96-SNAP cells  
BG-Abberior Flux 640  
Ms-Nup153 + 2<sup>ary</sup> Flux 680  
GLOX-MEA (10-15 mM)

pyMINFLUX workflow

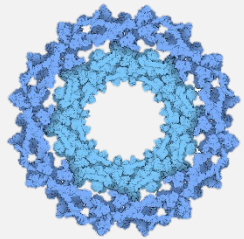


EFO filtering

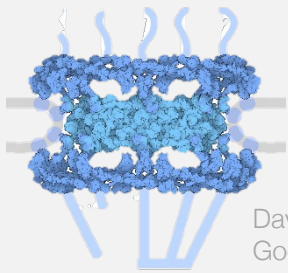


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view

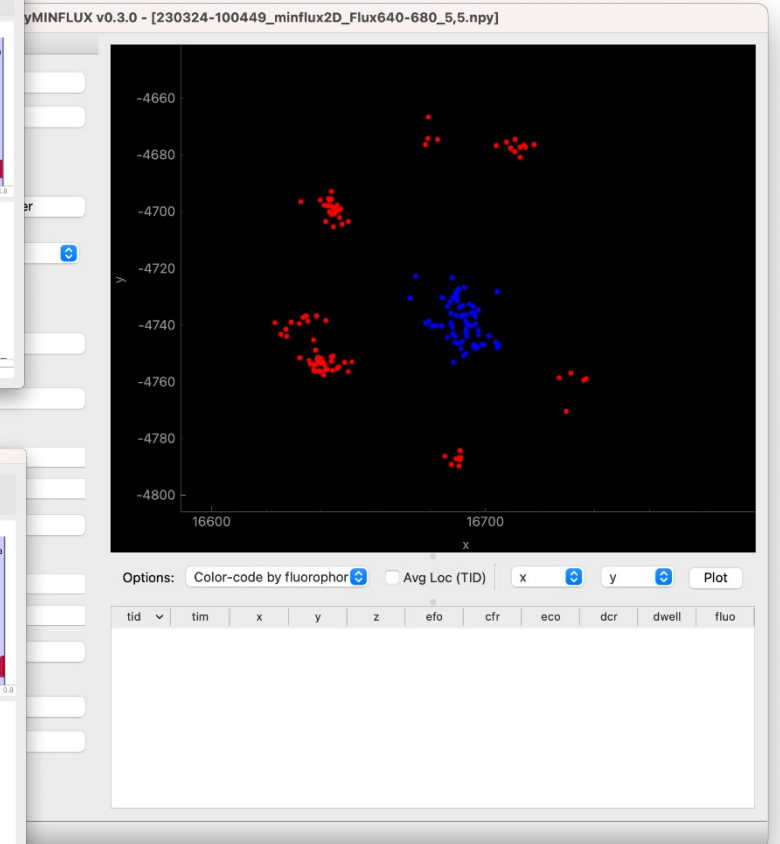
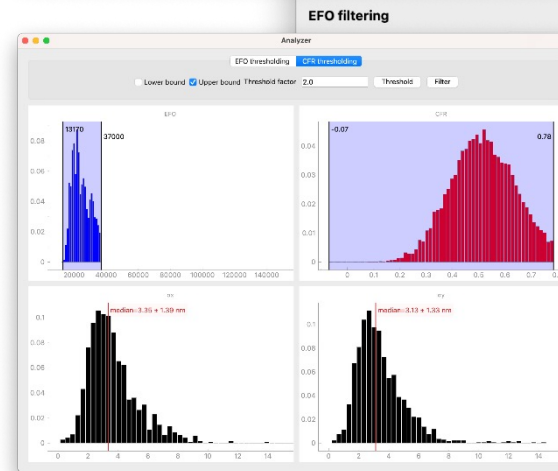
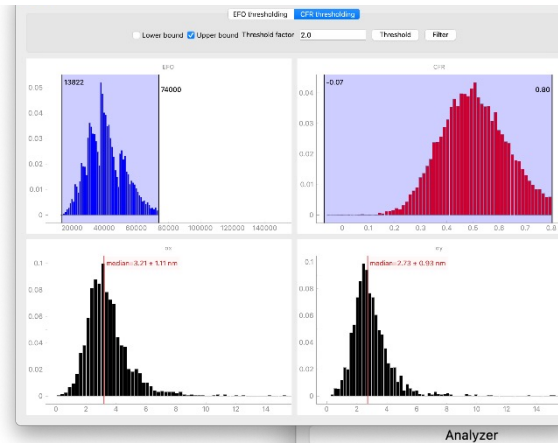
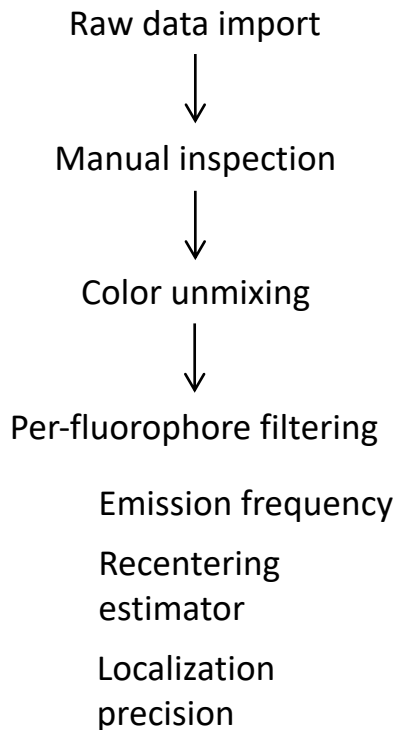


Side view

David Goodsell

U2OS Nup96-SNAP cells  
BG-Abberior Flux 640  
Ms-Nup153 + 2<sup>ary</sup> Flux 680  
GLOX-MEA (10-15 mM)

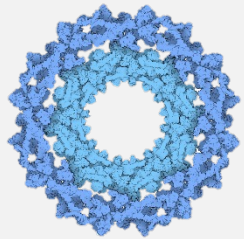
pyMINFLUX workflow



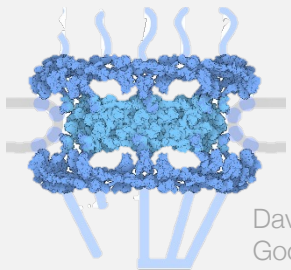


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view

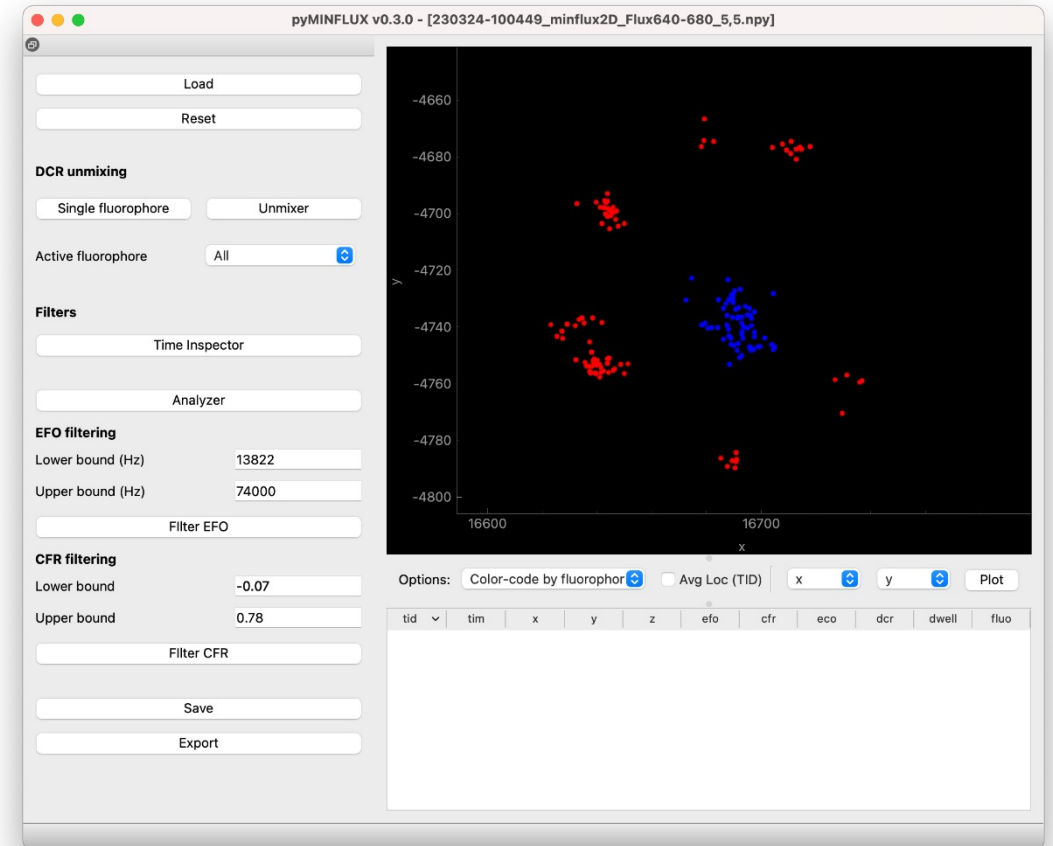
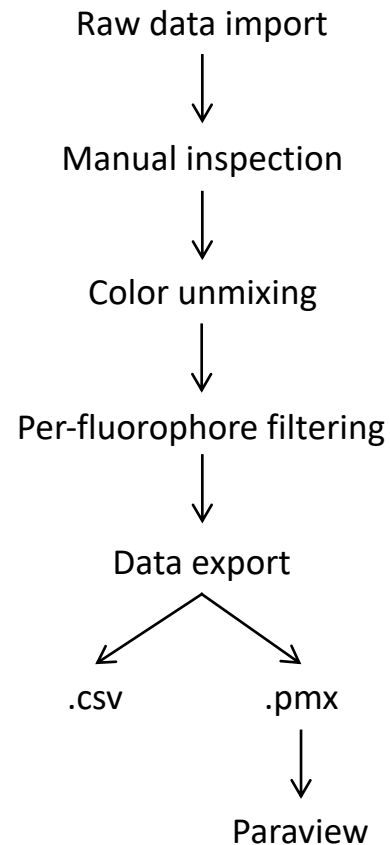


Side view

David  
Goodsell

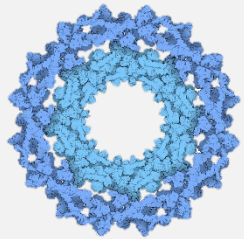
U2OS Nup96-SNAP cells  
BG-Abberior **Flux 640**  
Ms-Nup153 + 2<sup>ary</sup> **Flux 680**  
GLOX-MEA (10-15 mM)

pyMINFLUX workflow

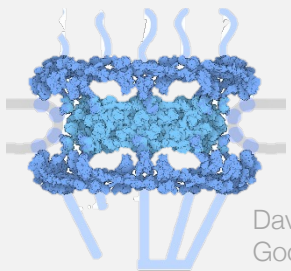


# Expert-driven projects :: pyMINFLUX

QC sample  
Nuclear pore complex



Top view

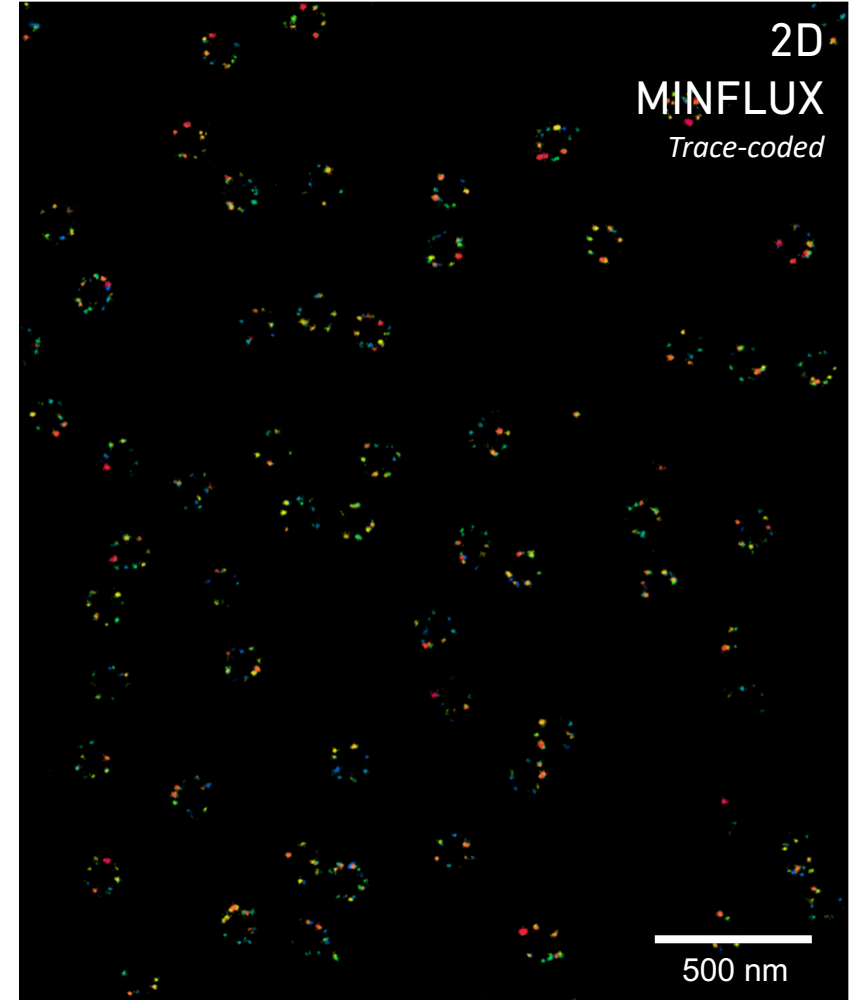
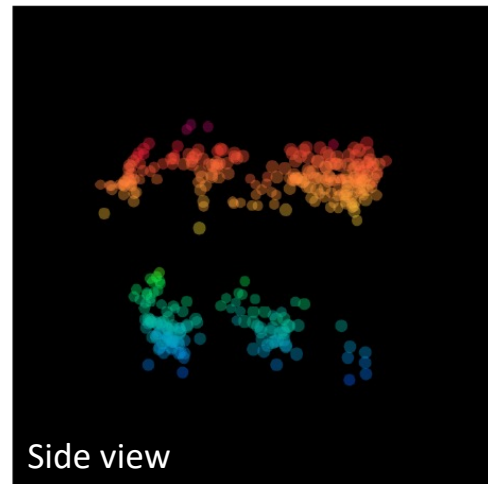
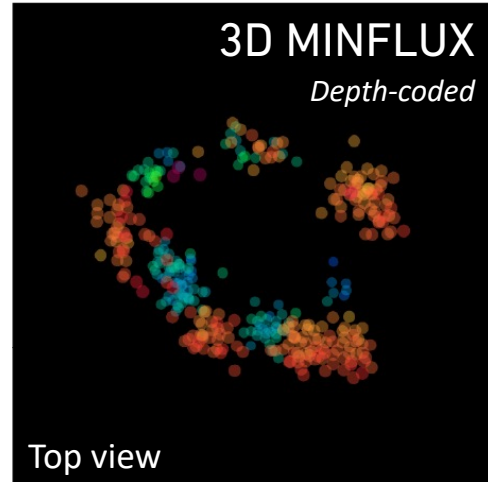
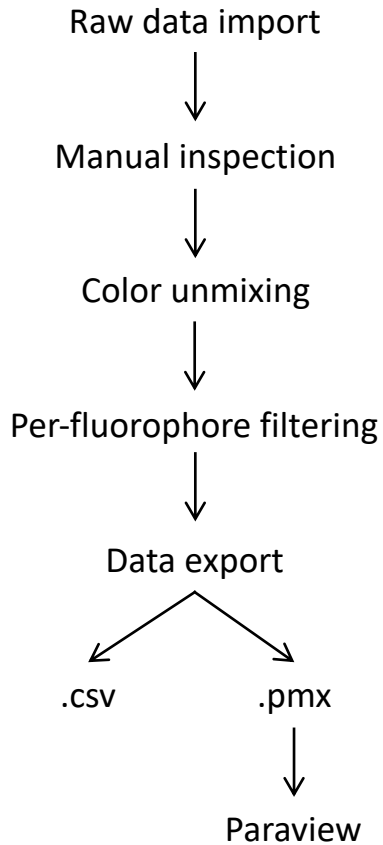


Side view

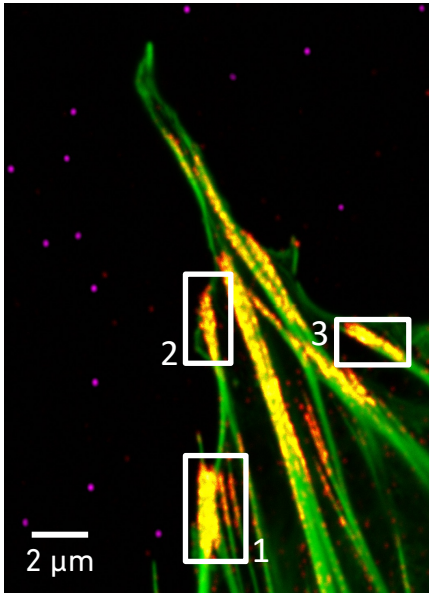
David  
Goodsell

U2OS Nup96-SNAP cells  
BG-Abberior Flux 640  
Ms-Nup153 + 2<sup>ary</sup> Flux 680  
GLOX-MEA (10-15 mM)

pyMINFLUX workflow



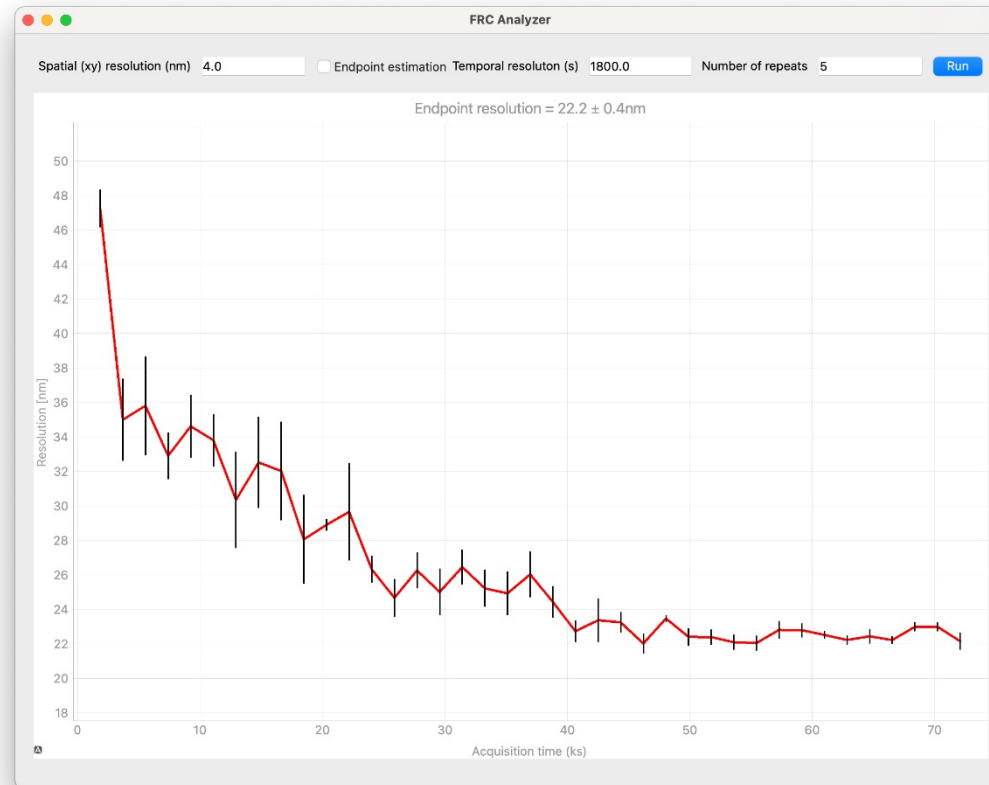
# Expert-driven projects :: pyMINFLUX



MEFs  
 Phalloidin-AF488  
 Rat-Integrin + 2<sup>ary</sup> AF568  
 Gold nanoparticles

Rb anti Paxillin  
 sdAb DNA-PAINT Atto 655

## Fourier-ring correlation analysis (“endpoint indicator”)



Nico Strohmeyer, D-BSSE

# Expert-driven projects :: pyMINFLUX

## Expert contributions

**Javier Casares Arias:** MINFLUX specialist, designed a lot of controlled experiments to test filtering and analysis strategies

**Aaron Ponti:** development of pyMINFLUX

## Users

- D-BSSE, Daniel Mueller Group (Michele Nava, Nico Strohmeyer, Matilde Lucioli, Krishna Kasuba)
- D-BSSE, Timm Schroeder Group (Germán Camargo)
- University of Basel, Thomas Ward Group (Michaela Slánská)
- University of Heidelberg (Charlotte Kaplan)

# Summary

There are different classes of (image analysis) projects with different types of requirements and target audiences:

- **User-specific projects** usually have a small scope and are solved in a tight feedback loop with the end user.
- **General-purpose projects** target large audiences with less specific sets of functionality and often require larger development teams and better software engineering practices.
- **Expert-driven projects** require tight collaboration between experts with different sets of field knowledge with the goal of creating tools that appeal to reasonably large but niche audiences.

# Acknowledgments

# Thank you for your attention!



**Thomas Horn** (head of SCF and LAF)

## Microscopy

Erica Montani  
Javier Casares Arias  
Tom Lummen

## Lab Automation

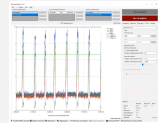
Sant Kumar  
Daniel Gerngross

## Flow Cytometry

Mariangela Di Tacchio  
Aleksandra Gumienny  
Chiara Cavallini

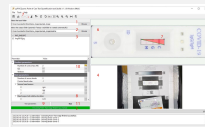
## SpectraSorter

Todd Duncombe, D-BSSE



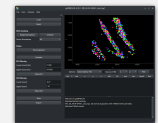
## pyPOCQuant

Fabian Rudolf, D-BSSE and BAG  
Andreas Cuny, D-BSSE



## pyMINFLUX

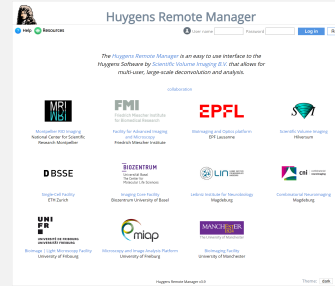
Javier Casares Arias, D-BSSE



## User-specific project

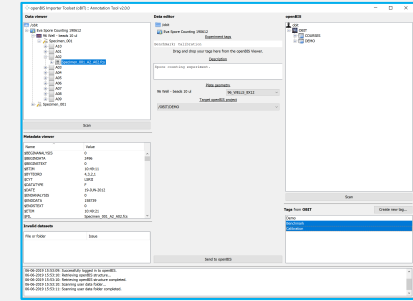
Gabriel Senn, D-BSSE

## HRM



Volker Bäcker, Montpellier Rio Imaging  
Daniel Sevilla, Scientific Volume Imaging  
Niko Ehrenfeuchter, Biozentrum  
Torsten Stöter, Leibniz Institute for Neurobiology  
Felix Meyenhofer, University of Fribourg  
Olivier Burri, EPFL  
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Alessandra Griffa, EPFL  
José Viña, Scientific Volume Imaging  
Kevin Namink, Scientific Volume Imaging  
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## oBIT



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Antti Luomi, SIS  
Jakub Straszewski, SIS  
Manuel Kohler, SIS  
Vernon Bailey, ITSC  
John Ryan, ITSC  
Vincenzo Spanò, ITSC  
Martin Fox, ITSC

**D BSSE**

All the users!