



Tuesday 3rd of March

Venue: (NOT SYMPOSIUM's Venue) Centre Broca Nouvelle Aquitaine

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| 14:00-15:00 | Helpdesk: Registration |
| 15:00 - 19:00 | SATELLITE MEETING for Bioimage Analysts |
| 15:00-16:45 | Session 1: Machine Learning in Microscopy <i>3x Talks:</i> Estibaliz Gomez de Mariscal, Dmitry Ershov, Dominik Kutra + Discussion |
| 16:45 | <i>Coffee</i> |
| 17:15-19:00 | Session 2: Biomage Analysis Facility Management <i>4x Talks</i> Stéphane Rigaud, Mafalda Sousa, Simon Noerrelykke, Perrine Paul-Gilloteaux + Discussion |
| 19:00- | <i>Free Evening</i> |

NEUBIAS 2020 at a glance

Wednesday 4th of March

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|-------|--|-------------------------------|
| 08:50 | INTRO/ | Welcome |
| 09:00 | Session 1 | 4x Talks + 13 FlashTalks (2") |
| 11:00 | <i>Coffee</i> | |
| 11:30 | Panel Discussion | |
| 12:50 | POSTERS | <i>with Lunch</i> |
| 14:30 | Session 2 | |
| | Keynote | SULIANA MANLEY |
| 15:15 | F1000Research, Companies Techbites (2") | |
| 15:40 | <i>Coffee</i> | |
| 16:10 | Company Workshops #1: Carl Zeiss, Nikon, ThermoFisher, Argolight | |
| | + OsSL | round 1 |
| 17:10 | OsSL | round 2 |
| 18:00 | OsSL | round 3 |
| 19:00 | <i>Free Evening</i> | |

Thursday 5th of March

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| 09:00 | Keynote | EMMA LUNDBERG |
| 09:45 | Session 3 | 2 Talks + Platinum Sponsors Talk (ZEISS, NIKON) |
| 11:00 | <i>Coffee</i> | |
| 11:25 | Session 4 | 5 Talks |
| 13:05 | POSTERS | <i>with lunch</i> |
| | 13:45 | NEUBIAS MC Meeting on invitation |
| 14:50 | Company Workshops #2: Carl Zeiss, Nikon, ThermoFisher, Argolight | |
| 15:50 | <i>Coffee</i> | |
| 16:40 | Session 5 | 2x Talks |
| 17:10 | CALL FOR HELP | |
| 18:40 | <i>Attendants registered to Gala dinner jump on the buses</i> | |
| 19:00 | <i>Buses Departure to Gala Dinner, sharp at 19:00 !</i> | |

Friday 6th of March

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| 09:00 | Keynote | KRISTIN BRANSON |
| 09:45 | Session 6 | <i>Special:</i> Deep Learning & ImageJ 3x Talks + 1 Talk |
| 10:55 | <i>Coffee</i> | |
| 11:25 | Session 7 | 3x Talks + 1x STSMs Report |
| 12:35 | <i>special session</i> | Bioimage Analysis, Perspectives |
| 13:30 | Closing comments | |
| 13:25 | Lunch, end and departure | |

NEUBIAS 2020 Symposium Detailed Program

Wednesday 4th of March 2020

08:00 – 08:50 Registration
08:50 – 09:00 Welcome by organizers *Florian Lvet (IINS)*
Fabrice Cordelières (BIC)
Sebastian Munck (VIB Leuven)

Session I Chair: *Marion Louveaux*

09:00 – 09:25 *invited* *Peter Bankhead*, University of Edinburgh (UK)
QuPath: Open source software (not just) for whole slide image analysis

09:25 – 09:40 *selected* *Allen Goodman & Beth Cimini*, The Broad Institute (Boston, USA)
Piximi: a free and open-source web app for object recognition

09:40 – 09:55 *selected* *Sébastien Tosi*, IRB Barcelona (ES)
LOBSTER: An environment to design bioimage analysis work flows for large and complex fluorescence microscopy data

09:55 – 10:10 *selected* *Christoph Moehl*, DZNE (Köln, DE)
Introducing YAPiC: An Open Source tool for biologists to perform complex image segmentation with deep learning

10:10 – 10:35 *invited* *Robert Haase*, MPI-CBG (Dresden, DE)
CLIJ: GPU-accelerated image processing for everyone

FLASH TALKS (Max 2'00")

selected *Estibaliz Gomez de Mariscal*, **Use of the p-value as a size-dependent function: model and applications**

Vanessa Isabell Jurtz, **Deep Learning reveals 3D atherosclerotic plaque distribution and composition**

David Barry, **GIANI: open-source software for automated analysis of 3D microscopy data**

Sylvain Prigent, **BioImage-IT: Design data analysis workflows using tools from different languages and keep track of the metadata**

Hannah Jeckel, **BiofilmQ, a software tool for quantitative image analysis of microbial biofilm communities**

Giada Bianchetti, **Microscopy pixel classification of intracellular sites of triglycerides and cholesteryl esters formation and storage through a machine-learning assisted polarity-driven segmentation workflow**

E. Josiah Lutton, **Quantitative analysis of macropinocytosis in Dictyostelium captured with light-sheet microscopy**

Carl-Magnus Svensson, Machine learning supported image analysis of microfluidic droplets: Using Random Forest classifiers and Bayesian inference for identification of experimental conditions

Dagmar Kainmueller, PatchPerPix for Instance Segmentation

Léo Guignard, Contact-area dependent cell communications and the morphological invariance of ascidian embryogenesis

Laura Nicolas-Saenz, Automatic, non-supervised image registration and segmentation for the study of tumor heterogeneity maps

Philippe Andrey, Statistical modeling of spatial interactions in biological patterns, with application to plant nuclear organization

Thomas Fish, GridSNAP: Automated Alignment of TEM Grid Mosaics for Correlative Imaging

11:05 - 11:35 Coffee

11:35 – 12:50 **Panel Discussion**

Moderator: Kota Miura

12:50 - 14:30 POSTER SESSION I with Lunch

Session II *Chair: Martin Jones*

Keynote Lecture

14:30 – 15:15 **SULIANA MANLEY**, École Polytechnique Fédérale de Lausanne (CH)
Superresolution fluorescence microscopy for multi-color 3D particle reconstruction

15:15 - 15:25 *Vicky Hellon, F1000Research (UK)*
F1000Research - Gateways
invited

15:25 – 15:40 **TECHBYTES**, 2.5" pitch talks by industry partners

15:40 - 16:10 Coffee

16:15 - 17:15

**Open source
Software Lounge
(OsSL)**

AND

Companies Workshop 1

3 rotations:

16:15 - 17:15

OsSL Round 1

with
Digital Poster by



DRVISION
Technologies LLC

4 parallel workshops:

16:15-17:15

Carl Zeiss Microscopy

Nikon

Thermo Fisher

Argolight



ThermoFisher
SCIENTIFIC



ARGOLIGHT
A Precision Company

17:15 - 18:10

OsSL Round 2

18:10 - 19:00

OsSL Round 3

19:00 onwards Free evening

Thursday 5th of March 2020

Session III

Chair: Bertrand Vernay

Keynote Lecture

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| 09:00 – 9:45 | <p>EMMA LUNDBERG SciLifeLab & KTH (Stockholm, SE) University of Stanford (USA)</p> <p>Image-based spatiotemporal dissection of the human proteome</p> |
| 9:45 – 10:10 <i>invited</i> | <p><i>Edward Cohen</i>, Imperial College (London, UK)</p> <p>Resolution limit of image analysis algorithms</p> |
| 10:10 – 10:25 <i>selected</i> | <p><i>James Klatzow</i>, EMBL-EBI (Cambridge, UK)</p> <p>Mesh Correspondence pipeline for Biological Shape Analysis</p> |
| 10:25 – 10:55 | <p><i>Sebastian Rhode</i>, Carl Zeiss Microscopy (DE)</p> <p>Opening the ZEN ecosystem for the Open Software Community – How to use your code and machine learning models inside ZEN using APEER and Intellesis</p> |
| Platinum Sponsors Talks | <p><i>Michael Davis</i>, NIKON (NL)</p> <p>NIS.ai - a new machine learning toolbox for Nikon's NIS-Elements software</p> |
| Session IV | <p>Coffee</p> <p>Chair: <i>Dagmar Kainmueller</i></p> |
| 11:30 – 11:55 <i>invited</i> | <p><i>Laure Blanc-Féraud</i>, CNRS (Sophia-Antipolis, FR)</p> <p>Fluorescence Microscopy 3D resolution improvement with MA-TIRF reconstruction.</p> |
| 11:55 – 12:10 <i>selected</i> | <p><i>Ko Sugawara</i>, IGFL / CNRS (Lyon, FR)</p> <p>Semi-automated cell tracking with incrementally trainable convolutional neural networks</p> |
| 12:10 – 12:25 <i>selected</i> | <p><i>Emmanuel Bouillhol</i>, Univ. Bordeaux (FR)</p> <p>Deep Learning approaches for reliable quantification of multi-omics cell imaging datasets to interrogate RNA and protein spatial and temporal subcellular interactions</p> |
| 12:25 – 12:40 <i>selected</i> | <p><i>Minh Son Phan</i>, CNRS/INSERM/école Polytechnique (Palaiseau,FR)</p> <p>Quantitative geometry methods for analyzing 3D neuronal trajectories with GeNePy3D</p> |
| 12:40 – 13:05 <i>invited</i> | <p><i>Tammy Riklin Raviv</i>, Ben Gurion Univ. (IL)</p> <p>Diving Deep into Cell Segmentation in Microscopy Videos</p> |

13:05 - 14:50 POSTER SESSION II with Lunch
+ 13:45 - 15:45 NEUBIAS MC Meeting (on invitation)

14:50 - 15:50 | 4 parallel workshops:

Carl Zeiss Microscopy

Nikon

Thermo Fisher

Argolight



ARGOLIGHT
A Precision Company

ThermoFisher
SCIENTIFIC

15:50 - 16:10 Coffee

Session V

Chair: Arrate Muñoz-Barrutia

16:10 – 16:35
invited

Marcel Mueller, Univ. Bielefeld (DE)
fairSIM - image processing for structured illumination microscopy

16:35 – 17:00
invited

Thomas Walter, MinesTech /Curie (Paris, FR)
Machine Learning for computational phenotyping: how to get around massive manual annotation

17:00 – 18:30

“Call For Help” aka Shout your problem, Analysis Clinics....

Chair: Jan Eglinger,

Team: Laure Plantard, Szymon Stoma, ...

Interactive Session: *Life Scientists and BioImage Analysts facing Image Analysis Roadblocks in their research projects have submitted their problem, Developers and Analysts will address them, build and benchmark solutions. 10' per project with plenary discussion*

19:00 Sharp

Bus departure to Gala Dinner

Wine Castle Visit + Dinner

Return Bus from 23h (about 40min drive) to a Bordeaux center and Venue

Friday 8th of February 2019



Session VI

Chair: Sebastian Munck

Keynote Lecture

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| 09:00 – 9:45 | KRISTIN BRANSON , HHMI Janelia Campus (Ashburn, USA) TBA |
| 9:45 – 10:40 | <i>Special Session: Deep Learning and ImageJ</i> |
| 9:45 - 10:00 <i>invited</i> | Deborah Schmidt , MPI-CBG (Dresden, DE) Deep dive with Fiji |
| 10:00 - 10h15 <i>selected</i> | Arrate Muñoz Barrutia , Univ. Carlos III (Madrid, ES) DeepImageJ: Bridging Deep Learning to ImageJ |
| 10:15 - 10h30 <i>selected</i> | Martin Weigert , MPI-CBG (Dresden, DE) Nuclei Detection and Segmentation in Fluorescence Microscopy Images with Learned Shape Representations + <i>Discussion</i> |
| 10:40 – 10:55 <i>selected</i> | Elisabeth Wetzer , Uppsala Univ. (SE) Cross-modal Representation Learning for Efficient Registration of Multiphoton and Brightfield Microscopy Images of Skin Tissue |
| 10:55 - 11:25 | Coffee |
| Session VII | Chair: Natasa Sladoje |
| 11:25 – 11:50 <i>invited</i> | Fabrice de Chaumont , Pasteur (Paris, FR) Real time tracking and analysis of mice in rich environment |
| 11:50 – 12:05 <i>selected</i> | Nadezhda Koriakina , Univ. Uppsala (SE) Uncovering hidden reasoning of convolutional neural networks in biomedical image classification by using attribution methods |
| 12:05 – 12:20 <i>selected</i> | Lassi Paavolainen , FIMM (Helsinki, FI) Learning representation of microscopy images by weakly-supervised deep learning |
| 12:20 – 12:45 | <i>Short-term Scientific Missions: Julia Fernandez-Rodriguez</i> Devrim Unay , Izmir University (TR) Automated detection and segmentation of bacteria from multimodal microscopy images Ana Stojiljkovic , University of Bern (CH) High-content characterisation of canine induced pluripotent stem cells |
| 12:40 – 13:20 | <i>Bioimage Analysis Perspectives, NEUBIAS TEAM</i> |
| 13:30 | <i>Closing Comments and Lunch</i> |