

From Images to Knowledge with ImageJ & Friends

EMBL CONFERENCE

We have moved our website to embl.org/events. The content below is no longer being updated.

EMBL Courses and Conferences during the Coronavirus pandemic

With the onsite programme paused, many of our events are now being offered in virtual formats.

Registration is open as usual for many events, with back-up plans in place to move further courses and conferences online as necessary. Registration fees for any events affected by the COVID-19 disruption are fully refundable.

More information for participants of events at EMBL Heidelberg can be found here.

Programme

Got something to say? Tweet it! #EMBLImageJ #I2K

HIDE ALL

Day 1 - Thursday 6 December 2018

Time	Speaker
11:00 - 12:45	Arrival and registration
12:45 - 13:00	Welcome and opening remarks
13:00 - 13:45	ImageJ Curtis Rueden - Laboratory for Optical and Computational Instrumentation, USA Tobias Pietzsch - Max Planck Institute of Molecular Cell Biology and Genetics, Germany Christian Dietz - KNIME, Germany

Time	Speaker
13:45-15:00	Session 1 - Light microscopy 1 Chair: Pavel Tomancak, Max Planck Institute of Molecular Cell Biology and Genetics, Germany
13:45-14:15	Using morphological profiling to extract the most from your images Beth Cimini - Broad Institute, USA
14:15-14:45	Title TBC Jeremy Freeman - Chan Zuckerberg Initiative, USA
14:45-15:00	Quantitative imaging of motile cilia to investigate fluid flows and chirality in the left-right organizer Willy Supatto - Ecole polytechnique, CNRS, INSERM, France
15:00-15:30	Coffee break and Meet the Invited Speaker: Beth Cimini, Jeremy Freeman
15:30-18:00	Session 2 - Machine learning Chair: Anna Kreshuk - EMBL Heidelberg, Germany
15:30-16:00	The road to deeper ilastik Anna Kreshuk - EMBL Heidelberg, Germany
16:00 - 16:15	KNIME Image Processing: Status Quo and Future Directions Christian Dietz - KNIME, Germany
16:15 - 16:30	Introducing Yapic: A New Tool for Biologists to Perform Complex Image Segmentation with Deep Learning Christoph Möhl - German Center for Neurodegenerative Disease (DZNE), Germany
16:30 - 17:00	Coffee break
17:00 - 17:30	Content-Aware Image Restoration for Light and Electron Microscopy Facilitates Quantitative Data Analysis Florian Jug - Max Planck Institute of Molecular Cell Biology and Genetics, Germany
17:30 - 17:45	Automated Annotation of C. Elegans: How much Training Data is Needed? Dagmar Kainmueller - Berlin Institute of Health, Max Delbrück Center for Molecular Medicine, Germany
17:45-18:00	Flash talks (1 slide/2mins each) Albert #23 / Botelho #31/ Bujak #35 / Meyenhofer #79 / Schmidt #95
18:00-19:30	Poster Session 1 (odd numbers)
19:30	Free evening

Day 2 - Friday 7 December 2018

Time	Speaker
09:00-10:15	Session 3 - Light microscopy 2 Chair: Florian Jug - Max Planck Institute of Molecular Cell Biology and Genetics, Germany
09:00-09:30	Correlative Light-Electron Microscopy Based High Resolution Nuclear Model of C. elegans Dauer Ella Bahry - Max Delbrück Center for Molecular Medicine, Germany
09:30-10:00	NanoJ-SQUIRREL: quantitative mapping and minimisation of super- resolution optical imaging artefacts Siân Culley - University College London, UK
10:00-10:15	BigStitcher: Reconstructing high-resolution image datasets of cleared and expanded samples Stephan Preibisch - Max Delbrück Center for Molecular Medicine, Germany
10:15-10:45	Coffee break and Meet the Invited Speaker: Stephan Preibisch
10:45-12:00	Session 4 - Electron Microscopy analysis Chair: Christian Tischer - EMBL Heidelberg, Germany
10:45-11:15	Mapping and analyzing neuronal circuits with CATMAID software Albert Cardona - Janelia Research Campus, USA
11:15-11:45	N5, ImgLib2-Cache, BigDataViewer, and Spark: my favorite open- source eco-system for super-large image data Stephan Saalfeld - Howard Hughes Medical Institute, USA
11:45-12:00	webKnossos: efficient online 3D data annotation Norman Rzepka - scalable minds GmbH, Germany
12:00-13:30	Lunch
13:30-15:00	5 Parallel Workshops - Session 1
15:00-15:30	Coffee break and Meet the Invited Speakers: Albert Cardona, Stephan Saalfeld
15:30-16:55	5 Parallel Workshops - Session 2
17:00-17:15	NEUBIAS talk Julien Colombelli, IRB Barcelona, Spain

Time	Speaker
17:15 - 17:30	A natural-language based syntax for animating 3D/4D imaging data Benjamin Schmid - Friedrich-Alexander University of Erlangen- Nuremberg, Germany
17:30 - 17:45	Mastodon a large-scale track-editing framework for Fiji Tobias Pietzsch - Max Planck Institute of Molecular Cell Biology and Genetics, Germany
17:45-18:00	Flash Talks (1 slide / 2mins each) Arzt #25 / Cheeseman #41 / Günther #54 / Jain #63 / Mais #75
18:00-19:30	Poster Session 2 (even numbers)
19:30-21:30	Conference Dinner
21:30-00:00	Conference Party

Day 3 - Saturday 8 December 2018

Time	Speaker
09:00-10:15	Sesion 5 - Atlas projects Chair: Christophe Zimmer - <i>Institut Pasteur, France</i>
09:00-09:30	Deep learning as a tool in microscopy data analysis and digital pathology Carolina Wählby - Uppsala University, Sweden
09:30-10:00	Archives, Resources and Atlases: An OME Perspective on Structuring Data for the Long Haul Josh Moore - University of Dundee, UK
10:00-10:15	The mitotic cell atlas Jean-Karim Heriche - EMBL Heidelberg, Germany
10:15-10:45	Coffee break and Meet the Invited Speaker: Josh Moore, Virginie Uhlmann, Christoph Zimmer
10:45-12:00	Session 6 - Tools and Methods Josh Moore - University of Dundee, UK
10:45-11:15	New computational lenses for single molecule super-resolution microscopy Christophe Zimmer - Institut Pasteur, France
11:15-11:45	Title TBC Virginie Uhlmann - EMBL-EBI, United Kingdom

Time	Speaker
11:45-12:00	Example of an end-to-end application of imaging and image analysis to solve a biological problem Pavel Tomancak, Max Planck Institute of Molecular Cell Biology and Genetics, Germany
12:00-12:15	Closing remarks and Poster Prizes
12:15-12:45	Packed lunch and departure
12:45	Bus departures downtown and to Frankfurt Airport