

Sunday, April 10

3:00 pm	Check-in
6:00 pm	Reception (<i>Lobby</i>)
7:00 pm	Dinner
8:00 pm	Welcome and Opening Remarks
8:05 pm	Introduction of Keynote by Sydney Brenner
8:10 pm	Keynote Lecture: John G. White , University of Wisconsin-Madison <i>The birth of connectomics</i>
9:10 pm	Refreshments available at Bob's Pub

NOTE:

Meals are in the **Dining Room**

Talks are in the **Auditorium**

Posters are in the **Lobby**

Monday, April 11

- 7:30 am Breakfast (*service ends at 8:45am*)
- 9:00 am Session 1**
Chair: Winfried Denk
- 9:00 am **Kristen M. Harris**, University of Texas at Austin
Recognizing structural differences in synaptic efficacy
- 9:10 am **Harald F. Hess**, Janelia Research Campus/HHMI
Expanding FIBSEM capabilities for whole fly brain connectome
- 9:20 am **Kenneth J. Hayworth**, Janelia Research Campus/HHMI
Random access, parallel FIBSEM imaging of the Drosophila brain
- 9:30 am **Stephan Saalfeld**, Janelia Research Campus/HHMI
Artifact correction in large EM volumes
- 9:40 am **Graham W. Knott**, Ecole Polytechnique Federale de Lausanne (EPFL)
Ultrastructural analysis of the aging brain
- 9:50 am Break
- 10:40 am Session 2**
Chair: Eve Marder
- 10:40 am **Juergen Tritthardt**, Max Planck Institute for Neurobiology
Speeding up SBEM data acquisition with piezo driven line scanning
- 10:50 am **Dirk Zeidler**, Carl Zeiss Microscopy
Advancing the throughput of a multi-beam SEM
- 11:00 am **Dan J. Bumbarger**, Allen Institute for Brain Science
Improvements in section handling for high throughput transmission electron microscopy
- 11:10 am **Shawn Mikula**, Max Planck Institute for Neurobiology
Mammalian whole-brain preparation for neural circuit reconstruction
- 11:20 pm **Sven Dorkenwald**, Max Planck Institute for Medical Research
An automated synaptic connectivity inference pipeline to generate annotated connectomes
- 11:30 am Short discussion or break for lunch
- 12:00 pm Lunch (*service ends at 1pm*)

1:30 pm	Session 3 Chair: Mitya Chklovskii
1:30 pm	Edward S. Boyden , Massachusetts Institute of Technology <i>Expansion microscopy</i>
1:40 pm	JoAnn Buchanan , Allen Institute for Brain Science <i>Protocol development for high throughput 3D electron microscopy</i>
1:50 pm	Mark H. Ellisman , University of California, San Diego <i>New labeling and imaging methods applied to multiscale and multimodal mapping of projections of melanopsin retinal ganglion cells to multiple brain regions</i>
2:00 pm	Stephen J. Smith , Allen Institute for Brain Science <i>Shotgun connectomics</i>
2:10 pm	Rachel O. Wong , University of Washington <i>Combined fluorescence imaging and serial block face scanning electron microscopy identifies a new cell type in the retina</i>
2:20 pm	Break
3:00 pm	Session 4 Chair: Viren Jain
3:00 pm	Alexander Borst , Max Planck Institute of Neurobiology <i>Functional analysis of the fly's optic lobe connectome</i>
3:10 pm	Lou K. Scheffer , Janelia Research Campus/HHMI <i>Analysis of EM reconstructions</i>
3:20 pm	Kara Fulton (Briggman Lab) , NINDS/NIH <i>Species dependent wiring of direction selective circuitry in the mammalian retina</i>
3:30 pm	Clay Reid , Allen Institute for Brain Science <i>Anatomy and function of an excitatory network in the visual cortex</i>
3:40 pm	Sebastian Seung , Princeton University <i>Unbiased anatomical classification of retinal ganglion cells via serial electron microscopy</i>
3:50 pm	Break
4:15 pm	Discussion (Moderated by Winfried Denk)
5:30 pm	Poster reception
7:00 pm	Dinner
8:00 pm	Keynote Lecture: Eve Marder , Brandeis University <i>Going from the connectome to understanding circuit dynamics</i>
9:00 pm	Refreshments available at Bob's Pub

Tuesday, April 12

- 7:30 am Breakfast (*service ends at 8:45am*)
- 9:00 am Session 5**
Chair: Gerry Rubin
- 9:00 am **Kevan AC Martin**, Swiss Federal Institute of Technology, Zurich
Brains as potential technology
- 9:10 am **Fabian Svara**, Max Planck Institute of Neurobiology
Modularity in the larval zebrafish spinal cord
- 9:20 am **Joergen M. R. Kornfeld**, Max Planck Institute of Neurobiology
Connectomic analysis of zebra finch HVC
- 9:30 am **Kevin Boergens**, Max Planck Institute for Brain Research
Local connectome statistics in mouse barrel cortex
- 9:40 am **Michale S. Fee**, Massachusetts Institute of Technology
High-resolution reconstruction of cortico-basal-ganglia circuits underlying vocal learning
- 9:50 am Break
- 10:40 am Session 6**
Chair: Fred Hamprecht
- 10:40 am **Marta Zlatic**, Janelia Research Campus/HHMI
Circuit principles of memory-based behavioral choice
- 10:50 am **Davi Bock**, Janelia Research Campus/HHMI
*Reconstructions of neuronal connectivity in the mushroom body of *Drosophila melanogaster* from a whole-brain transmission electron microscopy volume*
- 11:00 am **Shinya Takemura**, Janelia Research Campus/HHMI
*FIB-SEM connectome reconstructions of the visual and olfactory centers of the *Drosophila* brain*
- 11:10 am **Wei-Chung Allen A. Lee**, Harvard Medical School
Functional neuronal assemblies for sensory processing
- 11:20 am **Benjamin Titze**, Friedrich Miescher Institute for Biomedical Research
Towards structure-function analysis of higher olfactory circuits in zebrafish
- 11:30 am Short discussion or break for lunch
- 12:00 pm Lunch (*service ends at 1pm*)
- 1:00 pm Tour (*optional - meet at reception*)

- 2:00 pm** **Session 7**
Chair: Kevan Martin
- 2:00 pm **Rainer W. Friedrich**, Friedrich Miescher Institute for Biomedical Research
Reconstructing olfactory circuits and computations in zebrafish
- 2:10 pm **Christel Genoud**, Friedrich Miescher Institute for Biomedical Research
Conductive sample preparation for SBEM and applications in the zebrafish olfactory bulb
- 2:20 pm **Albert Cardona**, Janelia Research Campus/HHMI
Collaborative reconstruction of neural circuits with CATMAID
- 2:30 pm **Hanspeter Pfister**, Harvard University
High-throughput connectomics with MapRecurse
- 2:40 pm **Scott W. Emmons**, Albert Einstein College of Medicine
Whole-animal C. elegans adult connectomes
- 2:50 pm Break
- 3:30 pm** **Session 8**
Chair: Clay Reid
- 3:30 pm **Verena S. Kaynig**, Harvard University
Scaling up automatic neuron reconstruction to the cubic millimeter
- 3:40 pm **Stephen M. Plaza**, Janelia Research Campus/HHMI
Large-scale EM connectome reconstruction in the adult fly brain
- 3:50 pm **Jeff W. Lichtman**, Harvard University
Connectomics from the top down and/or the bottom up
- 4:00 pm **Dmitri Chklovskii**, Simons Center for Data Analysis
Derivation of neural circuits from the similarity matching principle
- 4:10 pm Break
- 4:30 pm Discussion (Moderated by Gerry Rubin)
- 5:45 pm Poster reception
- 7:00 pm Dinner
- 8:00 pm** **Pietro V. De Camilli**, HHMI/Yale University
Taking a look at the inner world of neurons
- 9:00 pm Refreshments available at Bob's Pub

Wednesday, April 13

- 7:30 am Breakfast
- 9:00 am Session 9** (*machine learning / segmentation*)
Chair: Stephan Saalfeld
- 9:00 am **Moritz Helmstaedter**, Max Planck Institute for Brain Research
Connectome phenotypes in the cerebral cortex
- 9:10 am **Fred A. Hamprecht**, University of Heidelberg
Automated tracing: Learning from few labels
- 9:20 am **Viren Jain**, Google
Automated reconstruction of EM connectomic data with large scale machine learning and computation
- 9:30 am **Srini C. Turaga**, Janelia Research Campus/HHMI
New machine learning algorithms for image segmentation, and statistical models for interpreting and connecting neural connectivity to neural activity
- 9:40 am **Anna Kreshuk**, University of Heidelberg
Exploiting prior knowledge for automated EM image analysis
- 9:50 am Break
- 10:30 am Session 10**
Chair: Kristen Harris
- 10:30 am **Michal Januszewski**, Google
Recurrent neural networks for EM data segmentation
- 10:40 am **Seymour Knowles-Barley**, Harvard University
Rhoana: Dense automatic neuron annotation - case studies
- 10:50 am Closing Discussion and Final Remarks
- 11:30 am Lunch and Departure
- 12:00 pm First shuttle to Dulles
- 1:00 pm Second shuttle to Dulles
- 2:00 pm Last shuttle to Dulles