

Brain Connectivity Mapping

IJCNN Tutorial

August 4, 2013

Yoonsuck Choe, Ph.D.

Brain Networks Laboratory

Department of Computer Science and Engineering

Texas A&M University

Joint work with: L. C. Abbott, J. Keyser, B. McCormick, D. Han, J. Kwon, D. Mayerich, D. E. Miller, J. R. Chung, C. Sung.

Introduction and Overview

Mapping Brain Connectivity



Organism

C57BL/6 mouse

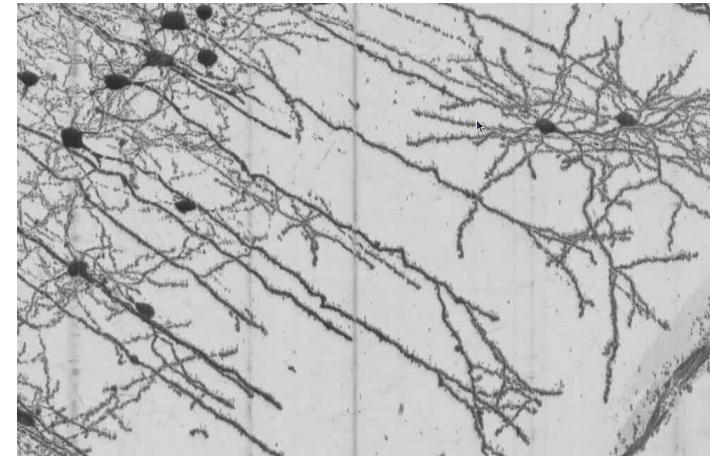
<http://mouseatlas.org>



Brain

Mouse brain

<http://nervenet.org>



Connectivity

Brain circuits (Mouse cortex)

- First step toward Understanding brain function: from structure to function.
- Approach: Omics

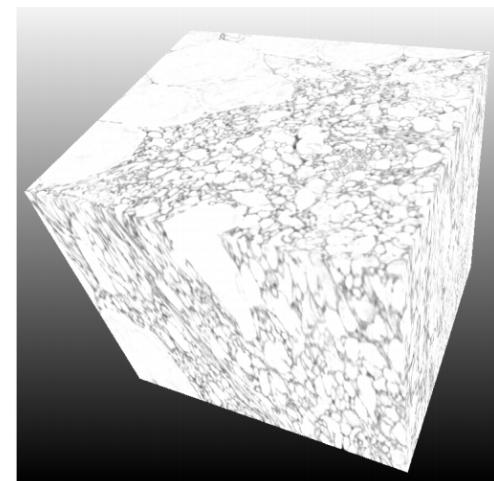
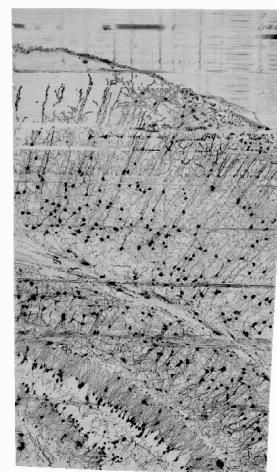
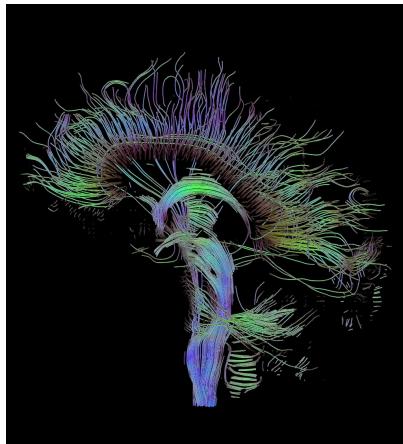
Age of the “Omics”

Biology has entered the age of “Omics”.

- “X-ome” means a complete collection of X
 - Derived from $-\omega\mu\alpha$ (-oma) in Greek.
 - “X-omics” means the study of “X-ome”.
- Examples:
 - genome, proteome, metabolome, physiome, etc.
- Why study “omics”?
 - Can understand how the whole system works.

Connectomics

Connectome: Collection of all connections between all neurons in the brain (Sporns et al. 2005; Sporns 2012; Seung 2012).



Imaging: Diffusion Tensor Imaging (DTI)

Scale: $\sim 10 \text{ cm cube}$

Whole Human brain

Resolution: $\sim 1 \text{ mm cube}$

Time: hours

See e.g. Hagmann et al. (2007)

Light Microscopy

$\sim 1 \text{ cm cube}$

Whole Mouse brain

$\sim 1 \mu\text{m cube}$

weeks

Mayerich et al. (2008)

Electron Microscopy

$\sim 100 \mu\text{m cube}$

Hundreds of neurons

$\sim 10 \text{ nm cube}$

year

Denk and Horstmann (2004)

Why Connectomics?

- Brain evolution is mostly evolution of the architecture (connectome), not the elements (neurons) (Swanson 2003)
- Current state of neuroscience is too specialized, local, and fragmented.
- Huge accumulation of (local) experimental (anatomical, physiological, genetic, behavioral) data.
- Need a framework to integrate the scattered data for a system-level understanding of the brain.

Current Status of Connectomics

- Nematode *C. elegans*: Only available connectome (White et al. 1986).
- Mostly focused on data acquisition (microscopy and imaging).
- Analysis framework leading behind.

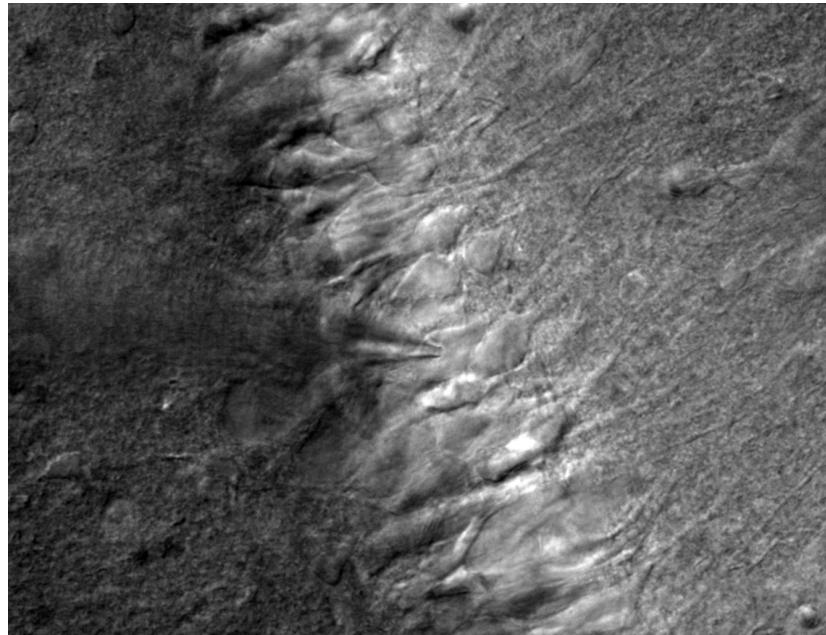
Overview

1. Staining and Labeling
2. Imaging
3. Data and Online Resources
4. Analysis
5. Wrap Up

Part I

Staining and Labeling

Staining and Labeling



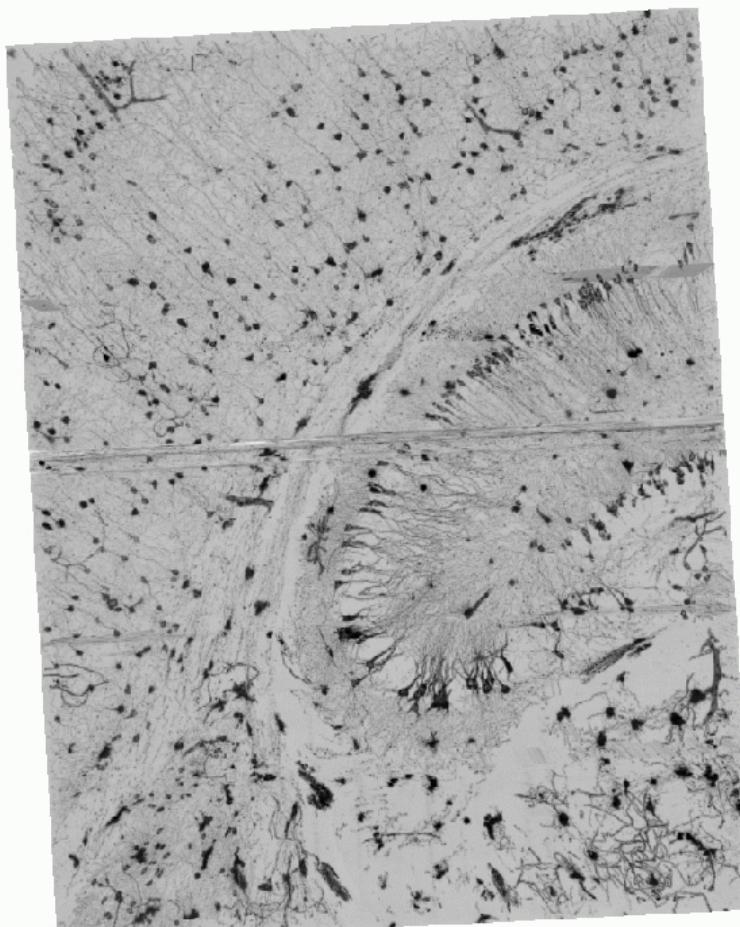
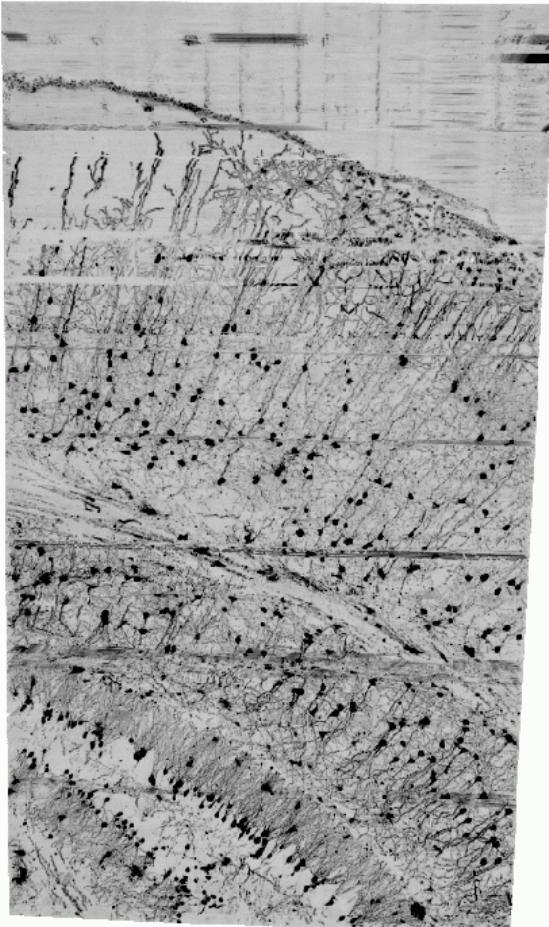
<http://commons.wikimedia.org/wiki/File:WholeCellPatchClamp.jpg>

- Need: Very low contrast between neurons and non-neuronal cells/tissue in the brain (see image above).
- Chemical stains and molecular labels are used to provide contrast.

Types of Stains/Labels

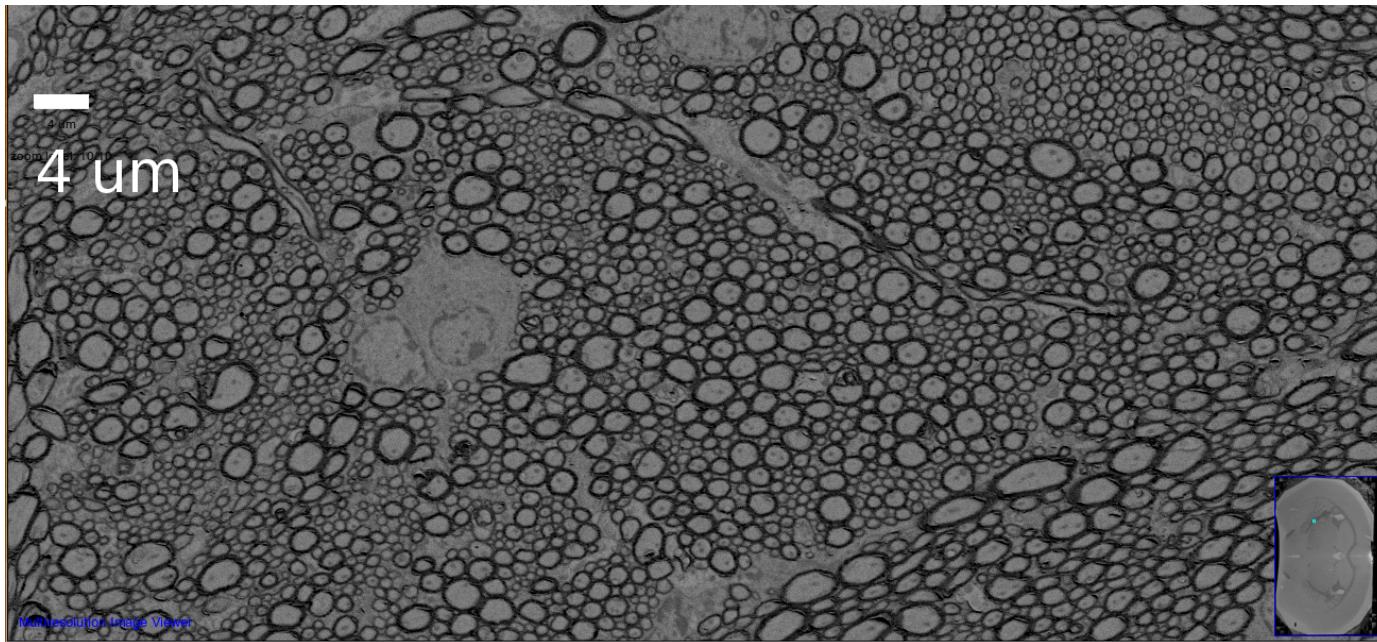
- Sparse (few neurons marked) vs. dense (all neurons marked)
- Random (random population marked) vs. targetted (specific cell types marked).
- For use with different imaging methods: light microscopy, electron microscopy, fluorescence microscopy, etc.

Golgi



- Stains whole neurons (axons unreliably stained): Sparse ($\sim 1\%$ stained), Random, Whole brains can be stained.
Ideal for light microscopy.

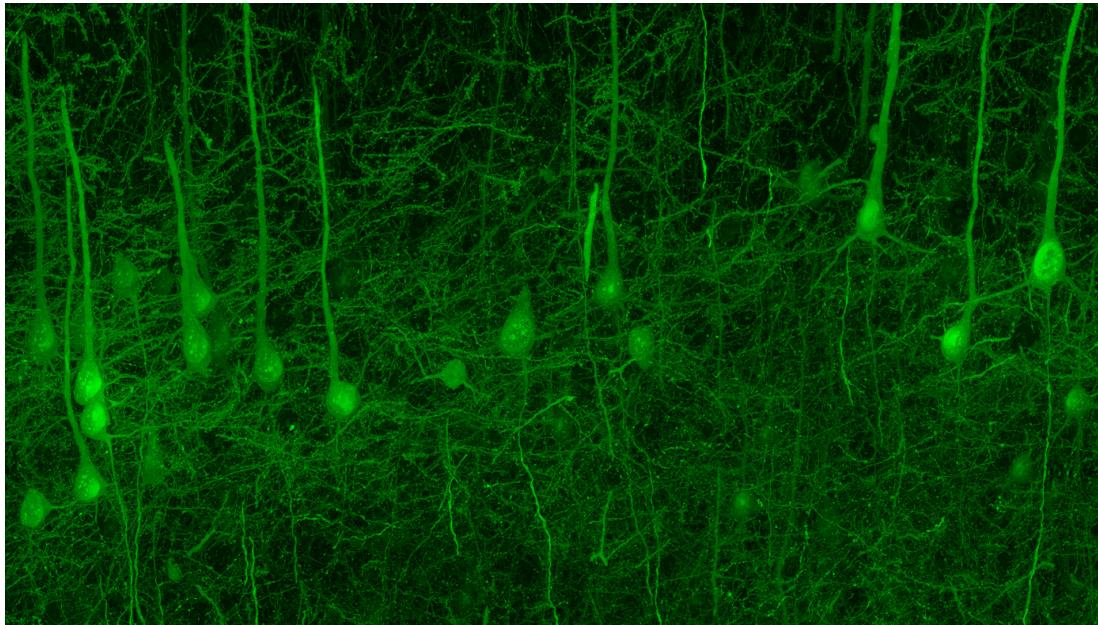
Osmium Tetroxide (OsO_4)



From <http://connectomes.org>. See Mikula et al. (2012). wbPATCO stain (OsO_4 variant)

- Stains lipid (all cell membranes): Dense, Unselective, Whole brains can be stained. Ideal for electron microscopy.

Immunofluorescence Labeling



http://smithlab.stanford.edu/Smithlab/Array_Tomography.html

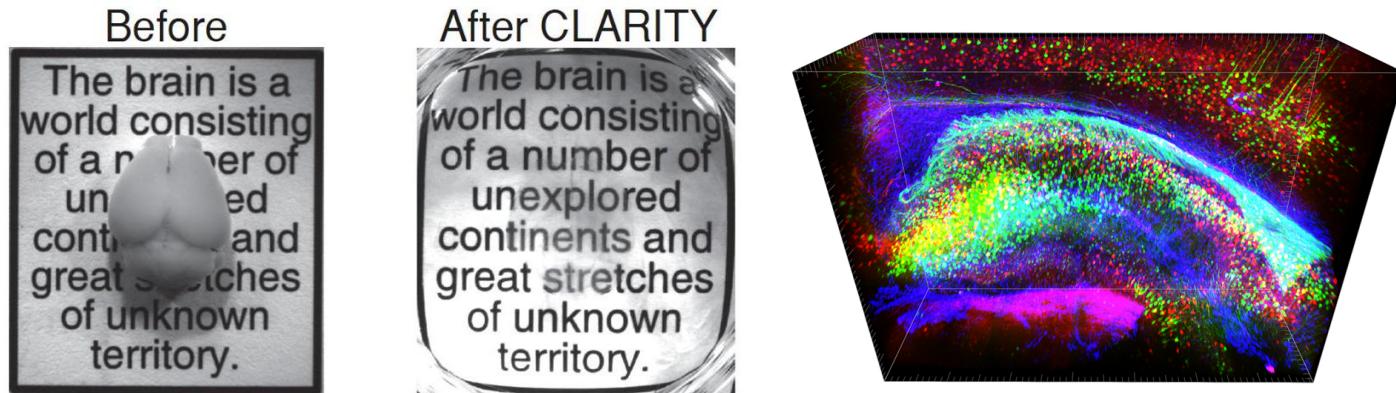
YFP expressed in whole neurons (false color added)

- Targets specific molecules (e.g. proteins): Sparse, Targetted
- Use antibody (to attach to antigen in the target) linked to fluorophore (directly or indirectly).

Tracer Injections

- Fills neurons near injection site (whole neurons):
Sparse (local to injection site), Unselective, Can span long distances.
- Anterograde (soma toward axon terminal),
Retrograde (axon terminal toward soma)
- Viral: anterograde or retrograde. Can cross synapses through infection to highlight higher-order connections (e.g., Pseudorabies virus)

Other Relevant Techniques



<http://directorsblog.nih.gov/the-brain-now-you-see-it-soon-you-wont/>

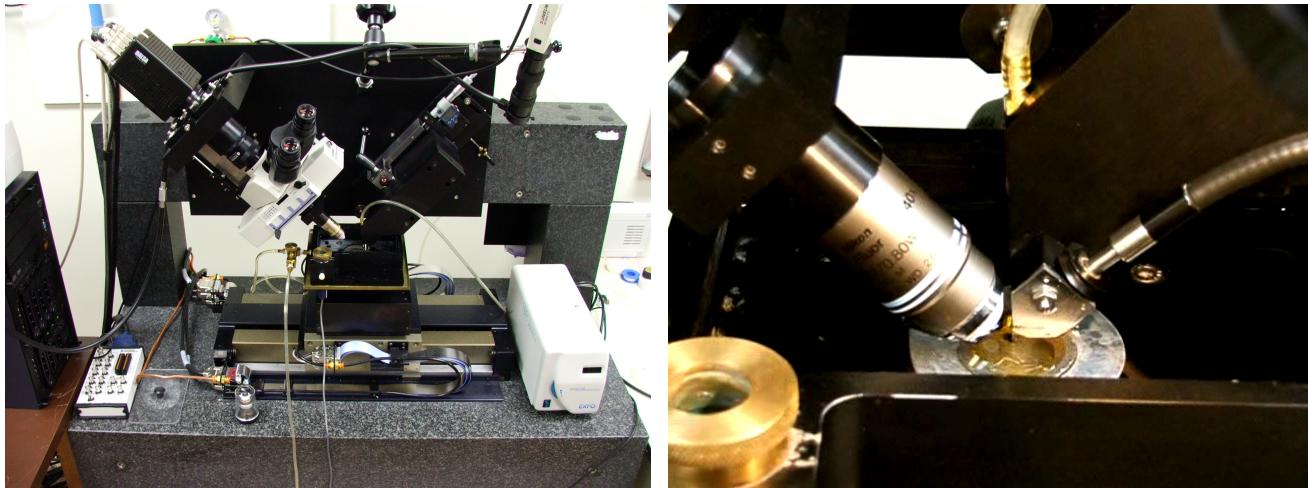
<http://clarityresourcecenter.org/> (Chung and Diesseroth 2013)

- Making brain tissue transparent: remove lipid, replacing with hydrogel for structural support.
- CLARITY: Allows imaging using multiple immunostains over large volumes of brain tissue.

Part II

Imaging

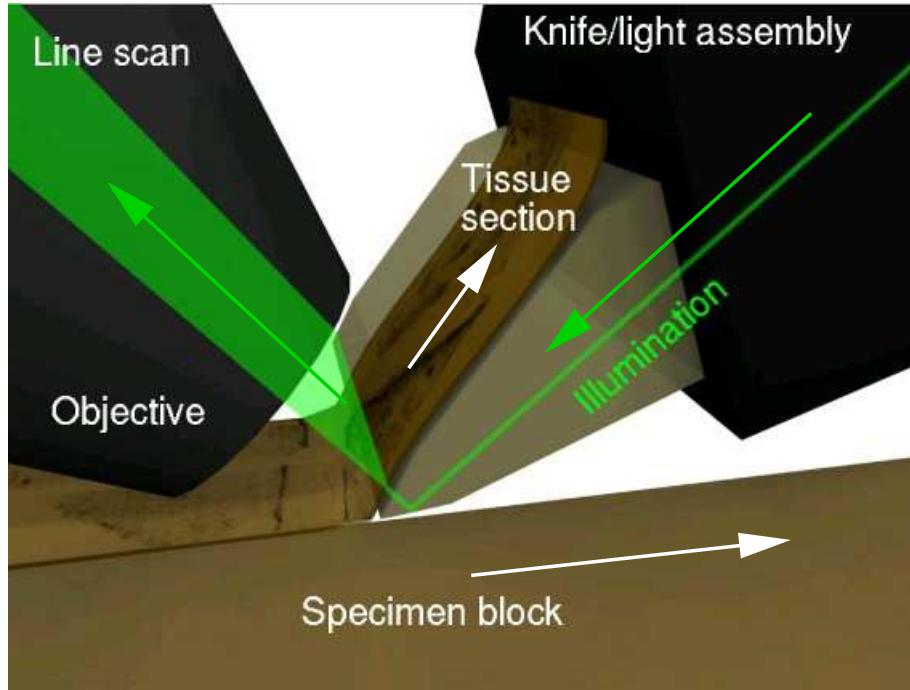
Knife-Edge Scanning Microscope



Mayerich et al. (2008); Chung et al. (2011)

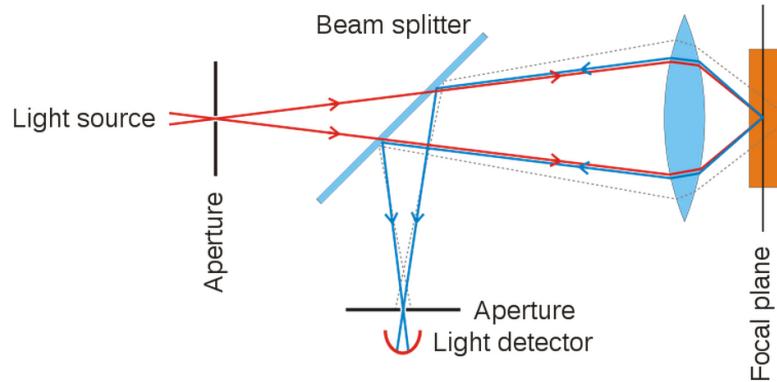
- Physical sectioning, as opposed to optical sectioning (e.g. confocal).
- Light microscopy, bright-field imaging (fluorescence in the works).
- Stains: Golgi (neuron morphology), Nissl (soma), India ink (vasculature). (Fluorescence imaging in the works.)
- $0.6 \mu\text{m} \times 0.7 \mu\text{m} \times 1 \mu\text{m}$ voxel resolution.
- Custom software for control, image capture (Kwon et al. 2008).
- Compare to MOST (based on KESM) (Li et al. 2010).

Imaging Principles of the KESM



- Image while cutting (line-scan at the tip of the knife).
- Transmission illumination through the diamond knife.
- Tissue thickness: 1 μm (or possibly less).

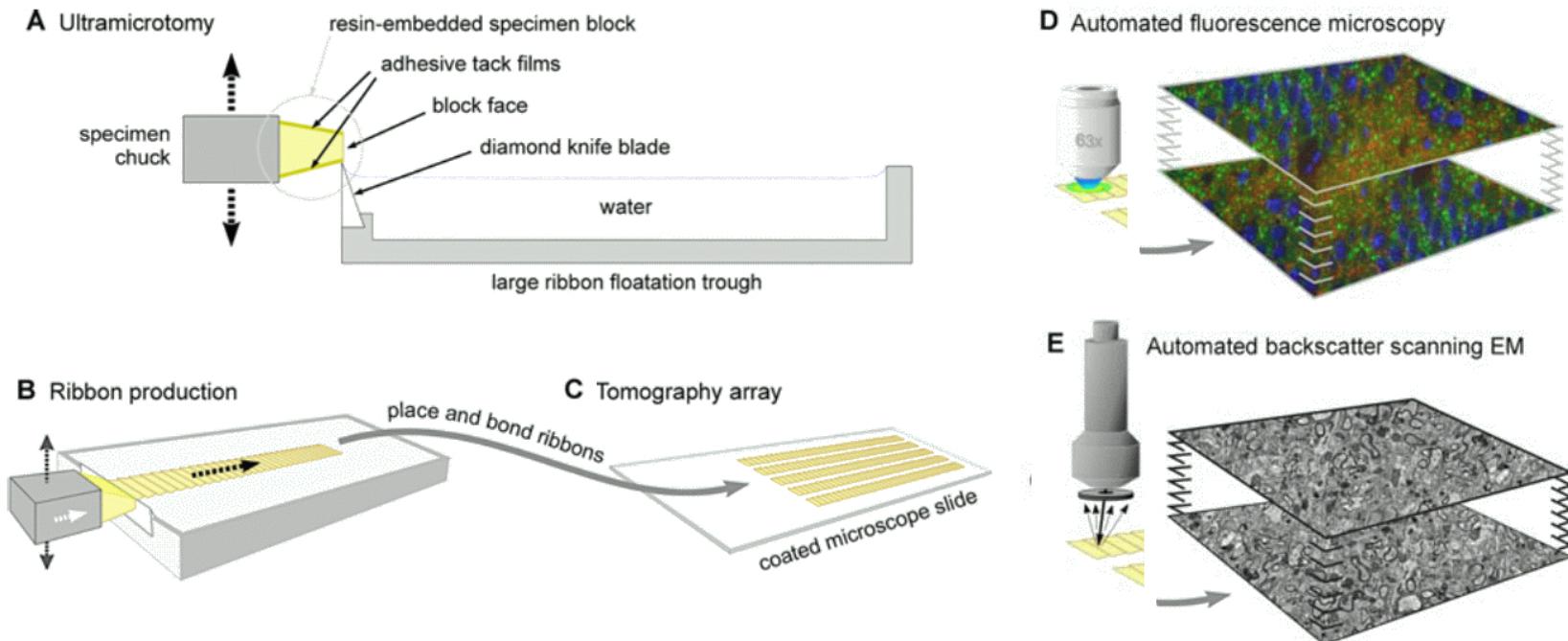
Confocal Microscopy



http://en.wikipedia.org/wiki/File:Confocalprinciple_in_English.svg

- Optical, not physical sectioning: Imaging at a specific focal depth. Scanning. Fluorescence imaging.
- Depth limit (max 1 mm) (Murray 2011).
- Also see two-photon (and multi-photon) imaging.

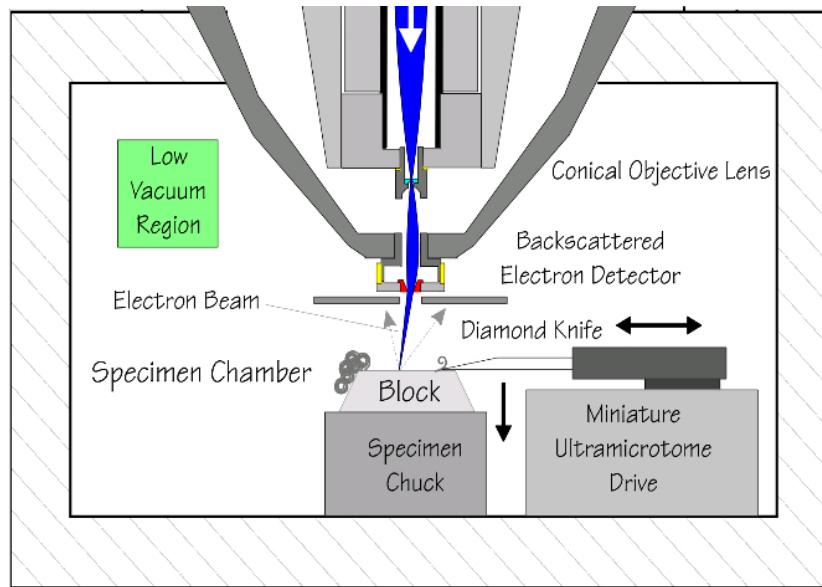
Array Tomography



Micheva and Smith (2007)

- Ultrathin sections transferred on glass slide.
- Repeated washing and staining allows perfectly registered volume data from multiple staining modalities.

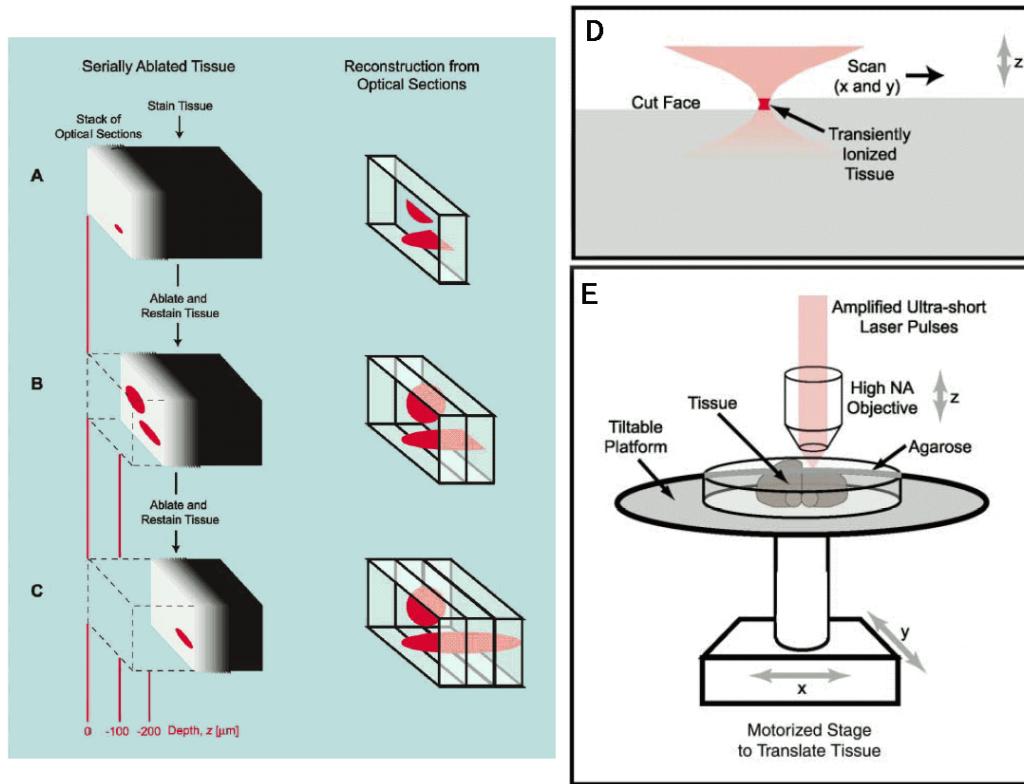
SBF-SEM (or SBEM)



Denk and Horstmann (2004)

- Microtome installed inside the vacuum chamber of an SEM.
- Commercially available from Gatan.

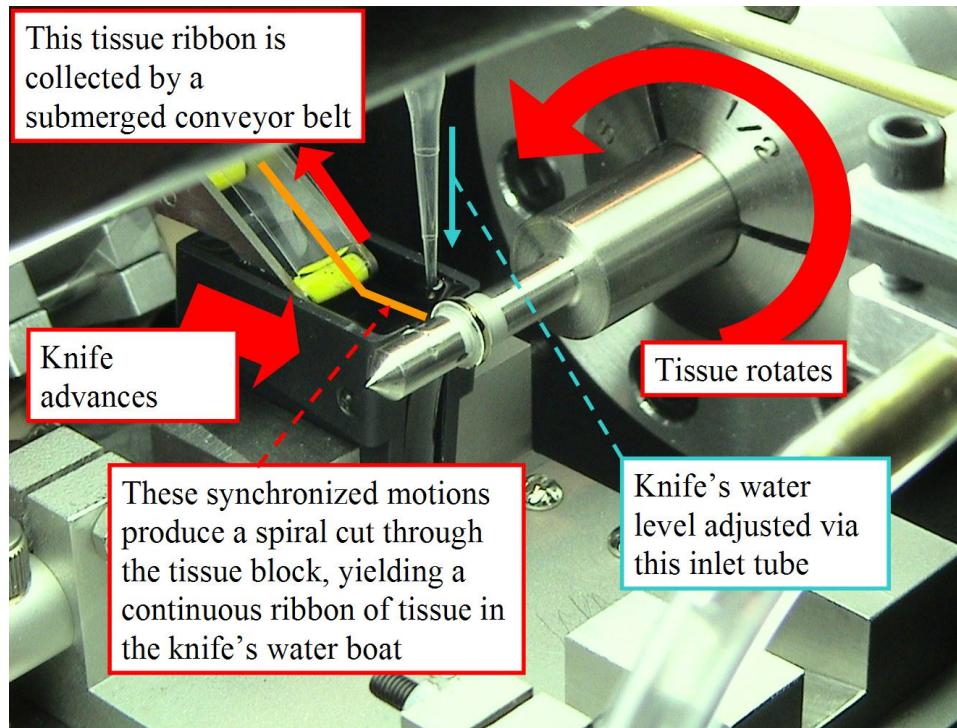
All-Optical Histology



Tsai et al. (2003)

- Hybrid of physical sectioning and optical sectioning (cf. Serial Two-Photon Tomography (Ragan et al. 2012)).
- Femtosecond laser pulses used to ablate $\sim 150 \mu\text{m}$ sections, followed by multiphoton imaging.

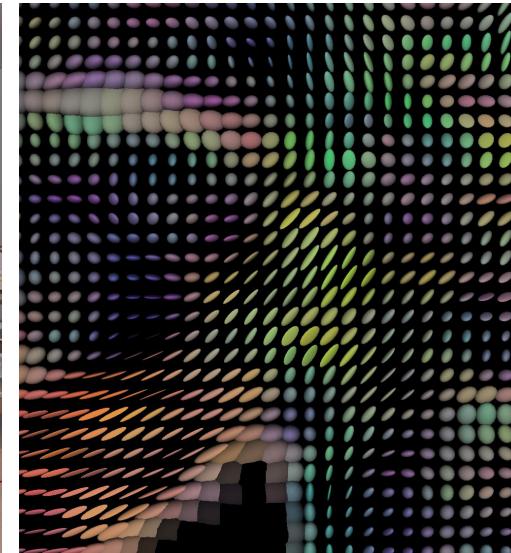
ATLUM



Hayworth et al. (2006)

- Continuous sectioning using a lathe.
- Sectioned tissue collected on adhesive tape.
- Post-staining and imaging of tape library with Transmission EM.

Diffusion Tensor Imaging (DTI)

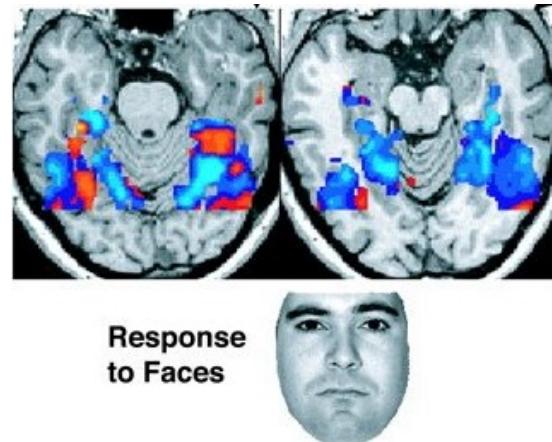


<http://en.wikipedia.org/wiki/File:MRI-Philips.JPG>

<http://en.wikipedia.org/wiki/File:DTI-axial-ellipsoids.jpg>

- Based on Magnetic Resonance Imaging (MRI). Low resolution ($\sim 100 \mu\text{m}$).
- Detect anisotropic diffusion patterns of water molecules along fiber tracts.

functional MRI (fMRI)



Haxby et al. (2001) (image cropped)

- Brain activity measured through BOLD (blood oxygen level dependent signal).
- Region-to-region connectivity can be inferred based on activity correlation or causality (dynamic causal model, Granger causal model): (Friston 2009).

Comparison

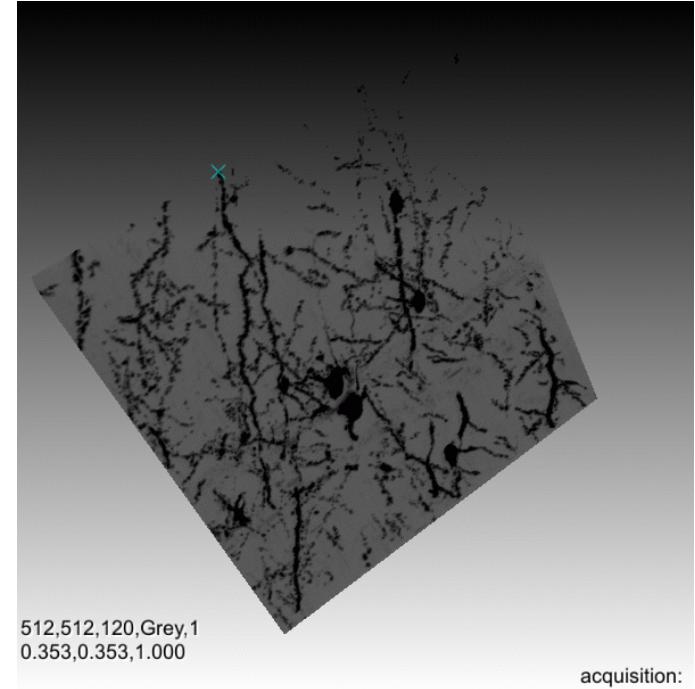
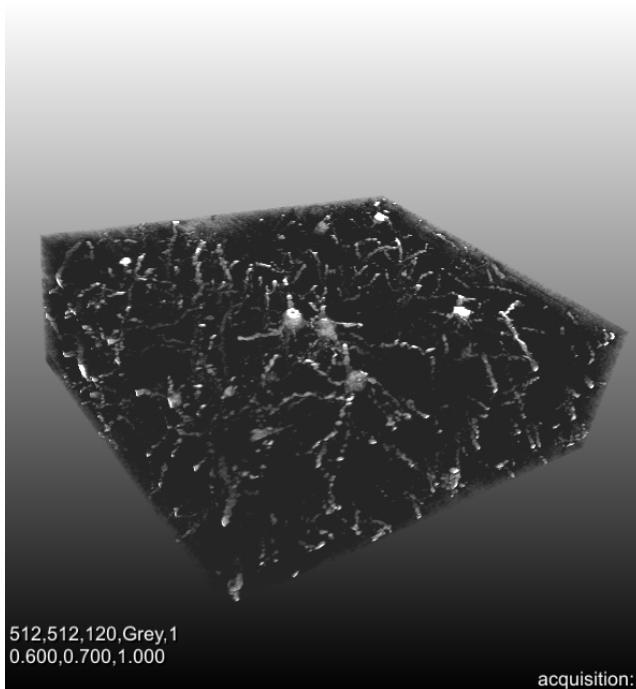
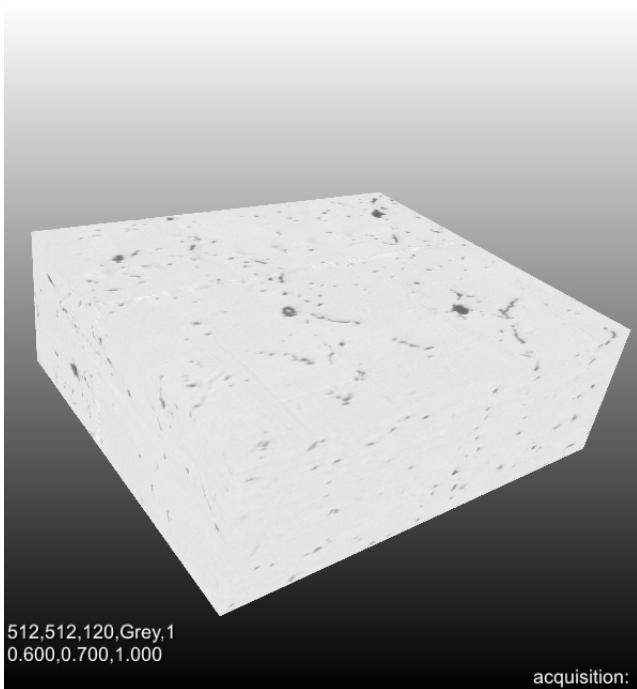
Table 1: Summary Comparison.

Method	nm-scale	μm-scale	High-Volume	High-Throughput
KESM (Mayerich et al. 2008) (cf. Li et al. 2010)	–	○	○	○
Confocal	–	○	–	–
All-Optical Hist. (Tsai et al. 2003)	–	○	○	–
Serial Two-Photon Tomography (Ragan et al. 2012)	–	○	○	–
Array Tomography (Micheva and Smith 2007)	○	○	–	–
SBF-SEM (Denk and Horstmann 2004)	○	–	–	–
ATLUM (Hayworth et al. 2006)	○	–	○	–
MRI/diffusion MRI (Jacobs et al. 1999; Hagmann et al. 2007)	–	–	○	○
nm-scale: ~ 10 nm (thickness of cell membrane)				
μm-scale: ~ 1 μm (diameter of dendrites, axons, capillaries, etc.)				
High-Volume: > 1 cm ³ (approximate volume of mouse brain and other organs)				
High-Throughput: < 100 hours (for ~ 50 scanned organs per year)				

Part III

Data and Online Resources (with Demo)

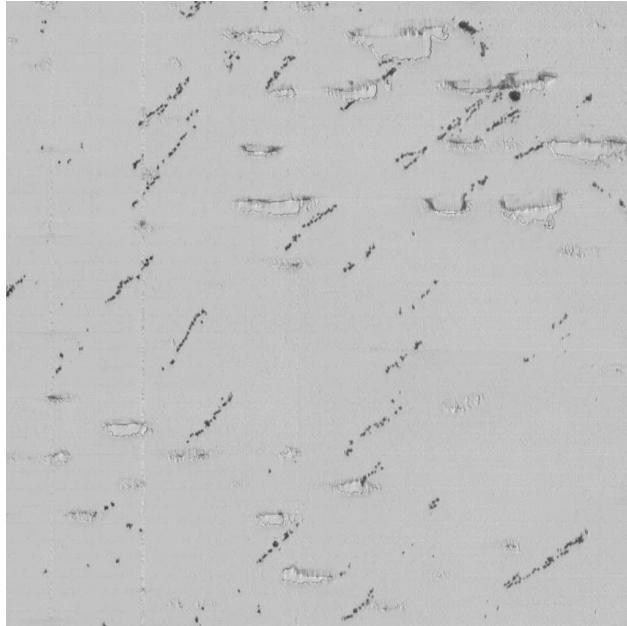
KESM Data



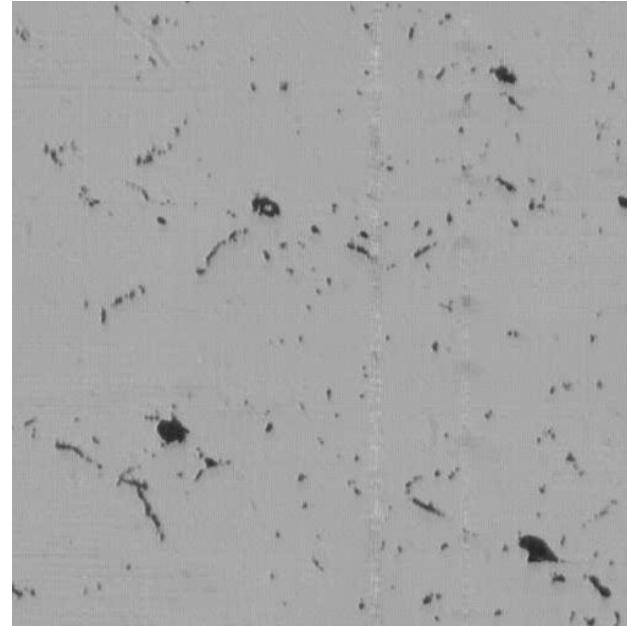
300 μm \times 350 μm \times 120 μm block

- Basically a huge 3D stack made up of 2D images.
- Details such as dendritic spines can be observed.

KESM Data (Image Stack)



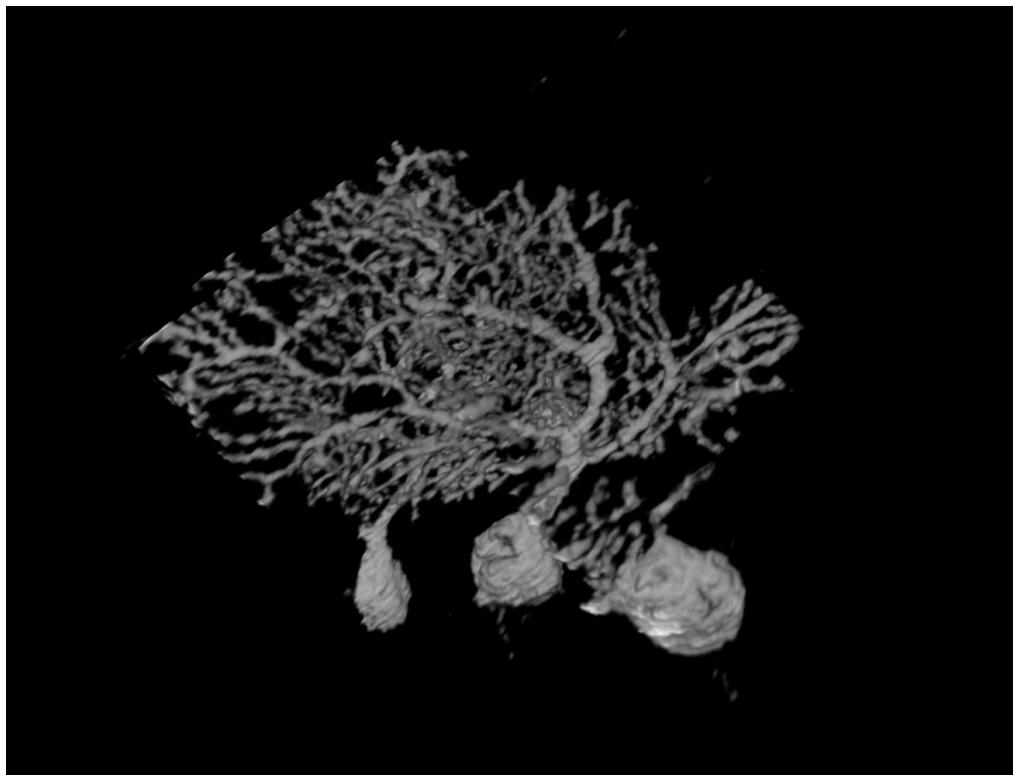
Cerebellum (Golgi)



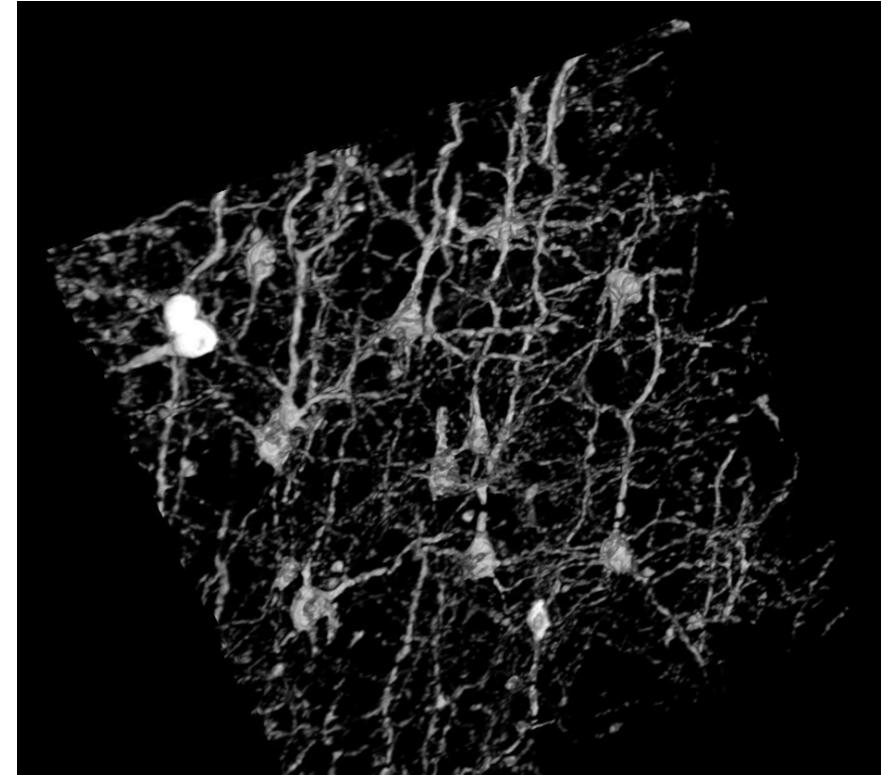
Cortex (Golgi)

- Flythrough of 3D stack: Looks like a movie in 2D.
- Each frame = 1 μm -thin section.

KESM: Volume Visualization



Golgi (Cerebelum)

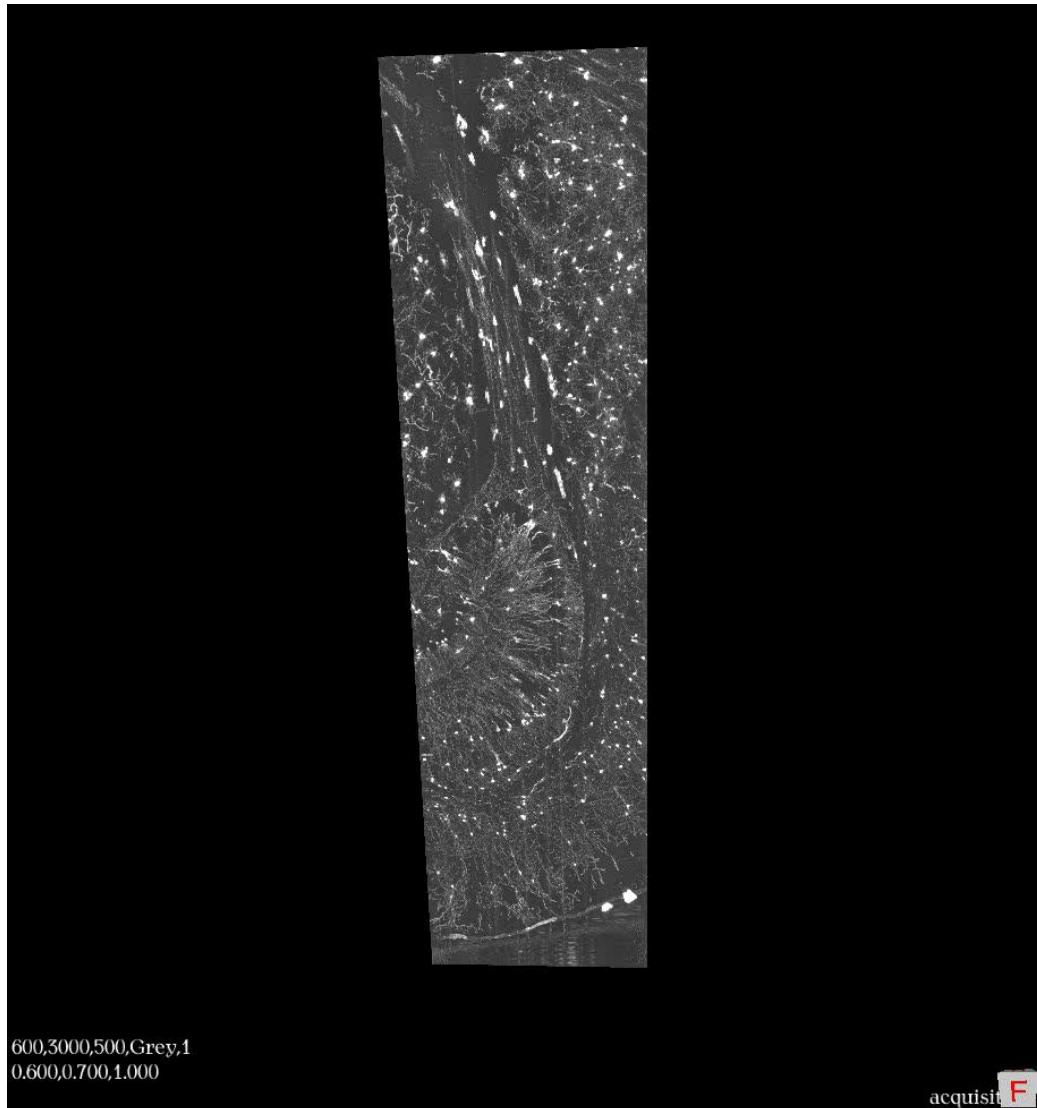


Golgi (Cortex)

3D visualization of

- Purkinje cells and pyramidal cells.

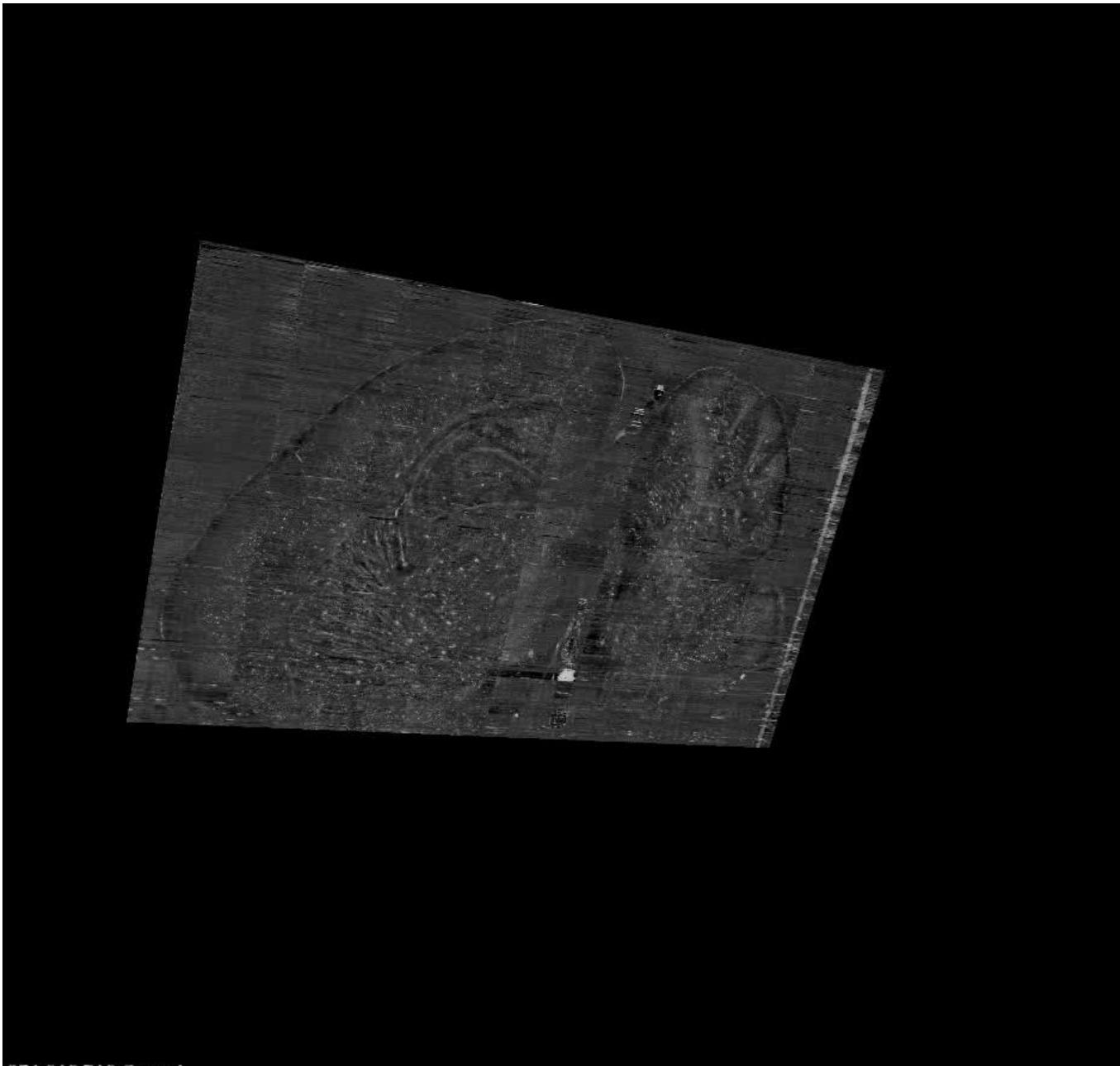
KESM: Local Circuits (Hippocampus)



600,3000,500,Grey,1
0.600,0.700,1.000

acquisit F

KESM Whole Brain: Neurons (Golgi)



KESM Brain Atlas

New Features

Customization

GoogleMaps™
Javascript API V2

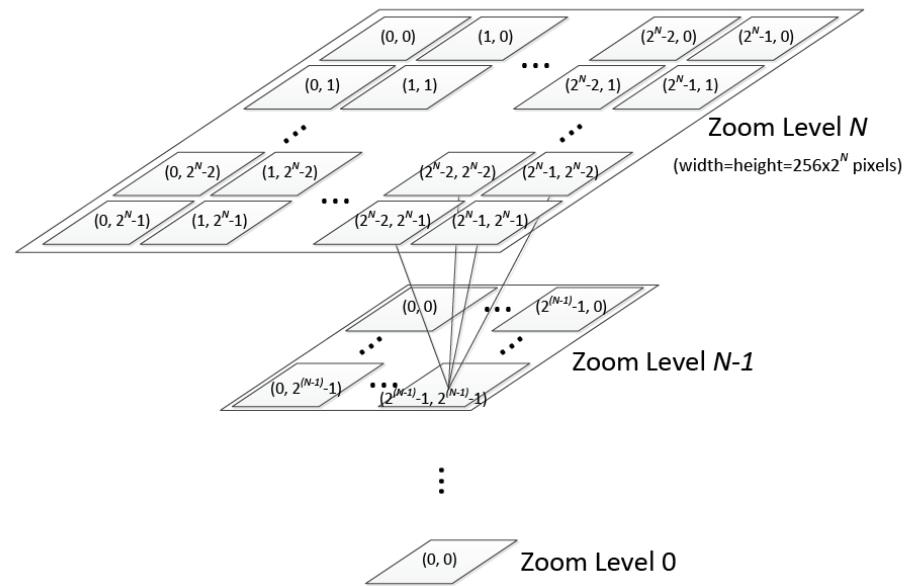
- Custom Tile
- Custom Overlay
 - Overlay Number
 - Overlay Interval
 - Zoomable Annotation
- Redraw

- Information Panel
- Scale Bar
- Map Capture
- Z-axis Navigation

API layers

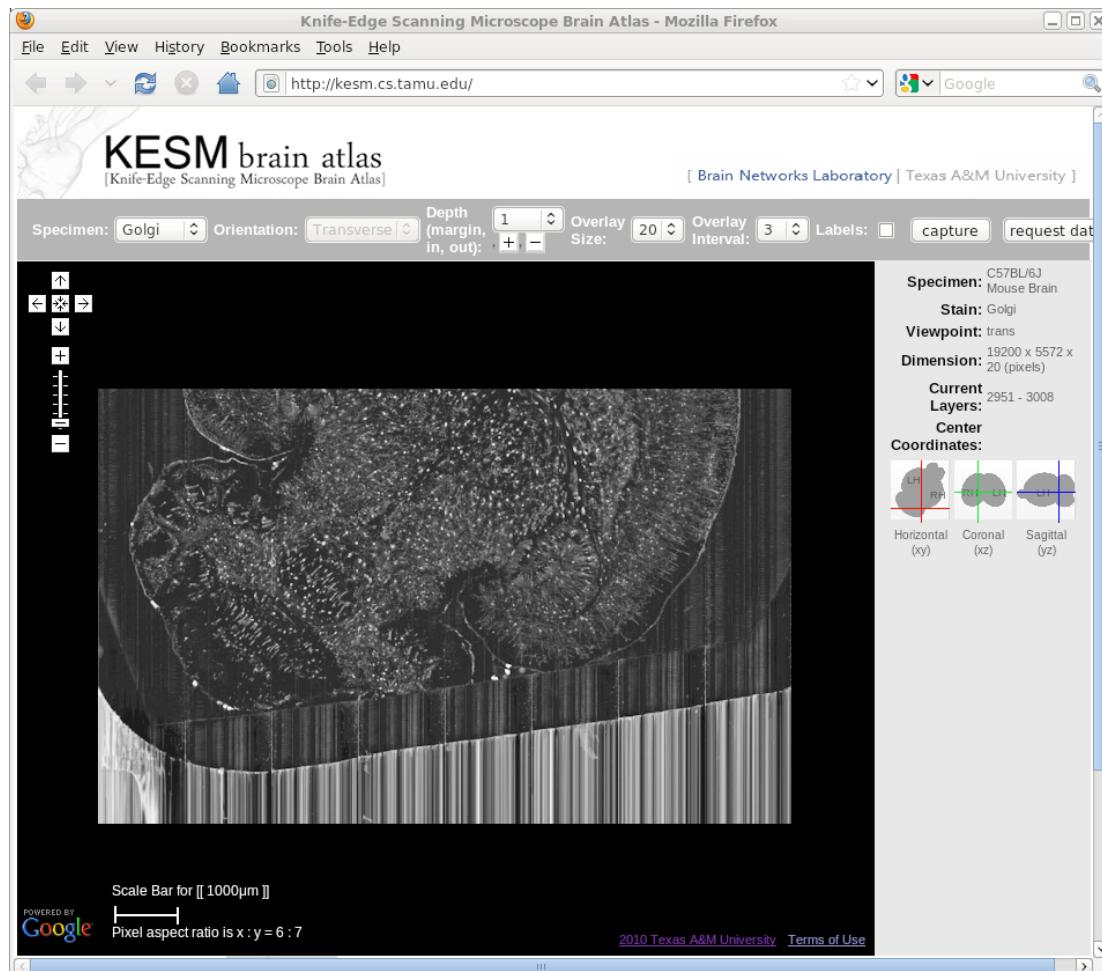
- Multi-scale tiles.
- Semi-transparent images.
- Google Maps API (v2).

→ KESM Brain Atlas



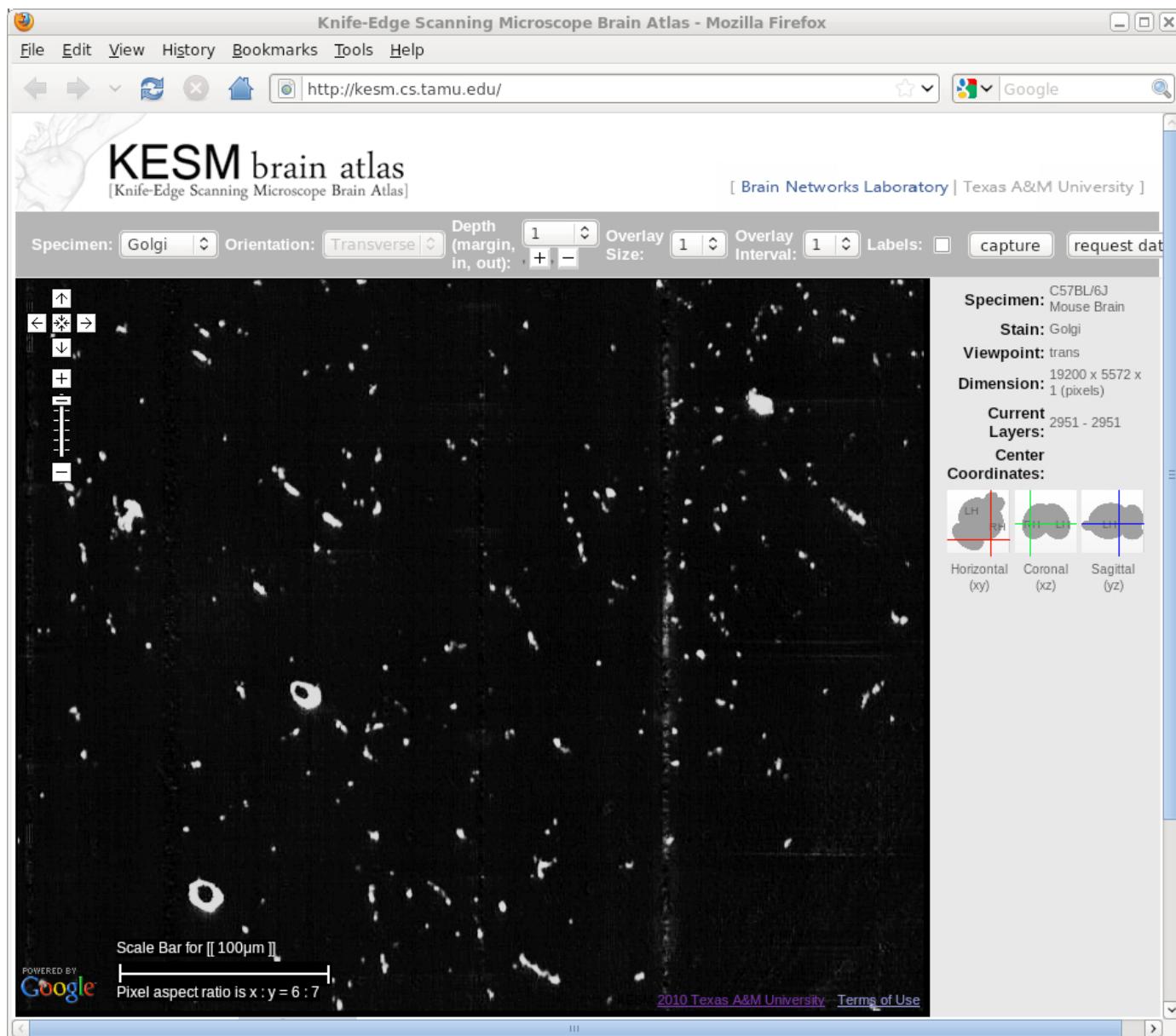
Tiling Scheme

KESM Brain Atlas (KESMBA)

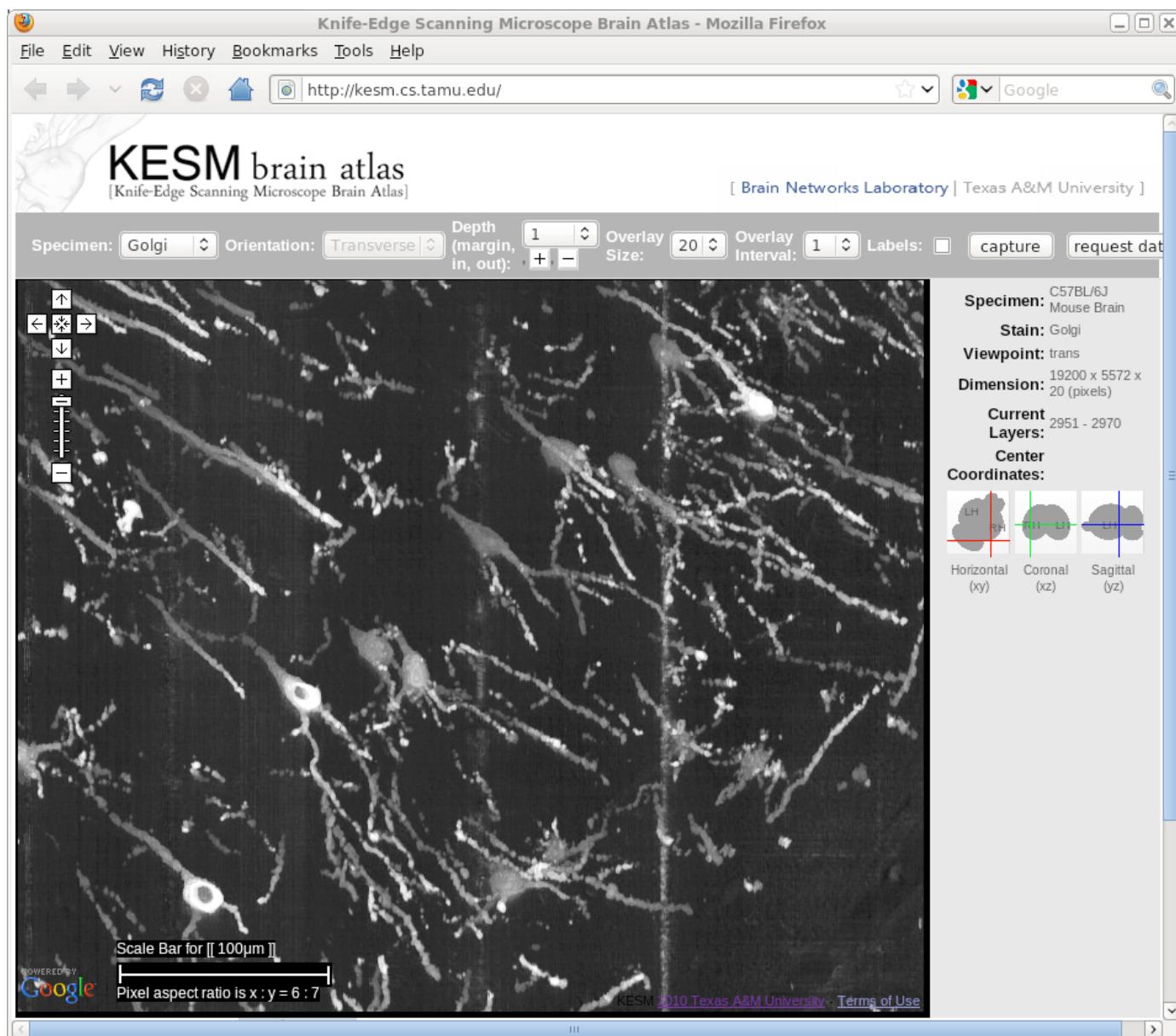


- <http://kesm.org> (Chung et al. 2011).
- Open to all! Even runs on smartphone browsers (can be slow).

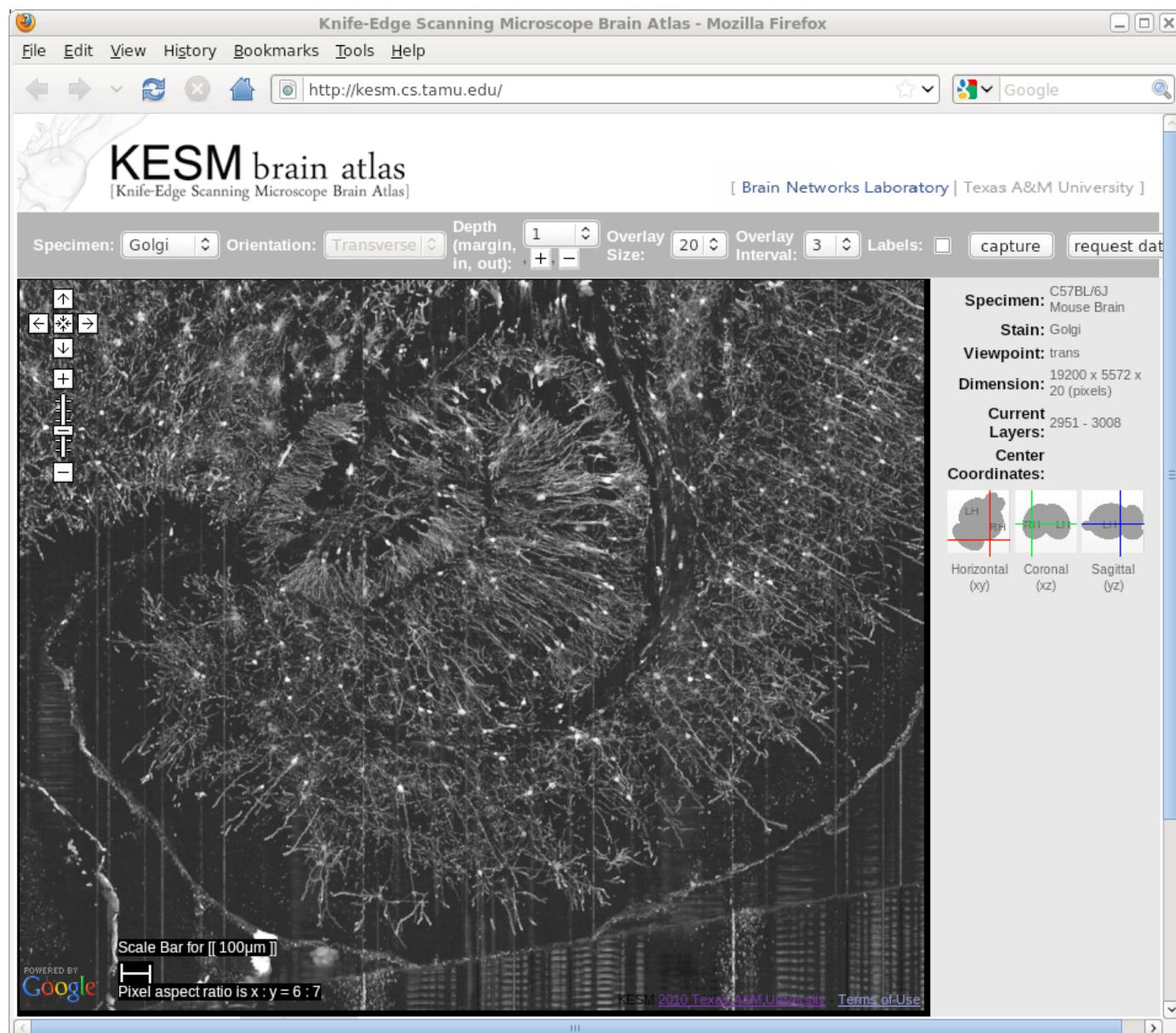
KESMBA: Single Overlay



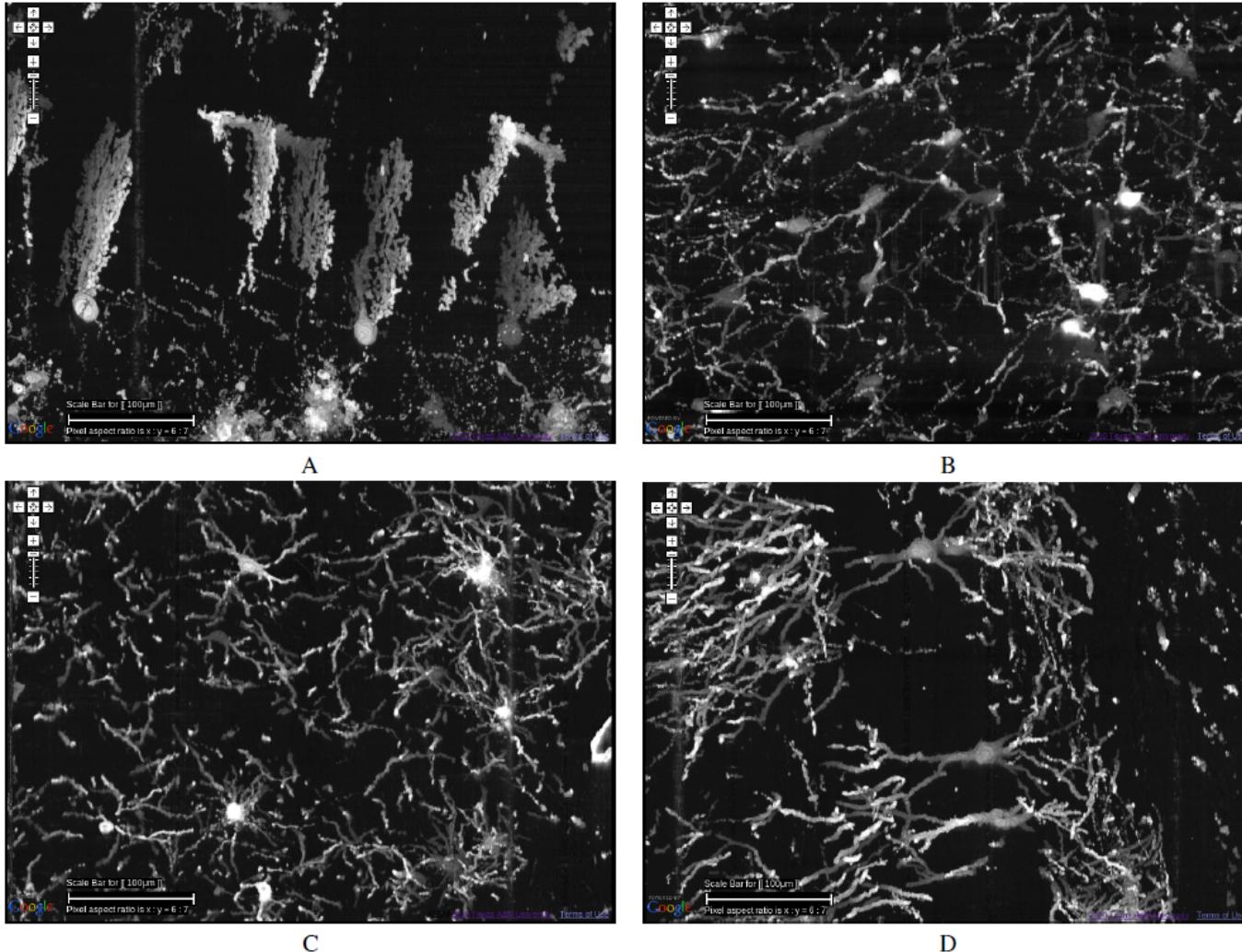
KESMBA: 20 Overlays



KESMBA: Zoomed Out

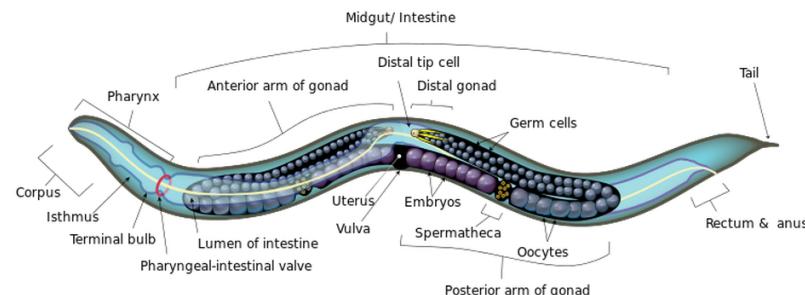


KESMBA: Some Samples



A: Cerebellum, B: Inferior colliculus, C: Thalamus, D: Hippocampus

OpenWorm Project



Enter the worm.

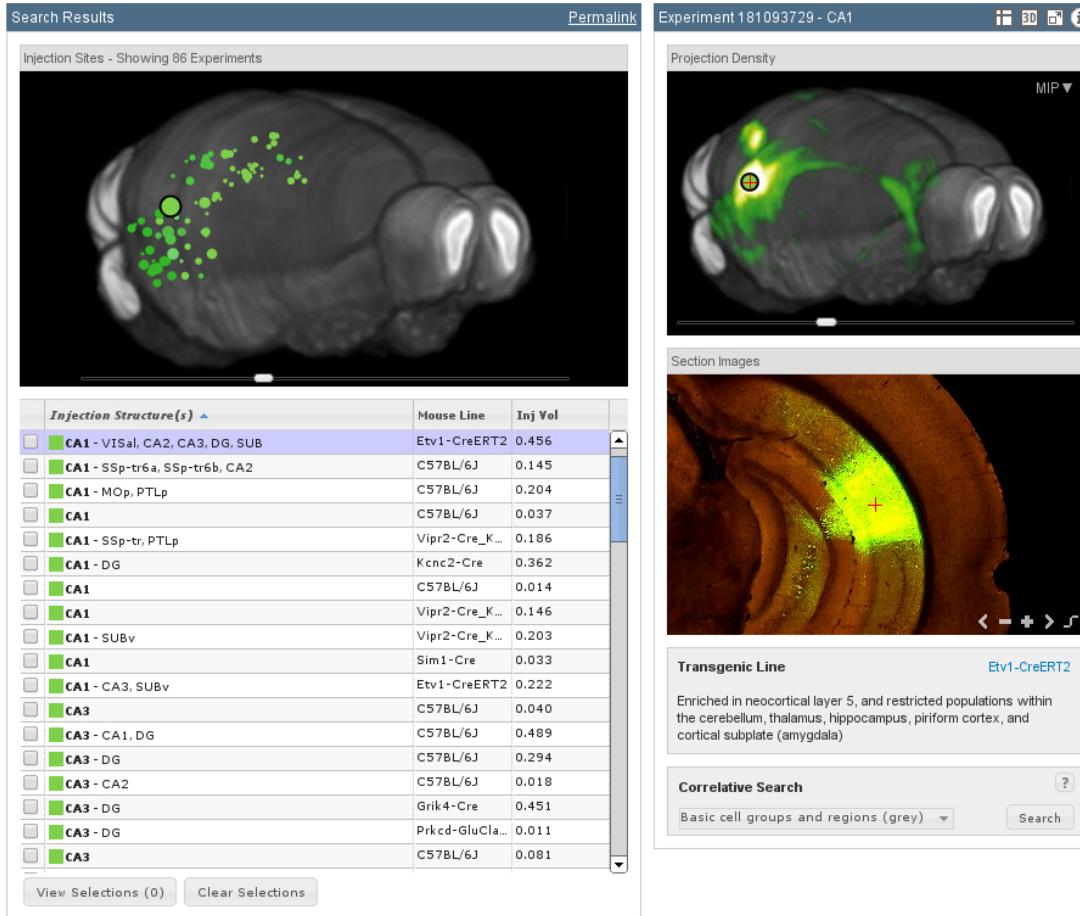


http://en.wikipedia.org/wiki/File:Caenorhabditis_elegans_hermaphrodite_adult-en.svg

<http://www.openworm.org/>

- **C. elegans connectome, downloadable in XML (NeuroML), for multicompartment models.**
- **Ultimate goal of constructing a detailed simulation of the whole worm.**

Allen Brain Atlas: Mouse Connectivity



<http://connectivity.brain-map.org>

- Tracer injection-based (1010 injection sites).
- Fluorescence microscopy.

Mouse Connectome Project (UCLA)

The image shows two screenshots of the Mouse Connectome Project website. The left screenshot is a list of 20 cases, each with details like Case ID, Plane, Coordinates, Injection Site Center, Description, Organism, and Tracers. The right screenshot is a detailed visualization of a coronal brain slice showing tracer pathways in green.

Case	Plane	Coordinate (x,y,z)	Injection Site Center	Description	Organism	Tracers
SWI20525-04A	CORONAL	(0.5, -2.7, 1.1) (0.0, -4.2, 2.3)	RSPv DR	Retrosplenial area, ventral part Dorsal nucleus raphe	mus musculus C57Bl/6J wild	fg/bda ctb/phal
SWI20525-02A	CORONAL	(0.4, -0.1, 0.8) (2.0, -0.1, 0.9)	ACAd SSp-II	Anterior cingulate area, dorsal part Primary somatosensory area, lower limb	mus musculus C57Bl/6J wild	fg/bda ctb/phal
SWI20525-01A	CORONAL	(3.62, -2.1, 0.7) (1.25, 2.8, 1.8)	SSp ORBI	Primary somatosensory area Orbital area, lateral part	mus musculus C57Bl/6J wild	fg/bda ctb/phal
SWI20404-04A	CORONAL	(0.37, 2.2, 1.25) (0.37, 2.2, 2.17)	PL ILA	Prelimbic area Infralimbic area	mus musculus C57Bl/6J wild	fg/bda fg/bda
SWI20404-03A	CORONAL	(0.37, 2.2, 1.25) (0.37, 2.2, 2.17)	PL	Prelimbic area Infralimbic area	mus musculus C57Bl/6J wild	bda phal
SWI20404-02A	CORONAL	(0.37, 2.2, 1.25) (0.37, 2.2, 2.17)	PL	Prelimbic area Infralimbic area	mus musculus C57Bl/6J wild	fg/bda ctb/phal
SWI20404-01A	CORONAL	(0.37, 2.2, 1.25)	PL	Prelimbic area	mus musculus C57Bl/6J wild	fg

<http://www.mouseconnectome.org/> (Hintiryan et al. 2012)

- Tracer injection-based (245 injection sites).
- Fluorescence microscopy.

Brain Architecture Project (CSHL)

The image shows two screenshots of the Brain Architecture Project website. The left screenshot is a list of injection sites, and the right is a 3D brain visualization.

Left Screenshot (List of Injection Sites):

Series Name	Viewer	Inj Region	Inj Coords (X, Y, Z)	Tracer	Preview
MouseBrain_1305 F		Frontal pole, cerebral cortex	x:1.00 y:3.24 z:1.50	AAV	Section Gallery
MouseBrain_1145 F		Somatomotor areas	x:1.20 y:2.84 z:2.30	RV	Section Gallery
MouseBrain_0958 F		Secondary motor area	x:1.25 y:2.84 z:1.52	AAV	Section Gallery

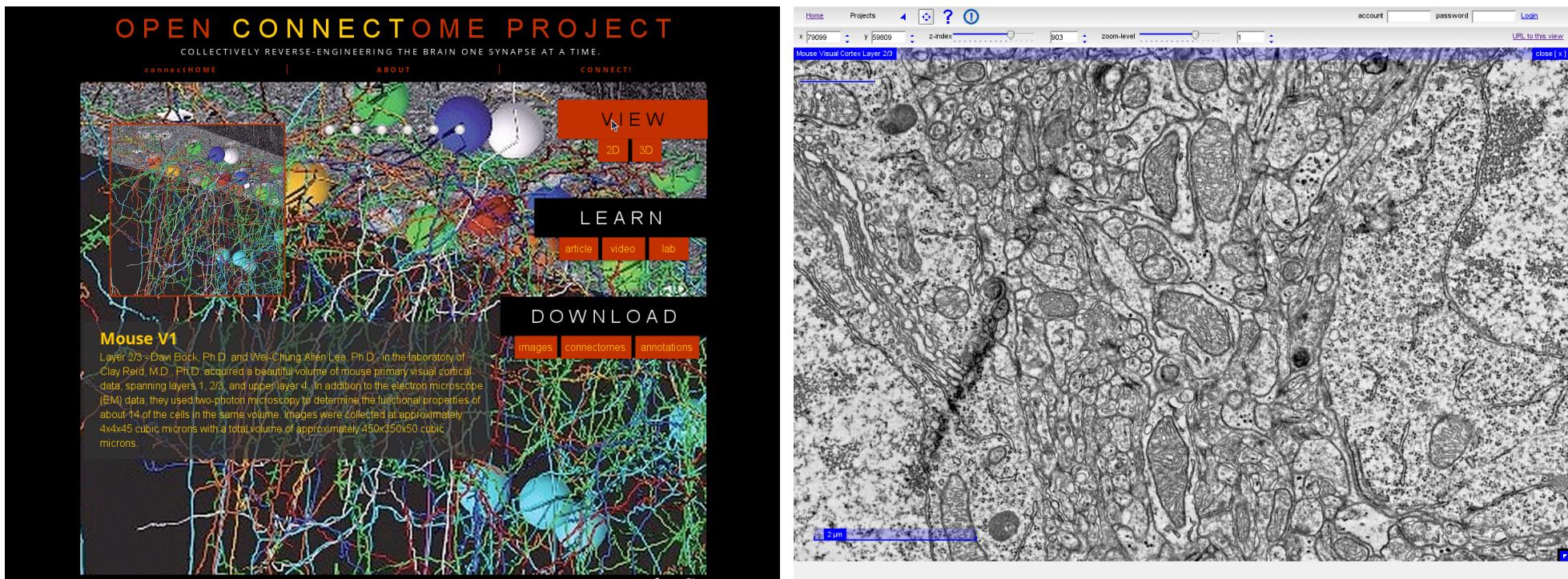
Right Screenshot (3D Brain Visualization):

A 3D rendering of a mouse brain showing injection sites. A red fluorescent signal is visible in the frontal pole and other regions. The interface includes a navigation menu on the left and a series of small brain section images at the bottom.

<http://brainarchitecture.org> (Mitra 2012)

- Tracer injection-based (235 injection sites, mouse).
- Fluorescence microscopy. Other species also available.

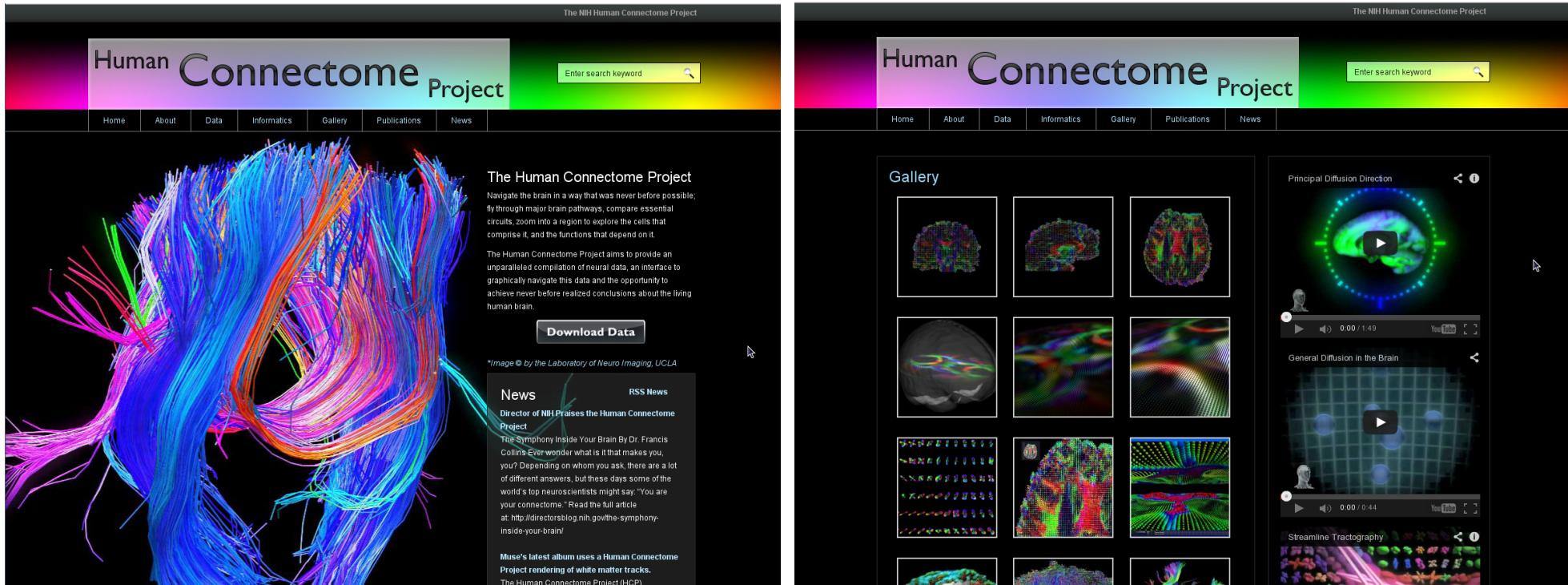
Open Connectome Project



<http://www.openconnectomeproject.org>

- EM data from mouse visual cortex (Bock et al. 2011).

Human Connectome Project

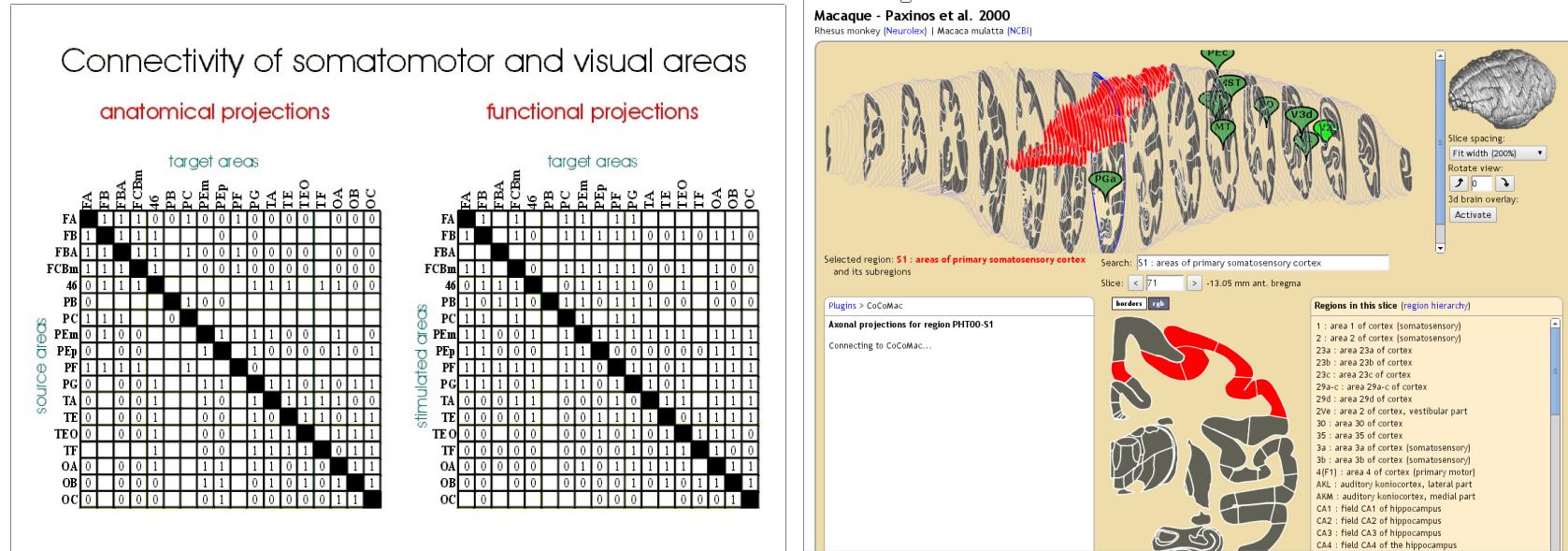


<http://www.humanconnectomeproject.org>

- DTI data from human (Van Essen et al. 2012).
- Also see (Hagmann et al. 2007).

CoCoMac

Permalink



<http://cocomac.org> <http://scalablebrainatlas.incf.org>

- Macaque brain connectivity (based on 2508 tracer injections, 39,748 connection details, collected from the literature).
- Second version under preparation:
<http://cocomac.g-node.org/>

UCLA Multimodal Connectivity DB

The UCLA Multimodal Connectivity Database (UMCD) is a web-based repository and analysis site for connectivity matrices that have been derived from de-identified neuroimaging data. Users can analyze connectivity matrices that have been shared publicly and upload their own matrices to share or analyze privately.

Analyze a Brain Network

To analyze any shared brain network, choose a study name, a network name from that study, a weighting scheme, and the percentage of edges to include, then click Analyze. Momentarily, you'll get a full breakdown of the chosen network, including study info and network analysis.

View/Download	Analyze	Study Name	Network Name	Atlas	Imaging Modality
View/Download	Analyze	ICBM	UCLA_ICBM_1004_DTI	Freesurfer 85	DTI
View/Download	Analyze	ICBM	ICBM_thickness_cormat	Freesurfer_68	Structural MRI
View/Download	Analyze	ICBM	ICBM_DTI_tractography	Freesurfer_68	DTI
View/Download	Analyze	ADHD200_CC200	KKI_2371032	Craddock 200	fMRI
View/Download	Analyze	ADHD200_CC200	KKI_2026113	Craddock 200	fMRI
View/Download	Analyze	ADHD200_CC200	KKI_3434578	Craddock 200	fMRI
View/Download	Analyze	ADHD200_CC200	KKI_8628223	Craddock 200	fMRI
View/Download	Analyze	ADHD200_CC200	KKI_1623716	Craddock 200	fMRI
View/Download	Analyze	ADHD200_CC200	KKI_1594156	Craddock 200	fMRI

<http://umcd.humanconnectomeproject.org>

- MRI-based connectivity database.

Part IV

Analysis

Geometric Reconstruction

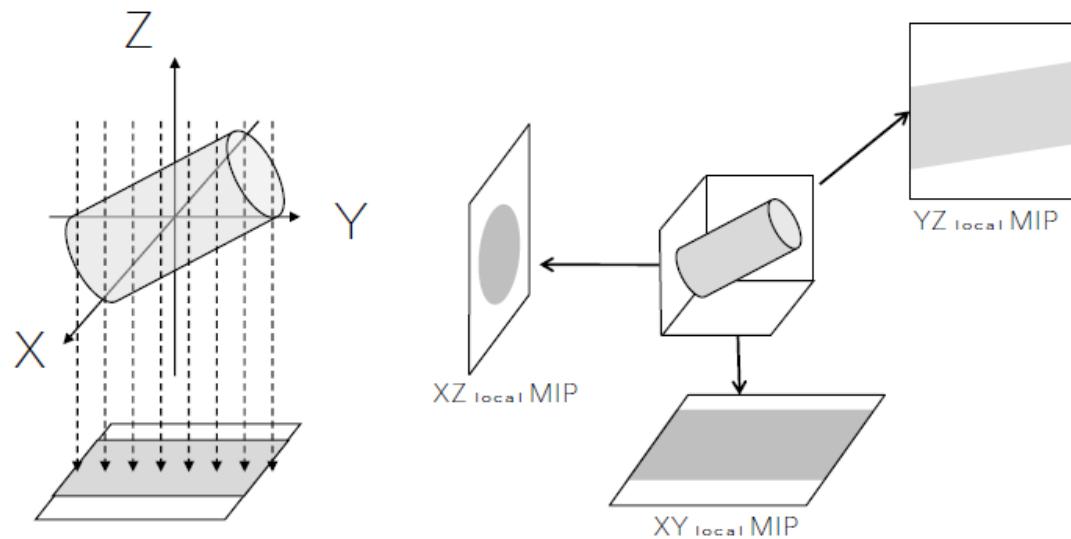
Raw data or volume visualization is not enough:

- We need to reconstruct the geometric structure of the objects in the data.
- Data can be huge (several TB): manual tracing is not an option.
- We need automated algorithms.

Reconstruction Approaches

- Segment-then-connect: the most common approach
- 3D convolutional network: Jain et al. (2010)
- Template-matching-based vector tracing: Al-Kofahi et al. (2002); Han et al. (2009b,a); Han (2009); Luisi et al. (2011)
- Semi-automated reconstruction: Yang and Choe (2011b)
- Topology-constrained reconstruction: Yang and Choe (2011a); Jain et al. (2007)
- Crowd sourcing: Eyewire.org (Seung and Burnes 2012).

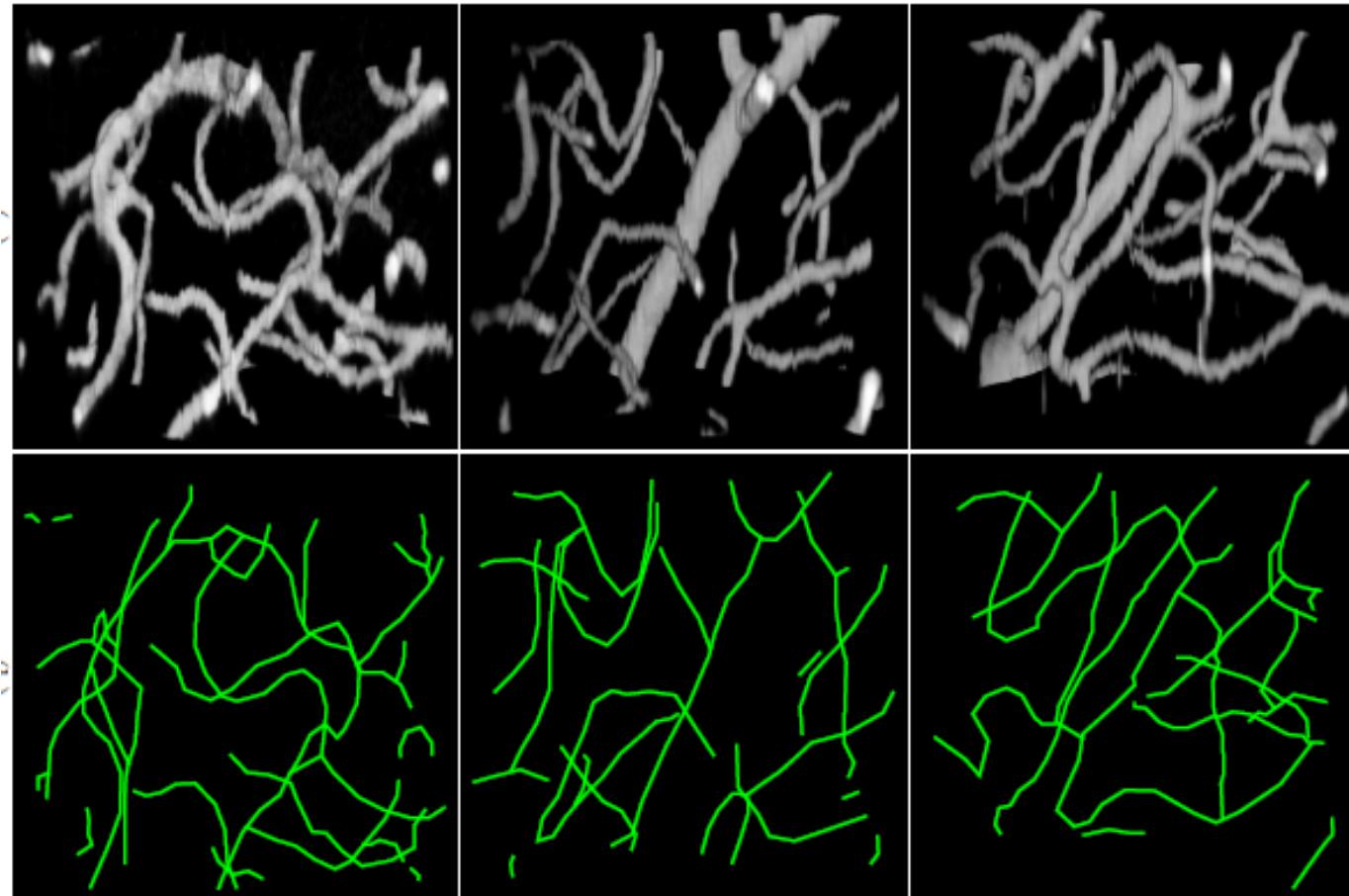
Tracing Example: MIP-Based Tracing



Han (2009)

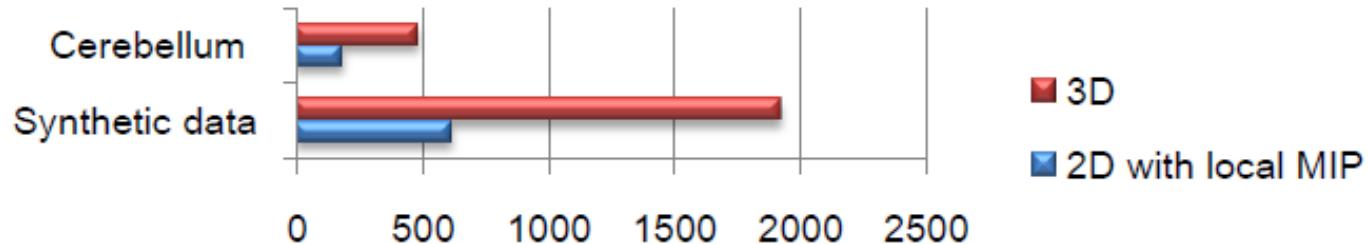
- Maximum-Intensity Projection (MIP).
- MIP-based tracing: Trace on projected 2D images.

MIP-Based Tracing Results



- KESM mouse vasculature data.

MIP-Based Tracing Performance



- MIP-based approach about $3\times$ faster than 3D version.

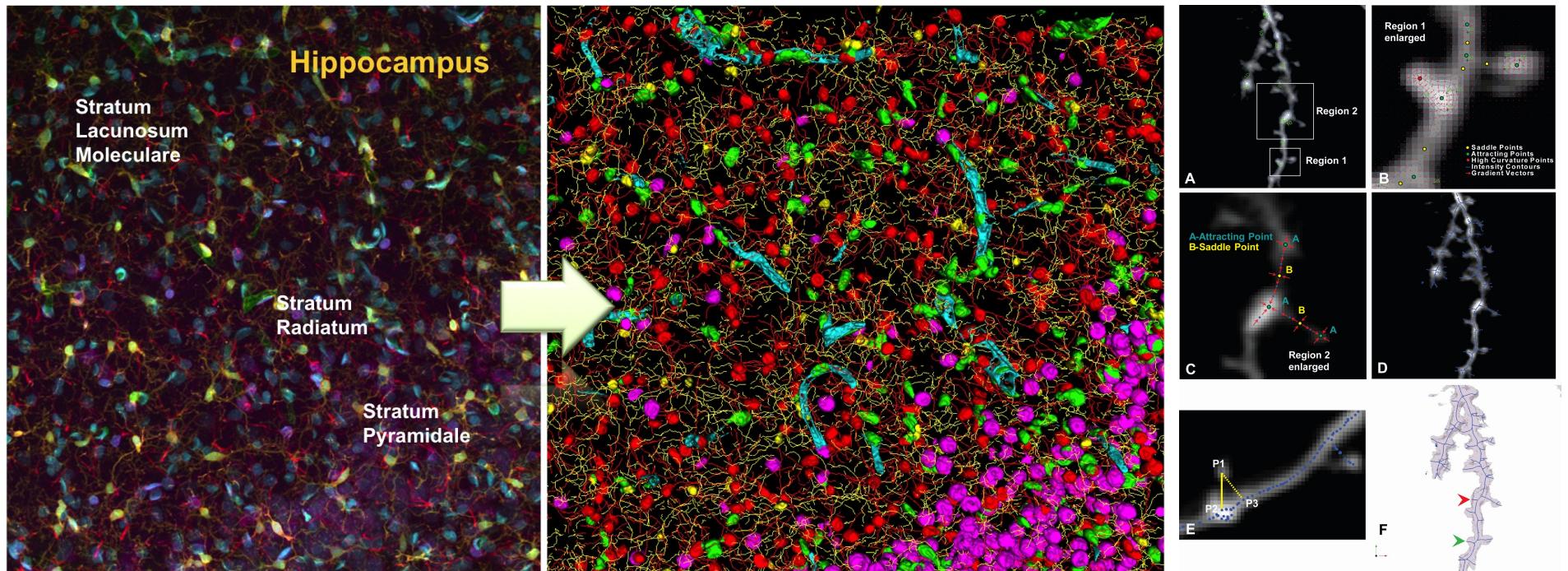
MIP-Based Tracing: Validation

	ϕ			φ		
	μ	σ	<i>p</i> -value	μ	σ	<i>p</i> -value
R1	0.1518	0.1762	0.4188	1.2131	0.3529	0.6853
R2	0.1325	0.1804		1.1294	0.3016	

Error

- Validation against small manual ground-truth (R1 and R2).
- ϕ = centerline deviation, φ = length difference.

FARSight Toolkit (U Houston)



<http://www.farsight-toolkit.org> (Luisi et al. 2011)

- 2D and 3D image analysis toolkit.

Other Reconstruction Tools

- KNOSSOS: 3D image data (mostly for EM) analysis tool (mostly manual). <http://www.knossostool.org/>
- EyeWire: crowd-sourcing EM reconstruction portal.
<http://eyewire.org/>
- Reconstruct: EM reconstruction tool (manual).
<http://synapses.clm.utexas.edu/tools/reconstruct/reconstruct.stm>
- Generic (yet powerful) tools:
 - ImageJ: <http://rsbweb.nih.gov/ij/>
 - ITK: <http://www.itk.org/>

Connectivity Analysis

- Graph-theory based analysis (Sporns 2002, 2011)
 - In-degree, out-degree, cluster index, power law
- Motif analysis (Milo et al. 2002).
 - Statistics of small sub-graph patterns.
- Dynamics (Thiel et al. 2003; Sporns and Tononi 2002)
- Large-scale simulation based on DTI (Izhikevich and Edelman 2008)
- Time is a crucial factor in connectivity analysis (Choe 2004).

Part V

Theoretical Insights

Wrap Up

Thinking Beyond Connectomics

Connections alone not enough:

- Sign: excitatory/inhibitory
- Weight: synaptic strength
- Delay: both conduction delay and integration time
- Molecular dynamics and gene expression
- Plasticity

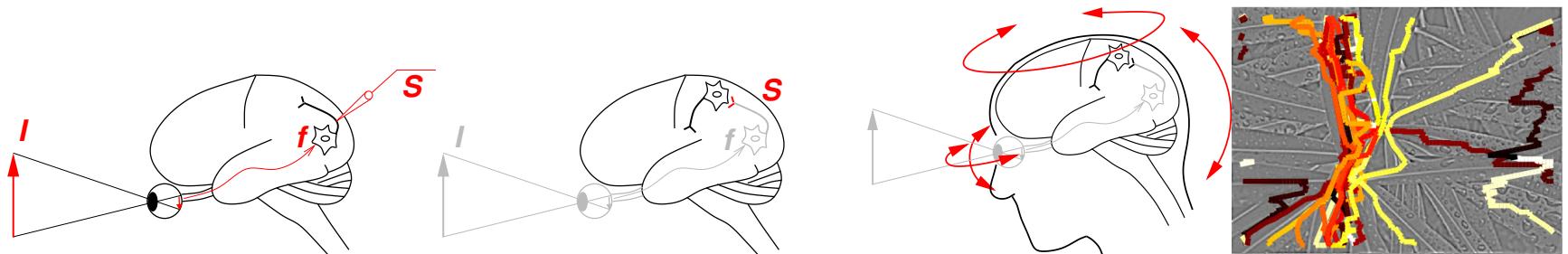
Is the Brain Enough? – Will Need the Body

- Brain is part of the body and a lot of function is performed by the spinal cord and the peripheral nervous system.
- To fully understand brain function, it must be understood in the context of the entire body.
- Imaging whole organisms may be necessary for a true understanding of brain function.

Risk of Doubling our Task?

- Without a proper theoretical framework for analysis, the resulting simulation can be as complex and hard to understand as the real brain.
- Such blind simulation could double our task.
- However, it has distinct merits:
 - Full read/write access and localized lesions.
 - Can investigate subjective phenomena such as consciousness (have the brain simulation study itself!).
 - Systematic, programmatic investigation becomes possible (automated science).

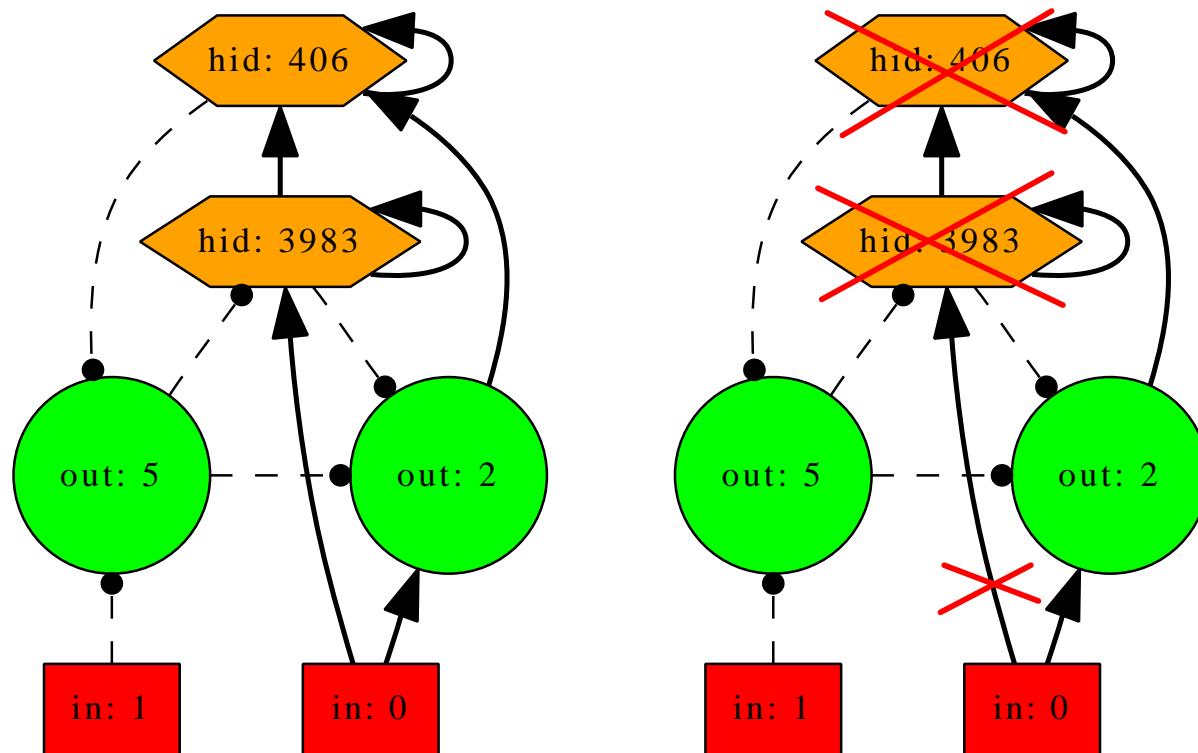
Conceptual Breakthroughs Needed



Choe and Smith (2006); Choe et al. (2007)

- Posing the right questions (Choe and Mann 2012): Internal perspective, problems faced by the brain itself.
- Sensorimotor perspective (Choe and Smith 2006; Choe et al. 2007).
- Developmental perspective.
- Evolutionary perspective (Chung and Choe 2011; Kwon and Choe 2008; Choe et al. 2012).
- Temporal perspective (Choe 2004; Lim and Choe 2008).

Inferring Function from a Brain Network: A Cautionary Tale



Analyze this!

- hid = hidden neuron, out = output neuron, in = input unit, arrow = excitatory connection, disc = inhibitory connection

Conclusion

- Understanding brain function requires a system-level investigation at a microscopic resolution.
- Innovative microscopy technologies are enabling a data-driven investigation linking the microstructure to the system level.
- A robust, accessible informatics platform is needed for knowledge discovery.
- Deep theoretical insights are needed to guide our investigation.

Acknowledgments

- People:
 - **BNL**: John Keyser, Louise C. Abbott, Bruce McCormick
 - **KESM**: Bruce McCormick, David Mayerich, Jaerock Kwon, Daniel Miller, Bernard Mesa (Micro Star), Todd Huffman (3Scan)
 - **KESM Brain Atlas**: Chul Sung, Ji Ryang Chung, Daniel Eng
- Funding:
 - IAMCS/KAUST (2010–2013); ● NSF CRCNS Data Sharing (#0905041, #1208174); ● NSF IDBR (#1256086); ● NIH/NINDS (#1R01-NS54252 & 03S1); ● NSF MRI (#0079874) and NSF ITR (#CCR-0220047), ● Texas Higher Education Coordinating Board (ATP #000512-0146-2001), ● TAMU CSE ● TAMU VPR ● 3Scan.

References

- Abbott, L. C. (2008). High-throughput imaging of whole small animal brains with the knife-edge scanning microscope. In *Neuroscience Meeting Planner, Washington, DC: Society for Neuroscience*. Program No. 504.2.
- Al-Kofahi, K. A., Lasek, S., Szarowski, D. H., Pace, C. J., Nagy, G., Turner, J. N., and Roysam, B. (2002). Rapid automated three-dimensional tracing of neurons from confocal image stacks. *IEEE Transactions on Information Technology in Biomedicine*, 6:171–187.
- Bock, D. D., Kerlin, A. M., Andermann, M. L., Hood, G., Wetzel, A. W., Yurgenson, S., Soucy, E. R., Kim, H. S., and Reid, R. C. (2011). Network anatomy and in vivo physiology of visual cortical neurons. *Nature*, 471:177–182.
- Choe, Y. (2004). The role of temporal parameters in a thalamocortical model of analogy. *IEEE Transactions on Neural Networks*, 15:1071–1082.
- Choe, Y., Abbott, L. C., Miller, D. E., Han, D., Yang, H.-F., Chung, J. R., Sung, C., Mayerich, D., Kwon, J., Micheva, K., and Smith, S. J. (2010). Multiscale imaging, analysis, and integration of mouse brain networks. In *Neuroscience Meeting Planner, San Diego, CA: Society for Neuroscience*. Program No. 516.3. Online.
- Choe, Y., Han, D., Huang, P.-S., Keyser, J., Kwon, J., Mayerich, D., and Abbott, L. C. (2009). Complete submicrometer scans of mouse brain microstructure: Neurons and vasculatures. In *Neuroscience Meeting Planner, Chicago, IL: Society for Neuroscience*. Program No. 389.10. Online.
- Choe, Y., Kwon, J., and Chung, J. R. (2012). Time, consciousness, and mind uploading. *International Journal on Machine Consciousness*, 4:257–274.

- Choe, Y., and Mann, T. A. (2012). From problem solving to problem posing. *Brain Mind Magazine*, 1:7–8.
- Choe, Y., and Smith, N. H. (2006). Motion-based autonomous grounding: Inferring external world properties from internal sensory states alone. In Gil, Y., and Mooney, R., editors, *Proceedings of the 21st National Conference on Artificial Intelligence(AAAI 2006)*, 936–941.
- Choe, Y., Yang, H.-F., and Eng, D. C.-Y. (2007). Autonomous learning of the semantics of internal sensory states based on motor exploration. *International Journal of Humanoid Robotics*, 4:211–243.
- Chung, J. R., and Choe, Y. (2011). Emergence of memory in reactive agents equipped with environmental markers. *IEEE Transactions on Autonomous Mental Development*, 3:257–271.
- Chung, J. R., Sung, C., Mayerich, D., Kwon, J., Miller, D. E., Huffman, T., Abbott, L. C., Keyser, J., and Choe, Y. (2011). Multiscale exploration of mouse brain microstructures using the knife-edge scanning microscope brain atlas. *Frontiers in Neuroinformatics*, 5:29.
- Chung, K., and Diesseroth, K. (2013). CLARITY for mapping the nervous system. *Nature Methods*, 10:508–513.
- Denk, W., and Horstmann, H. (2004). Serial block-face scanning electron microscopy to reconstruct three-dimensional tissue nanostructure. *PLoS Biology*, 19:e329.
- Friston, K. (2009). Causal modelling and brain connectivity in functional magnetic resonance imaging. *PLoS Biology*, 7:e1000033.
- Hagmann, P., Kurant, M., Gigandet, X., Thiran, P., Wedeen, V. J., Meuli, R., and Thiran, J.-P. (2007). Mapping human whole-brain structural networks with diffusion MRI. *PLoS ONE*, 2:e597.

- Han, D. (2009). *Rapid 3D Tracing of the Mouse Brain Neurovasculature with Local Maximum Intensity Projection and Moving Windows*. PhD thesis, Department of Computer Science, Texas A&M University.
- Han, D., Choi, H., Park, C., and Choe, Y. (2009a). Fast and accurate retinal vasculature tracing and kernel-isomap-based feature selection. In *Proceedings of the International Joint Conference on Neural Networks*, 1075–1082. Piscataway, NJ: IEEE Press.
- Han, D., Keyser, J., and Choe, Y. (2009b). A local maximum intensity projection tracing of vasculature in Knife-Edge Scanning Microscope volume data. In *Proceedings of the IEEE International Symposium on Biomedical Imaging*, 1259–1262.
- Haxby, J. V., Gobbini, M. I., Furey, M. L., Ishai, A., Schouten, J. L., and Pietrini, P. (2001). Distributed and overlapping representations of faces and objects in ventral temporal cortex. *Science*, 293(5539):2425–2430.
- Hayworth, K. J., Kasthuri, N., Schalek, R., and Lichtman, J. W. (2006). Automating the collection of ultrathin sections for large volume TEM reconstructions. *Microscopy and Microanalysis*, 12(Suppl. S02):86–87.
- Hintiryan, H., Gou, L., Zingg, B., Yamashita, S., Lyden, H. M., Song, M. Y., Grewal, A. K., Zhang, X., Toga, A. W., and Dong, H.-W. (2012). Comprehensive connectivity of the mouse main olfactory bulb: Analysis and online digital atlas. *Frontiers in Neuroanatomy*, 6(30).
- Izhikevich, E. M., and Edelman, G. M. (2008). Large-scale model of mammalian thalamocortical systems. *Proceedings of the National Academy of Sciences, USA*, 105:3593–3598.
- Jacobs, R. E., Ahrens, E. T., Dickinson, M. E., and Laidlaw, D. (1999). Towards a microMRI atlas of mouse development. *Computerized Medical Imaging and Graphics*, 23:15–24.

- Jain, V., Murray, J. F., Roth, F., Turaga, S., Zhigulin, V., Briggman, K. L., Helmstaedter, M. N., Denk, W., and Seung, H. S. (2007). Supervised learning of image restoration with convolutional networks. In *IEEE 11th International Conference on Computer Vision (ICCV 2007)*, 1–8.
- Jain, V., Seung, H. S., and Turaga, S. C. (2010). Machines that learn to segment images: a crucial technology for connectomics. *Current Opinion in Neurobiology*, 20:653–666.
- Kwon, J., and Choe, Y. (2008). Internal state predictability as an evolutionary precursor of self-awareness and agency. In *Proceedings of the Seventh International Conference on Development and Learning*, 109–114. IEEE.
- Kwon, J., Mayerich, D., Choe, Y., and McCormick, B. H. (2008). Lateral sectioning for knife-edge scanning microscopy. In *Proceedings of the IEEE International Symposium on Biomedical Imaging*, 1371–1374.
- Li, A., Gong, H., Zhang, B., Wang, Q., Wan, C., Wu, J., Liu, Q., Zeng, S., and Luo, Q. (2010). Micro-optical sectioning tomography to obtain a high-resolution atlas of the mouse brain. *Science*. In press.
- Lim, H., and Choe, Y. (2008). Extrapolative delay compensation through facilitating synapses and its relation to the flash-lag effect. *IEEE Transactions on Neural Networks*, 19:1678–1688.
- Luisi, J., Narayanaswamy, A., Galbreath, Z., and Roysam, B. (2011). The FARSIHT trace editor: An open source tool for 3-d inspection and efficient pattern analysis aided editing of automated neuronal reconstructions. *Neuroinformatics*, 9:305–315.
- Mayerich, D., Abbott, L. C., and McCormick, B. H. (2008). Knife-edge scanning microscopy for imaging and reconstruction of three-dimensional anatomical structures of the mouse brain. *Journal of Microscopy*, 231:134–143.

- Mayerich, D., Kwon, J., Sung, C., Abbott, L. C., Keyser, J., and Choe, Y. (2011). Fast macro-scale transmission imaging of microvascular networks using KESM. *Biomedical Optics Express*, 2:2888–2896.
- McCormick, B. H. (2004). System and method for imaging an object. USPTO patent #US 6,744,572 (for Knife-Edge Scanning; 13 claims).
- Micheva, K., and Smith, S. J. (2007). Array tomography: A new tool for imaging the molecular architecture and ultrastructure of neural circuits. *Neuron*, 55:25–36.
- Mikula, S., Binding, J., and Denk, W. (2012). Staining and embedding the whole mouse brain for electron microscopy. *Nature Methods*, 9:1198–1201.
- Milo, R., Sen-Orr, S., Itzkovitz, S., Kashtan, N., Chklovskii, D., and Alon, U. (2002). Network motifs: Simple building blocks of complex networks. *Science*, 298:824–827.
- Mitra, P. P. (2012). Technical white paper: Mouse brain architecture project. Technical report, Cold Spring Harbor Laboratory.
- Murray, J. M. (2011). Methods for imaging thick specimens: Confocal microscopy, deconvolution, and structured illumination. *Cold Spring Harbor Protocols*, 2011(12):1399–1437.
- Ragan, T., Kadiri, L. R., Venkataraju, K. U., Bahlmann, K., Sutin, J., Taranda, J., Arganda-Carreras, I., Kim, Y., Seung, H. S., and Osten, P. (2012). Serial two-photon tomography for automated *ex vivo* mouse brain imaging. *Nature Methods*, 9:255–258.

- Seung, H. S. (2012). *Connectome: How the Brain's Wiring Makes Us Who We Are*. Boston, MA: Houghton Mifflin Harcourt.
- Seung, S., and Burnes, L. (2012). Eyewire. <http://eyewire.org/>.
- Sporns, O. (2002). Graph theory methods for the analysis of neural connectivity patterns. In Kötter, R., editor, *Neuroscience Databases: A Practical Guide*. Boston, MA: Kluwer Publishers.
- Sporns, O. (2011). *Networks of the Brain*. Cambridge, MA: MIT Press.
- Sporns, O. (2012). *Discovering the Human Connectome*. Cambridge, MA: MIT Press.
- Sporns, O., and Tononi, G. (2002). Classes of network connectivity and dynamics. *Complexity*, 7:28–38.
- Sporns, O., Tononi, G., and Kötter, R. (2005). The human connectome: A structural description of the human brain. *PLoS Computational Biology*, 1:e42.
- Swanson, L. W. (2003). *Brain Architecture: Understanding the Basic Plan*. Oxford: Oxford University Press.
- Thiel, A., Schwegler, H., and Eurich, C. W. (2003). Complex dynamics is abolished in delayed recurrent systems with distributed feedback times. *Complexity*, 8:102–108.
- Tsai, P. S., Friedman, B., Ifarraguerri, A. I., Thompson, B. D., Lev-Ram, V., Schaffer, C. B., Xiong, Q., Tsien, R. Y., Squier, J. A., and Kleinfeld, D. (2003). All-optical histology using ultrashort laser pulses. *Neuron*, 39:27–41.

Van Essen, D., Ugurbil, K., Auerbach, E., Barch, D., Behrens, T., Bucholz, R., Chang, A., Chen, L., Corbetta, M., Curtiss, S., Penna, S. D., Feinberg, D., Glasser, M., Harel, N., Heath, A., Larson-Prior, L., Marcus, D., Michalareas, G., Moeller, S., Oostenveld, R., Petersen, S., Prior, F., Schlaggar, B., Smith, S., Snyder, A., Xu, J., and Yacoub, E. (2012). The human connectome project: A data acquisition perspective. *NeuroImage*, 62(4):2222 – 2231.

White, J. G., Southgate, E., Thomson, J. N., and Brenner, S. (1986). The structure of the nervous system of the nematode *caenorhabditis elegans*. *Philosophical Transactions of the Royal Society of London B*, 314:1–340.

Yang, H.-F., and Choe, Y. (2011a). Ground truth estimation by maximizing topological agreements in electron microscopy data. In *Proceedings of the 7th International Symposium on Visual Computing (LNCS 6938)*, 371–380.

Yang, H.-F., and Choe, Y. (2011b). An interactive editing framework for electron microscopy image segmentation. In *Proceedings of the 7th International Symposium on Visual Computing (LNCS 6938)*, 400–409.