

# Whole Brain Mapping with X-rays

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台大物理 27/10/2020

# Collaborators: A true international effort!

## • Taiwan

- Keng S. Liang, Ting-Kuo Lee, Maw-Kun Wu, Shih-Chang Lee, En-Te Hwu, Ming-Li Chu, Chih-Hsiun Lin (Academia Sinica): phase retrieval, reconstruction, magnetic nanoparticles, microfluidity
  - Ann-Shyn Chiang, Chia-Wei Li (Life Science, Tsing Hua U.): cell biology, shell fish, fossil, fire fly
  - Hong-Ming Lin (Tatung U): electron chemistry, battery
  - Yu-Tai Ching (NCTU), David Chien (UCS-St. Marcos): Image Processing, reconstruction
  - Chung-Shi Yang, Kelvin K. C. Tsai (NHRI): nanomedicine, microangiogenesis
  - Ann Chen, Maria Ka, Dueng-Yuan Hong (TSGH): CKD vessel imaging
  - Y. F. Hu, S. K. Tsai (TTY Pharmaceuticals): (drug delivery, pharmacokinetics)
  - Y. C. Yang, Hong-I Yeh, Yu-Jen Chen (Mackay Memorial Hospital): (artery disease, tumor development)
- 
- Japan: T. Ishikawa, Y. Kohmura, K. Sawada, Y. Joti (RIKEN/SPring-8 Center)
  - Korea: Jung-Ho Je (POSTECH), Doyoung Noh (GIST), Jun Lim, Jae-Hong Lim (PAL)
  - China: Jun Hu, Chunhai Fan, Lihua Wang, Xiaoqing Cai, Chichao Zhang, Ying Zhu
  - Singapore: Eng Soon Tok, Alvin Teo, Chian-Ming Low, Gan-Moog Chow, H. O. Moser (NUS)—drug delivery, polymer blend, electrochemistry, neurobiology.
  - France: Cyril Petibois, Sophie Javerzat, Michele Moenner (U. Bordeaux, brain tumor angiogenesis), Patrick Soukiassian
  - USA: Yong S. Chu, Wah-Keat Lee, Qun Shen, B. Lai, (APS), Wen-An Chiou (U. Maryland), John Boeckl (US Air Force Lab)—TXM
  - Switzerland:
    - G. Margaritondo (EPFL)
    - Rolf Gruetter (EPFL)

# NanoX Team (Current Members)

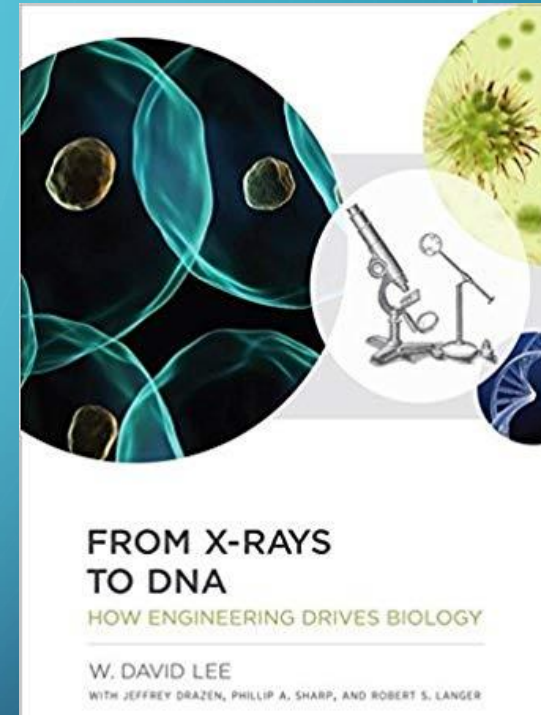
- X-ray microscopy:
  - Hsiang-Hsin Chen (陳翔欣), Shun-Min Yang (楊舜閔), Tsung-Tse Lee (李宗澤), Cheng-Huan Hsu (徐晟桓), Ching-Yu Chiou(邱鏡宇)
- X-ray Nanosynthesis:
  - Ming-Tsang Lee (李旻倉)
- Nanofabrication of X-ray optics:
  - Mei-Chun Chen (陳玫君), Yu-Ting Jian (簡郁庭)
- Bioimaging:
  - Shun-Min Yang (楊舜閔), Hsiang-Hsin Chen (陳翔欣), Yi-Yun Chen (陳怡云), Chia-Ru Chang (張家如), Ya-Sian Wang (王雅嫻), Shiou-Jin Chiou (邱繡謹), Cheng Jyun Yang (楊程鈞)
- Visitors: Cyril Petibois, Pei-Feng Chen, Keng S. Liang, Benoit Recur

# Innovation in physics and instrumentation has opened new eras of biology and medicine

- Microscopy
- Spectroscopy
- X-rays scattering
- Mass spectrometry
- Nuclear Medicine
- NMR, SPECT, PET, CT, ultrasound, OCT
- ...

## And, More Recently..

- Laser confocal scanning microscopy + fluorescent protein
- Protein crystallography
- Cryo-electron microscopy

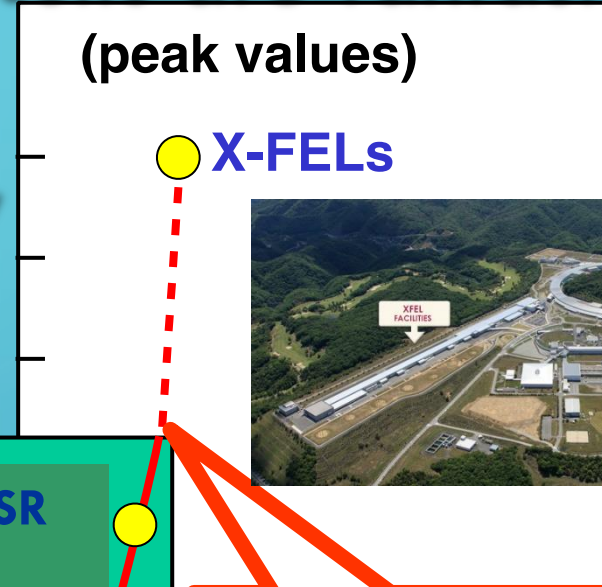
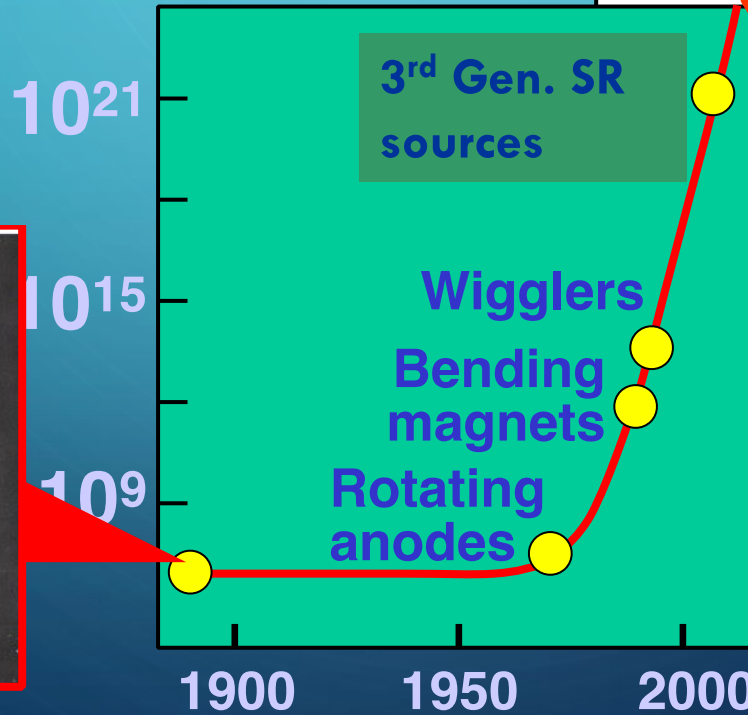
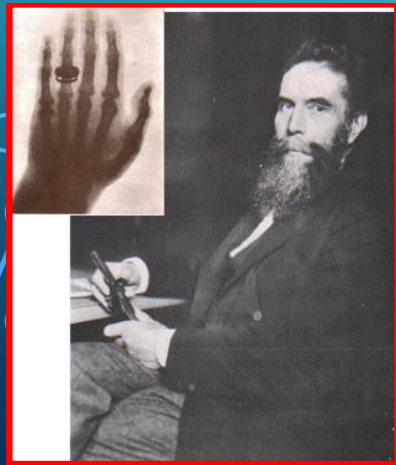


# The breakthroughs in x-ray imaging

- Phase contrast
- Nano-resolution
- Elemental contrast
- High speed for 3D imaging

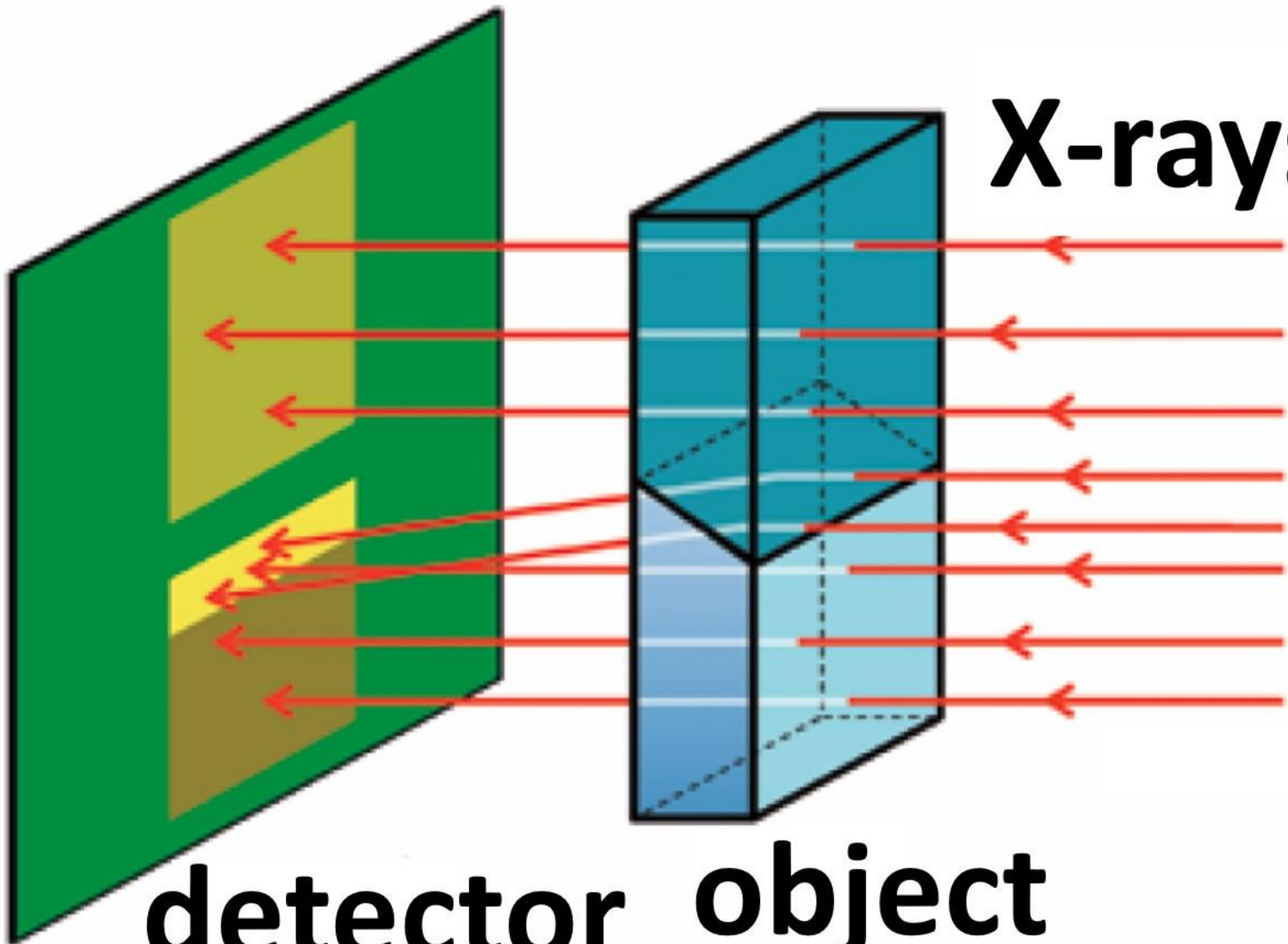
# It took decades to obtain the first x-ray free electron lasers, but the results are fantastic:

Historical increase of the x-ray source brightness (in standard units)



Taiwan Photon Source

**X-rays**



**detector object**

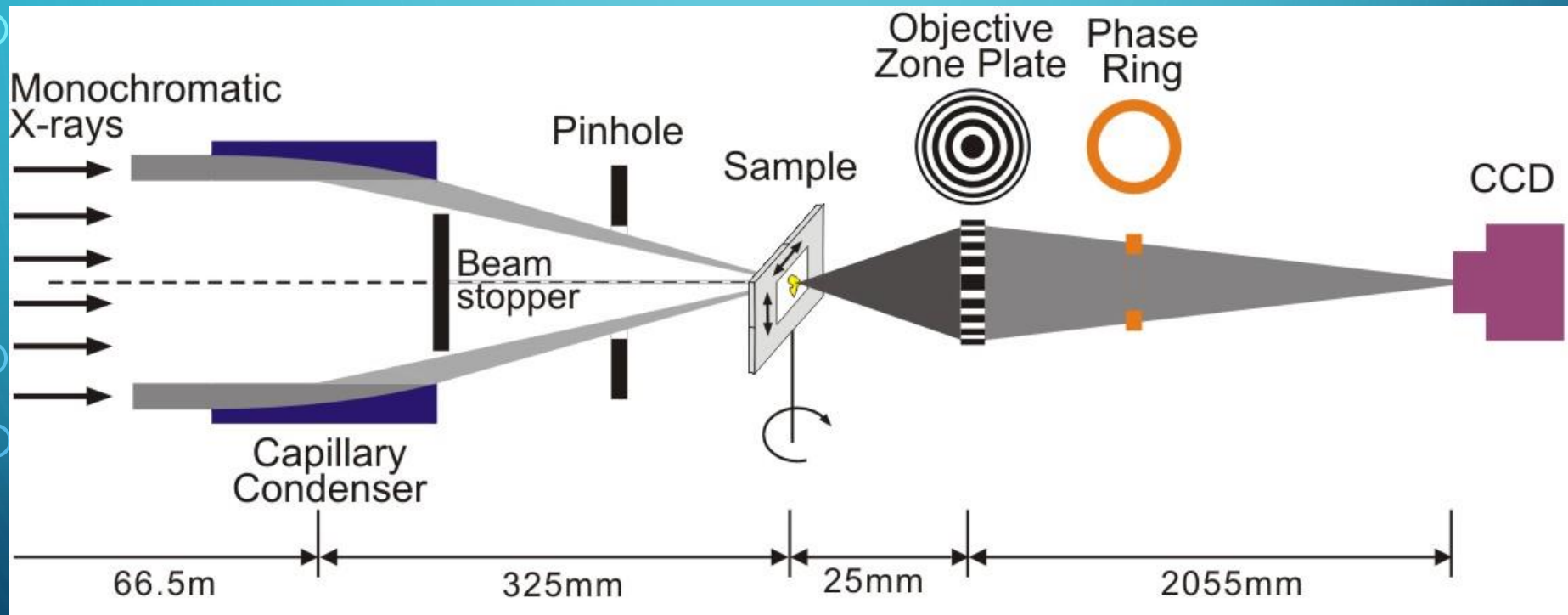
D= 1mm



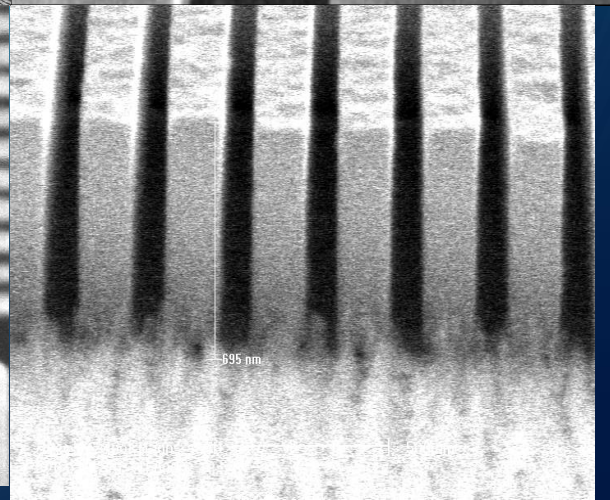
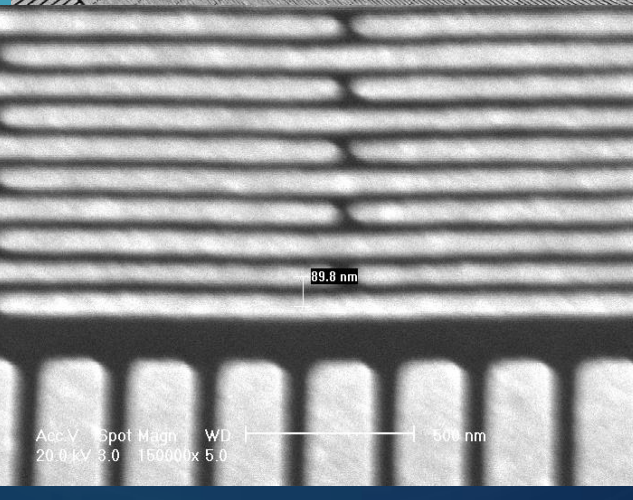
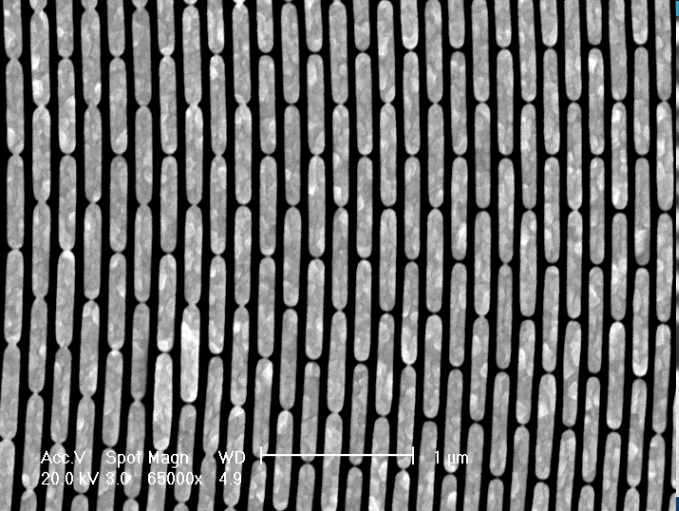
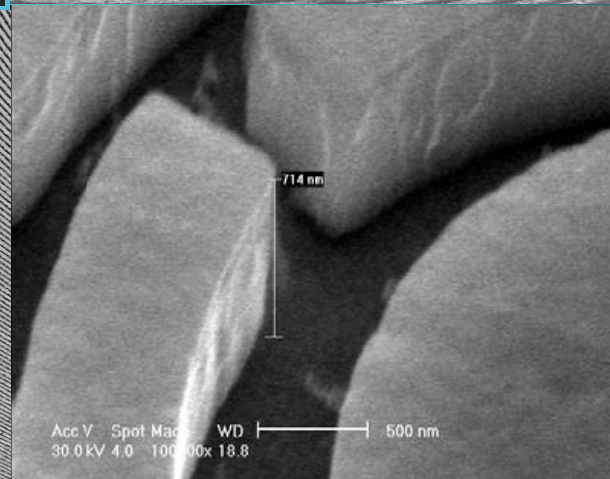
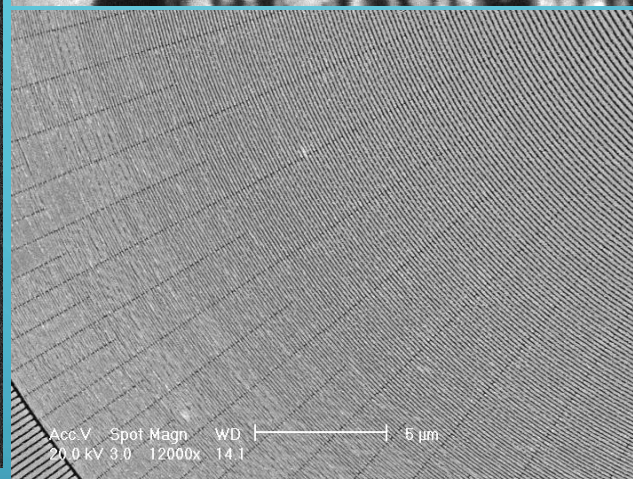
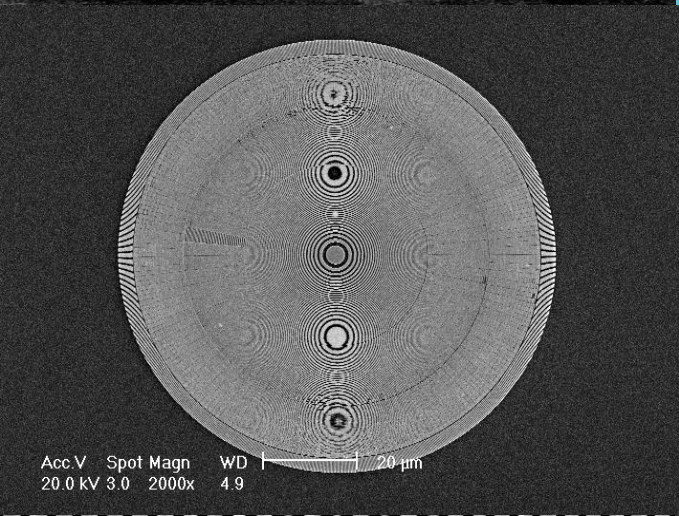
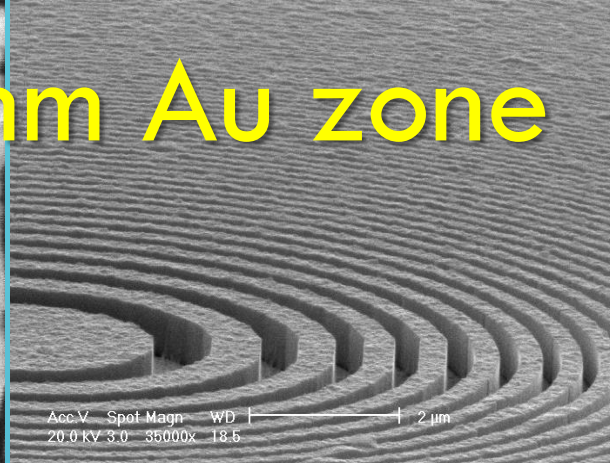
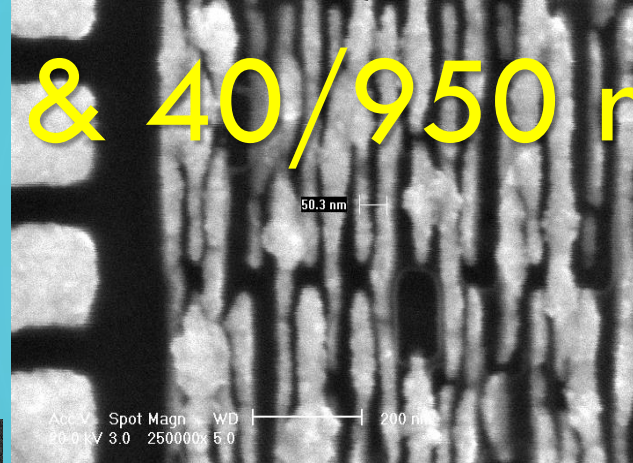
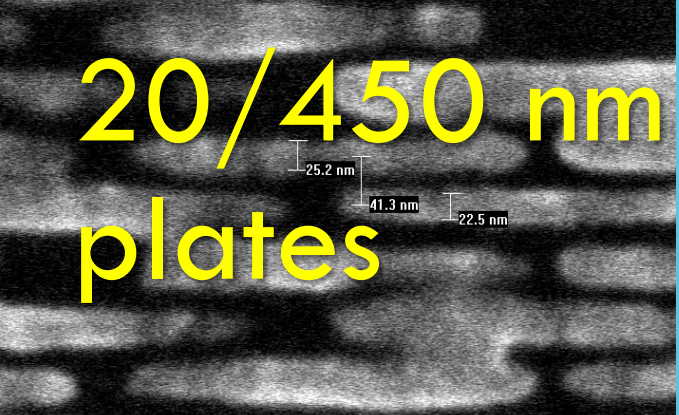


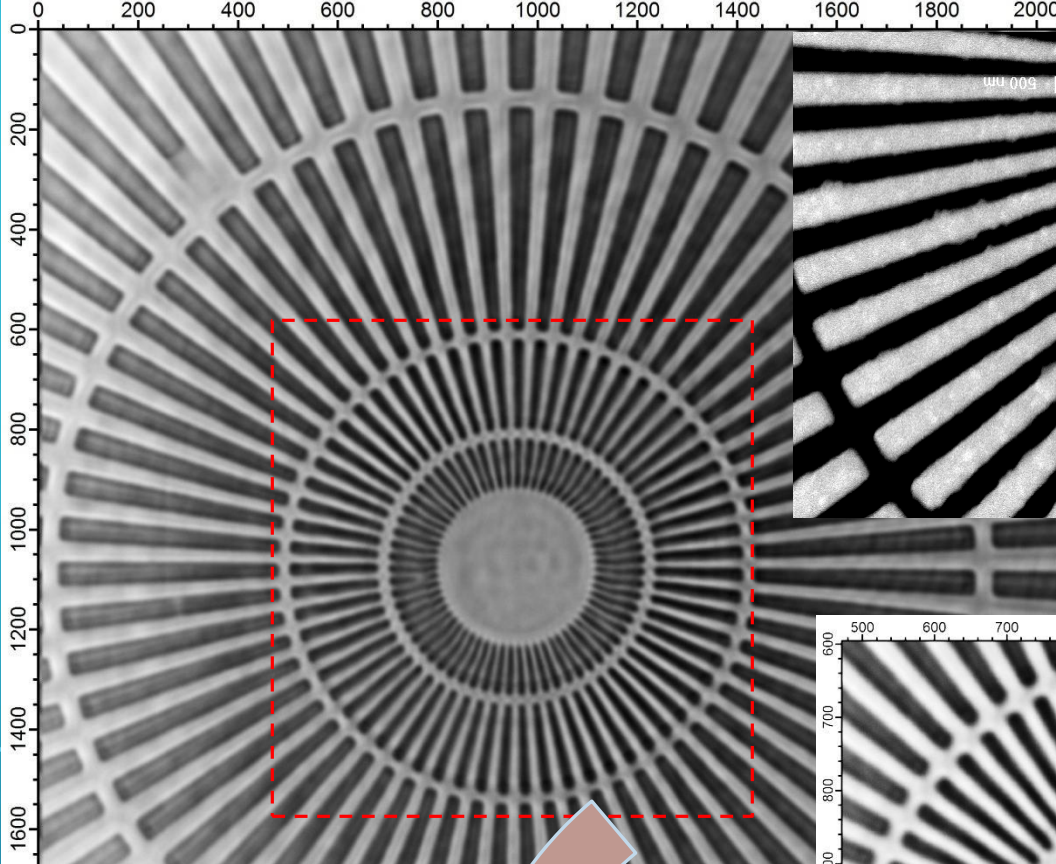
# The second breakthrough: nanoscale resolution

# Transmission x-ray microscopy with Fresnel zone plate optics



# 20/450 nm & 40/950 nm Au zone plates





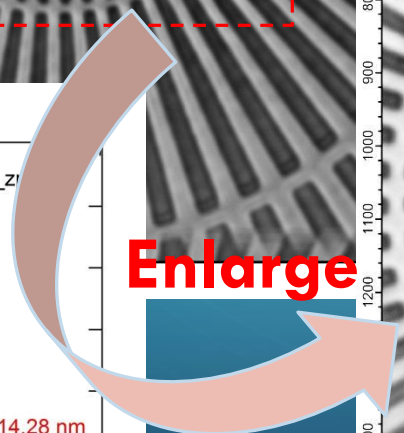
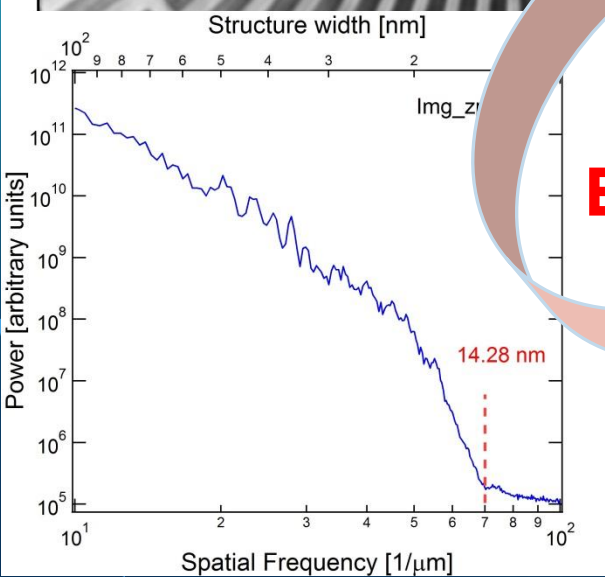
## Star Pattern Image

Finest line width: 20nm  
Au thickness: 300nm

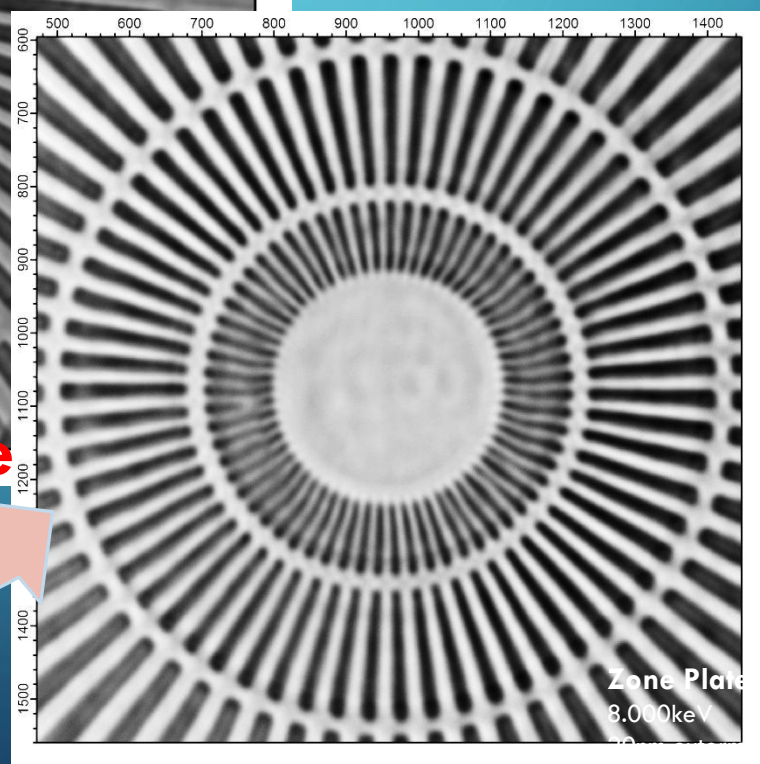
Pixel size: 2.51 nm  
FOV: 5.14um

No binning  
8 sec exposure  
60 frames averaged

4 pieces of diffuser



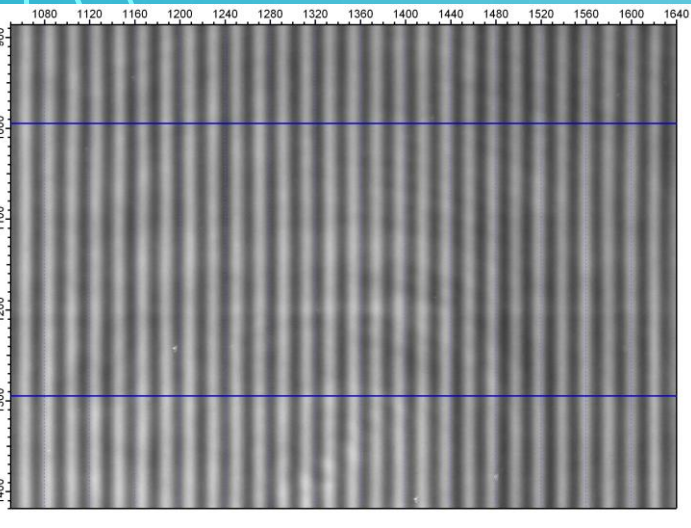
**Enlarge**



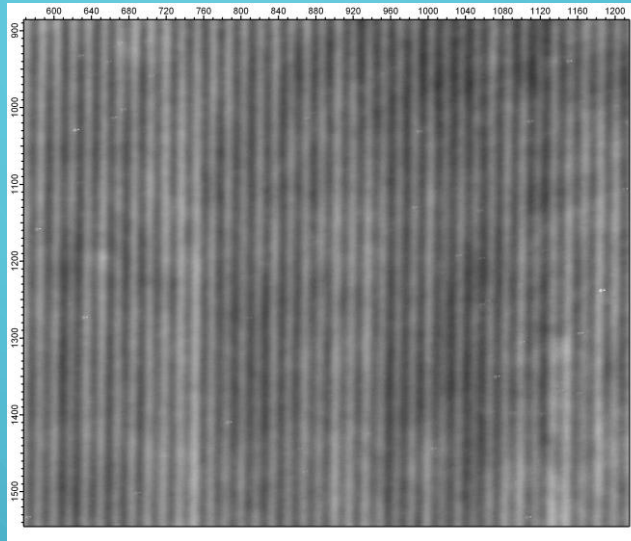
Zone Plate applied (ZP m2-6):  
8.000keV  
70mm diameter  
50um diameter  
400nm thickness

# Si/W multilayer test patterns

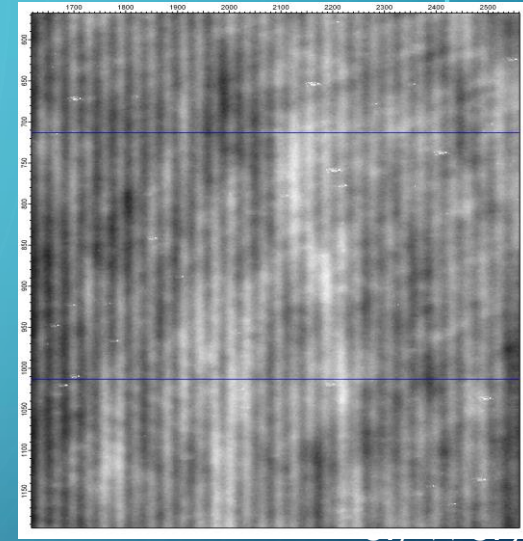
H. R. Wu, et al. Nanoresolution Radiology of Neurons, J. Phys. D 45, 242001 (2012).



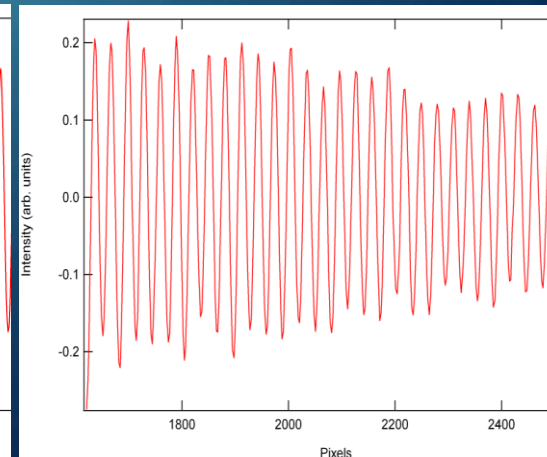
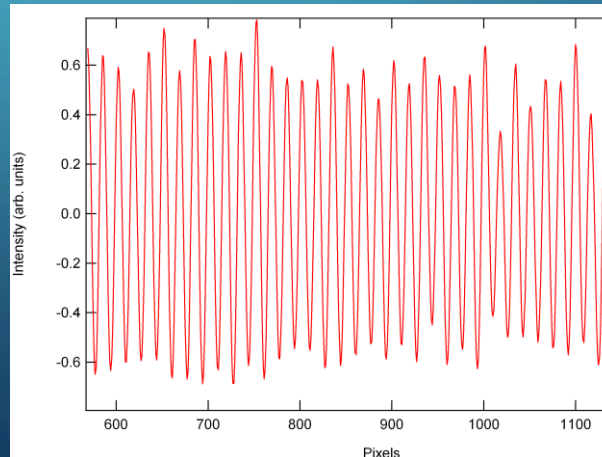
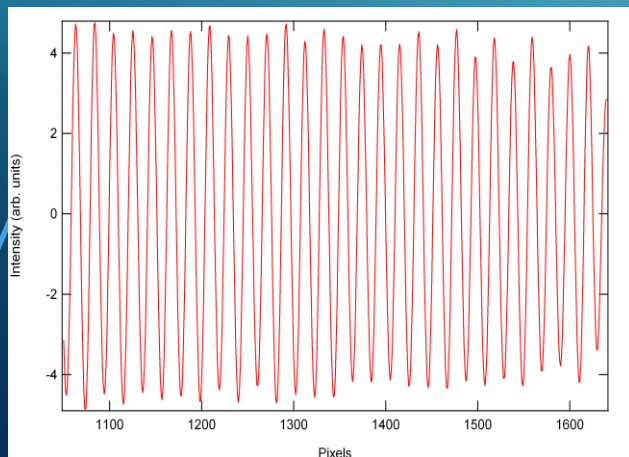
Finest line width = 25nm



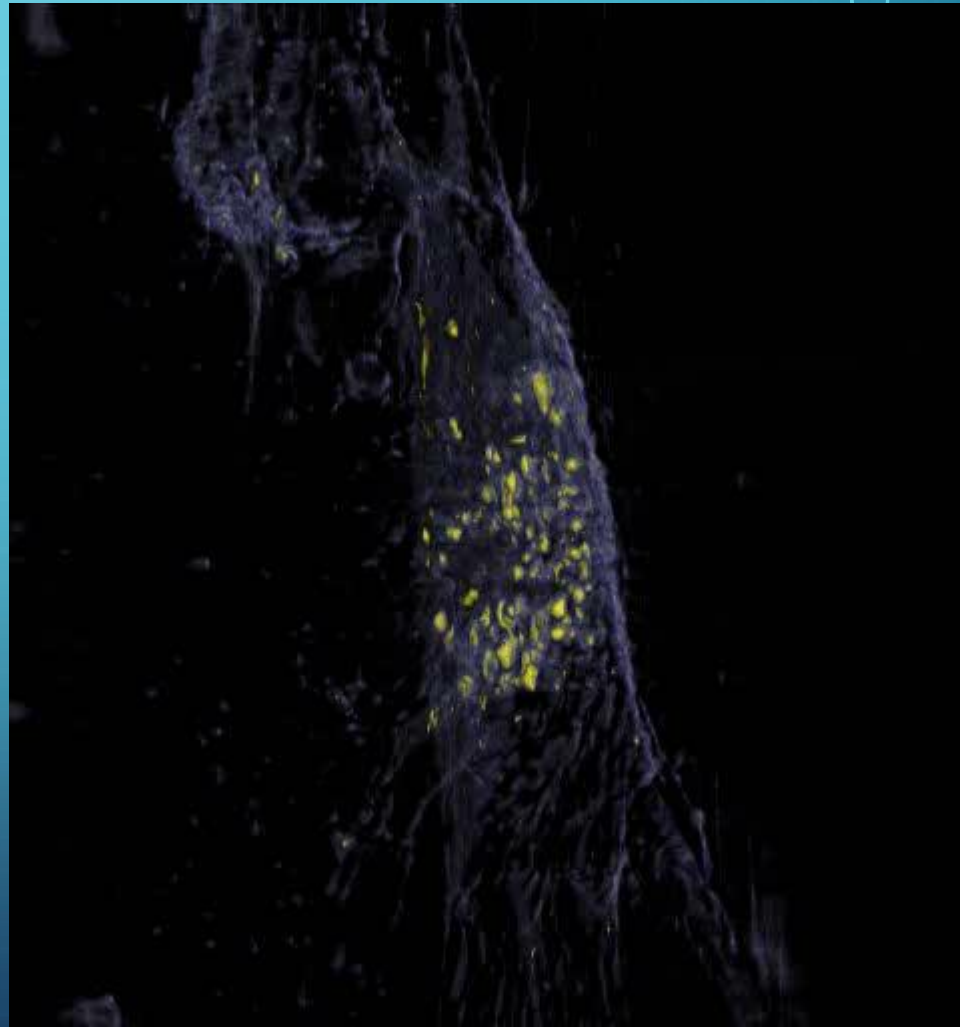
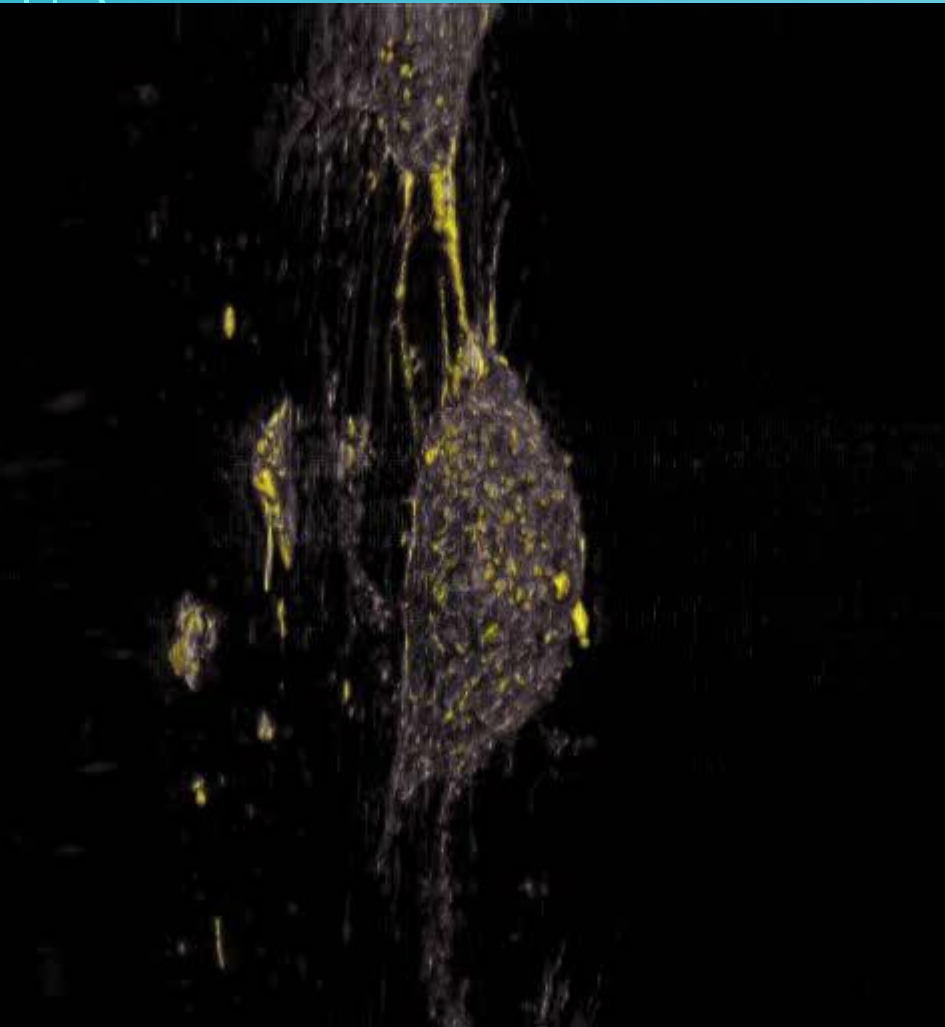
20 nm



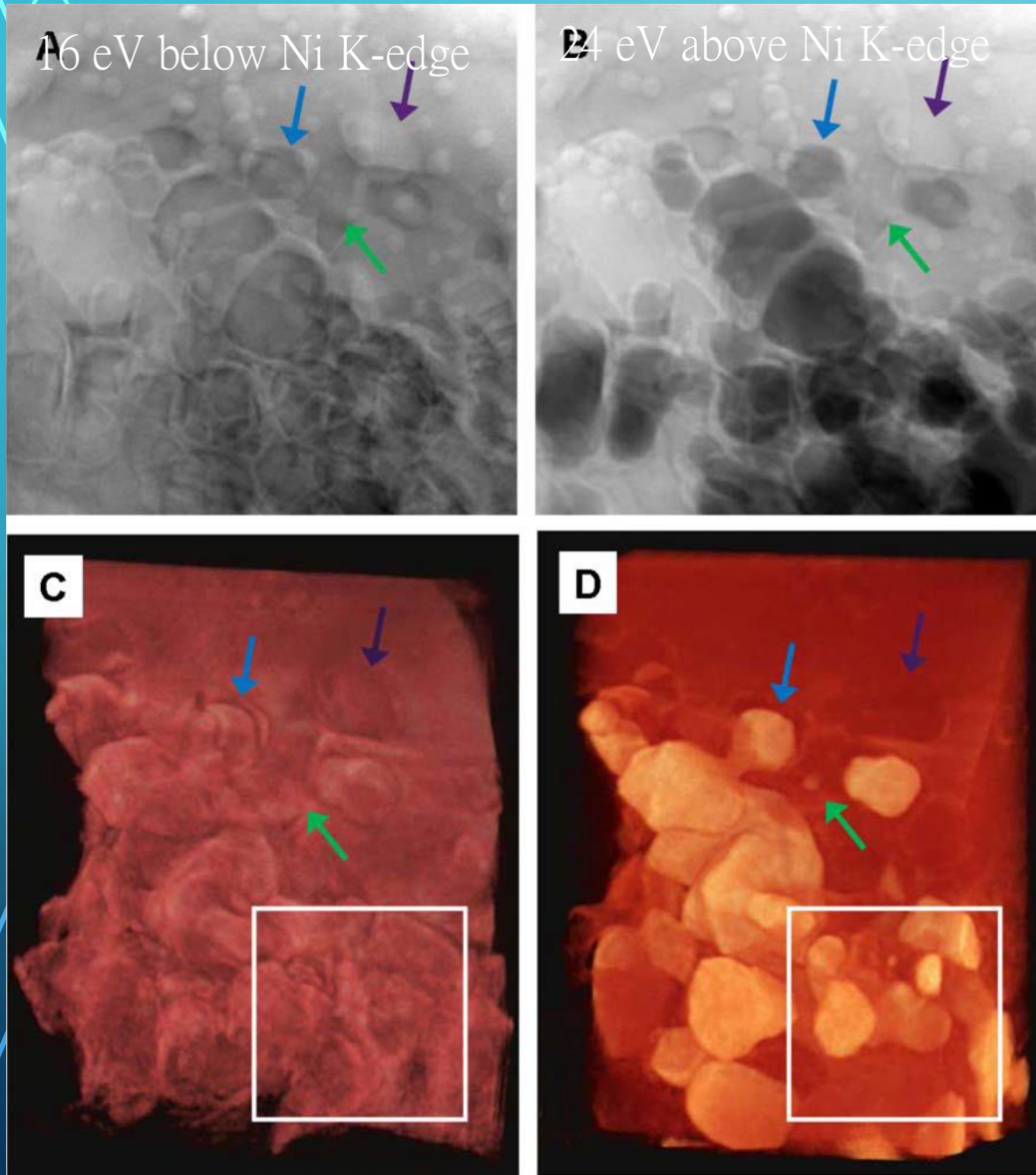
15 nm



# HeLa cells with AuNPs on culture dish



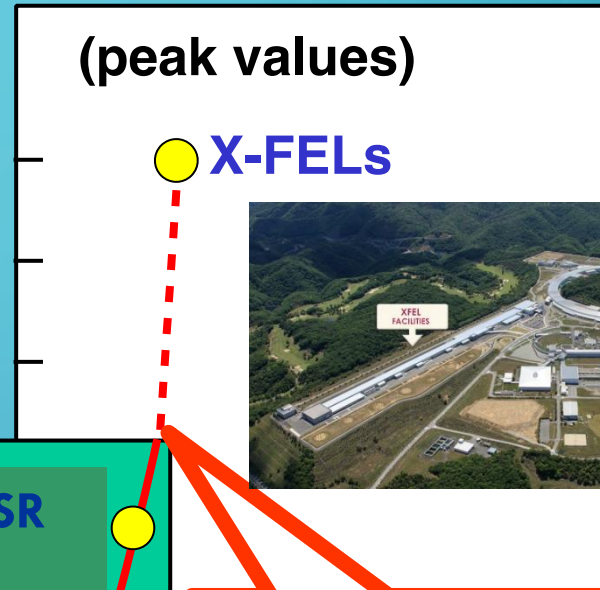
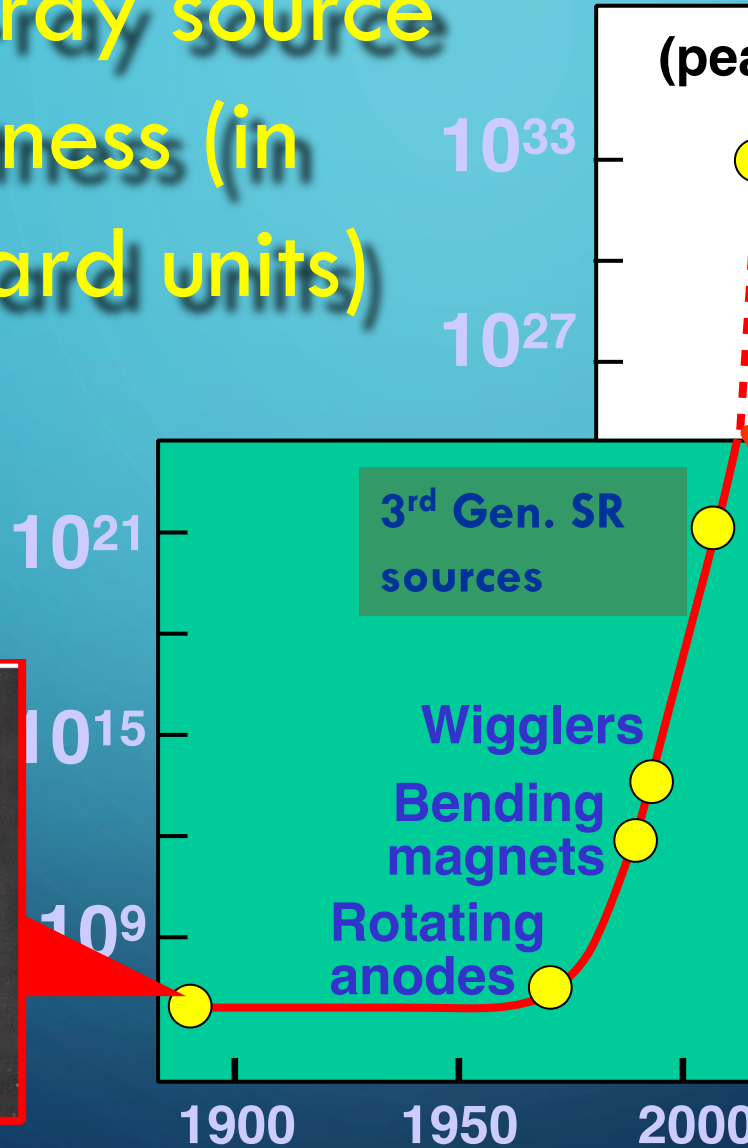
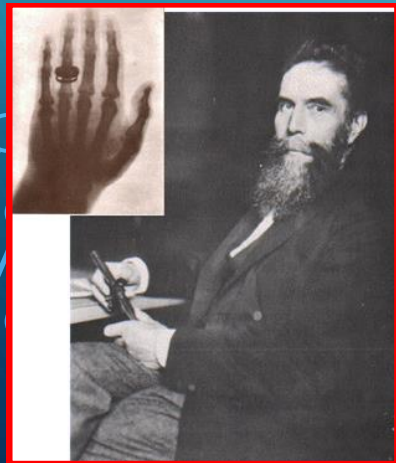
# Element mapping with nanoresolution



J Electrochem Soc 157,  
B783 (2010) **Figure 2**

- Ni – YSZ (yttria-stabilized zirconia) anode
- Ni K-edge: 8.333 keV
- Spatial resolution: 38.5 nm
- Box size: 5  $\mu\text{m}$

# Historical increase of the x-ray source brightness (in standard units)



Taiwan Photon Source

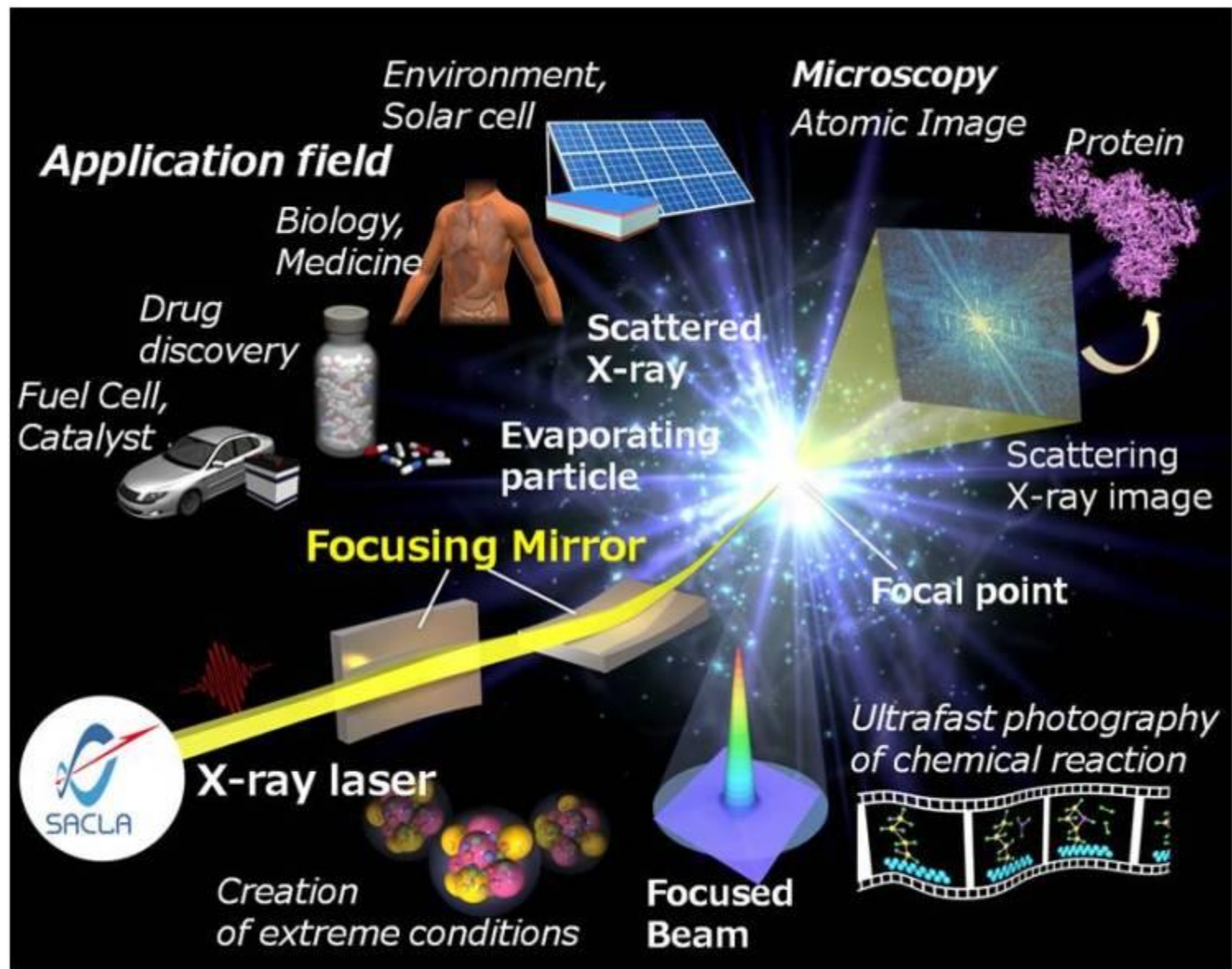




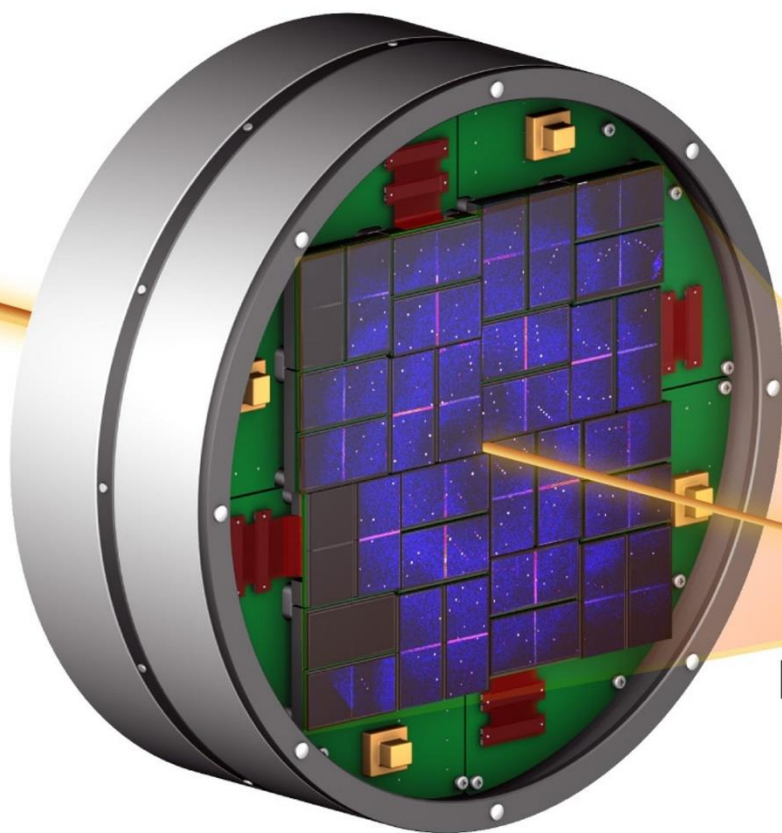
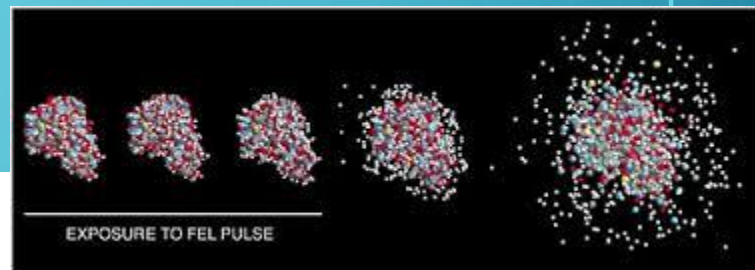
# Phase contrast: the first breakthrough

The background is a dark blue gradient with faint, large circular patterns. In the corners, there are white line-art graphics resembling circuit traces or fiber optic paths, with small circles at the end of the lines.

# The XFEL: Toward the ultimate x-ray source

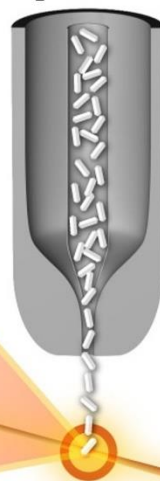


# “Detect and Destroy”



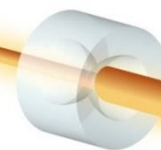
**CSPAD detector**  
( $z = 93$  mm)

**Liquid Jet**



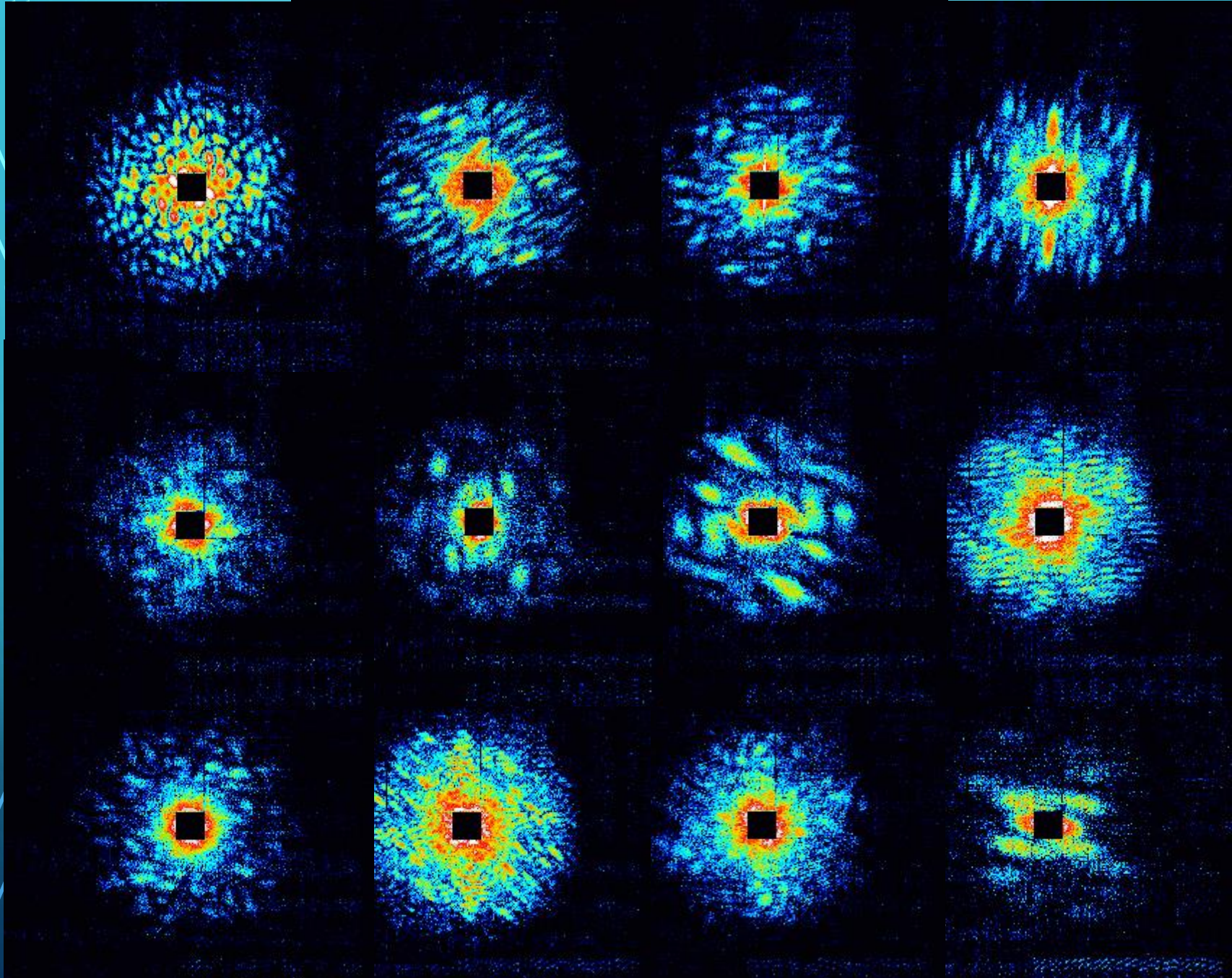
**Interaction Point**  
( $10 \mu\text{m}^2$  focus)

**Be lenses**

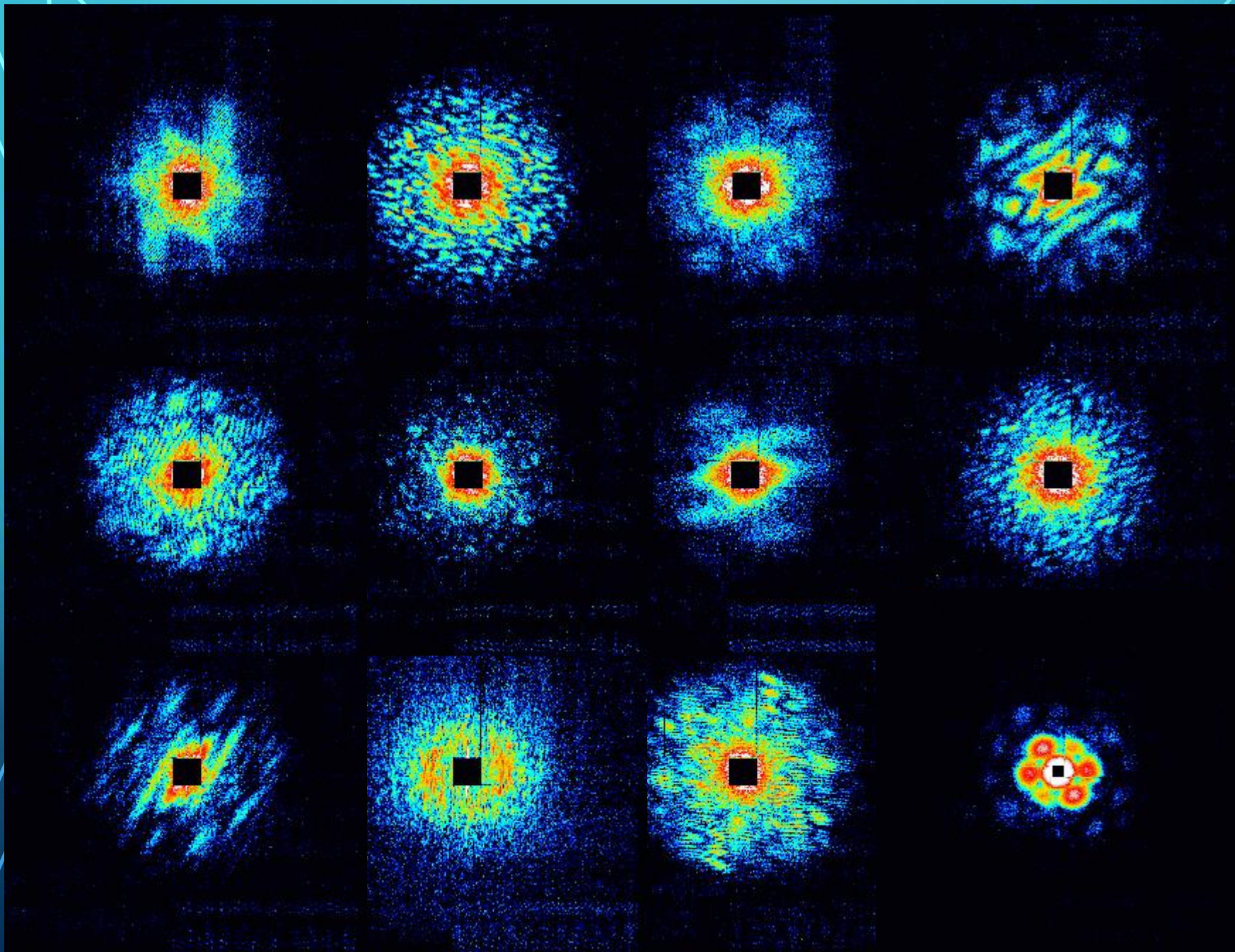


**Undulator**  
(420 m upstream)

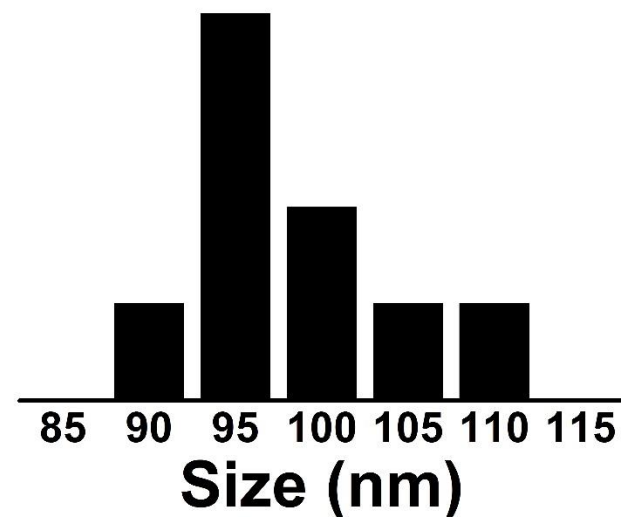
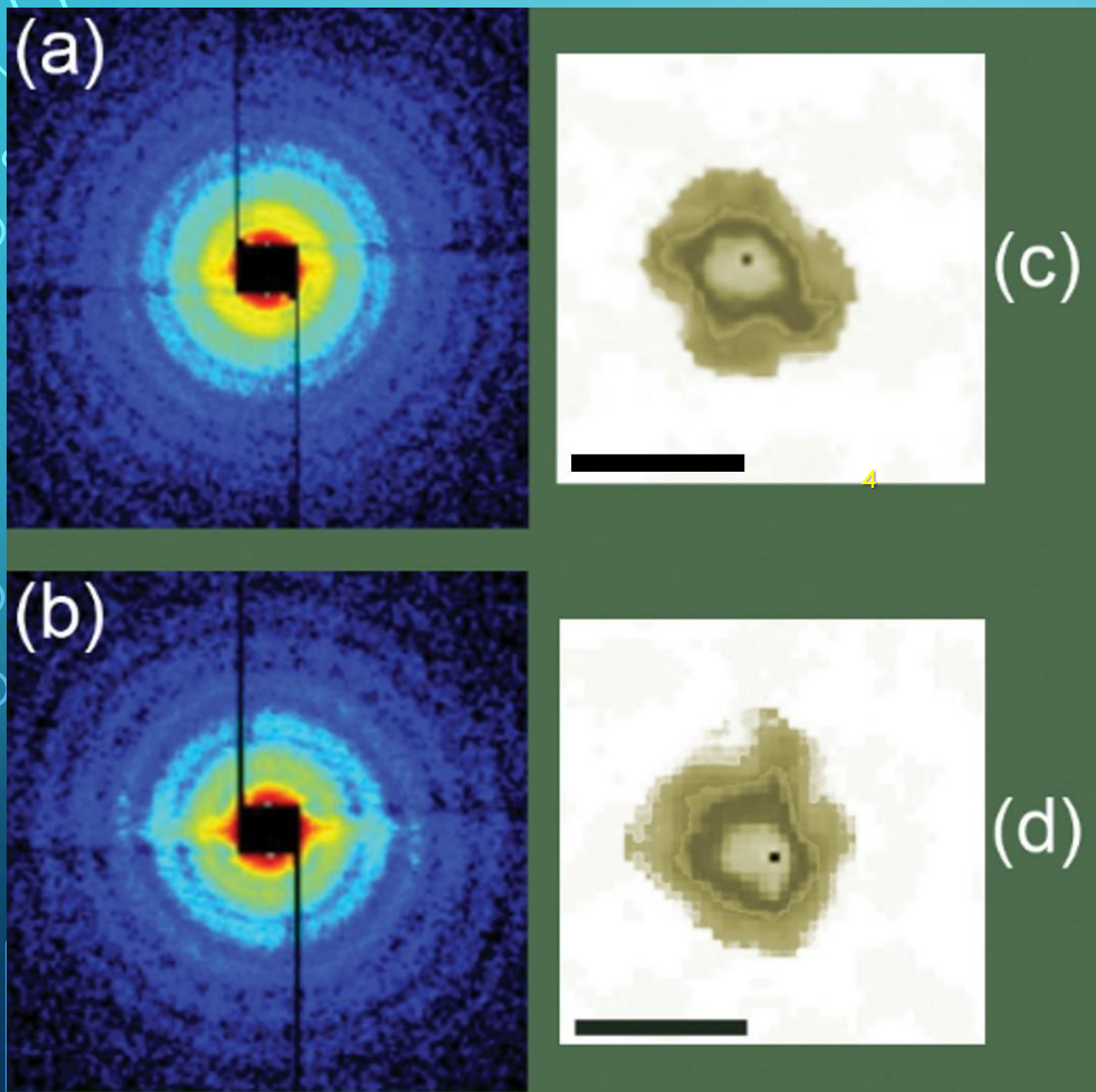




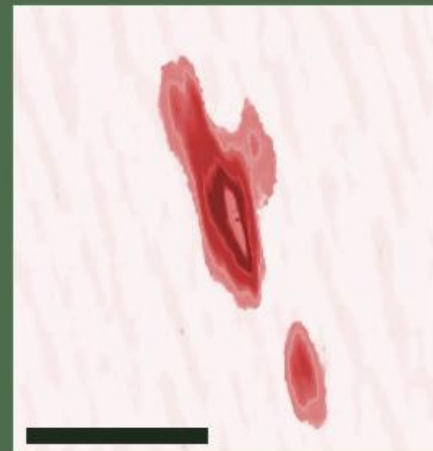
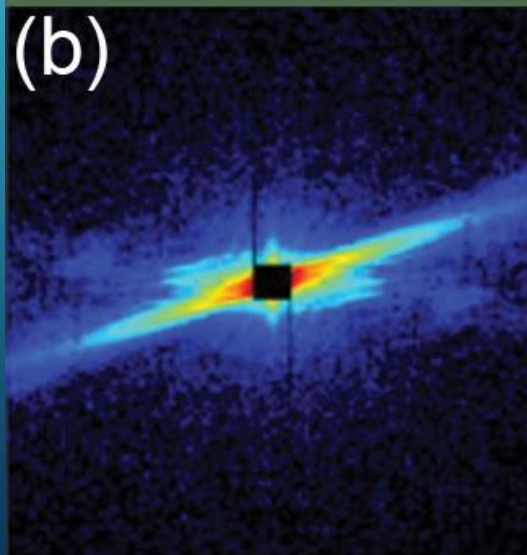
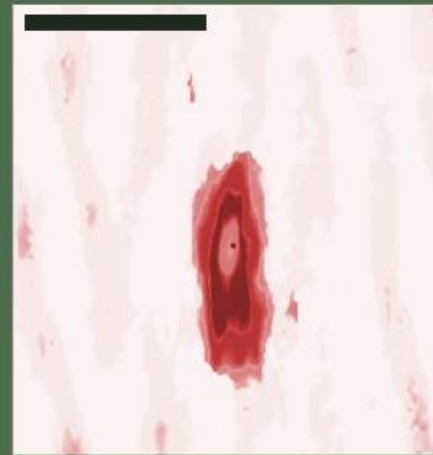
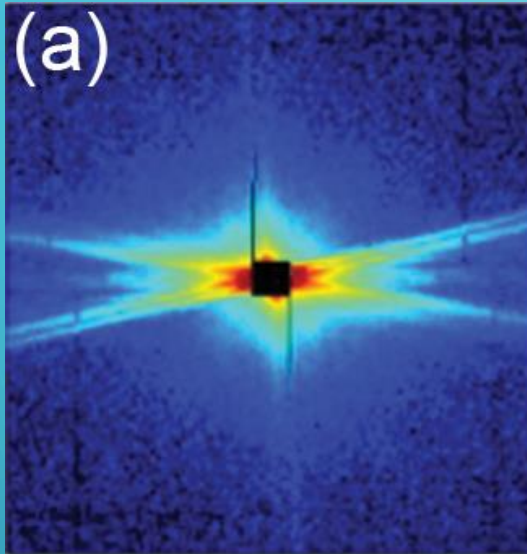
In collaboration with Prof. Yoshinori Nishino of Hokkaido University and SACLA



# Blank liposome ((NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> inside)

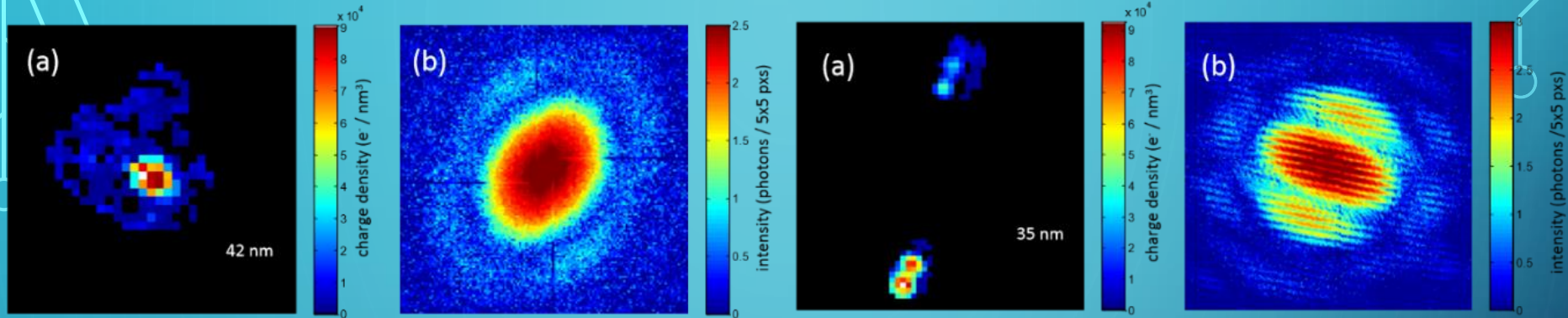


# Doxorubicin loaded liposome



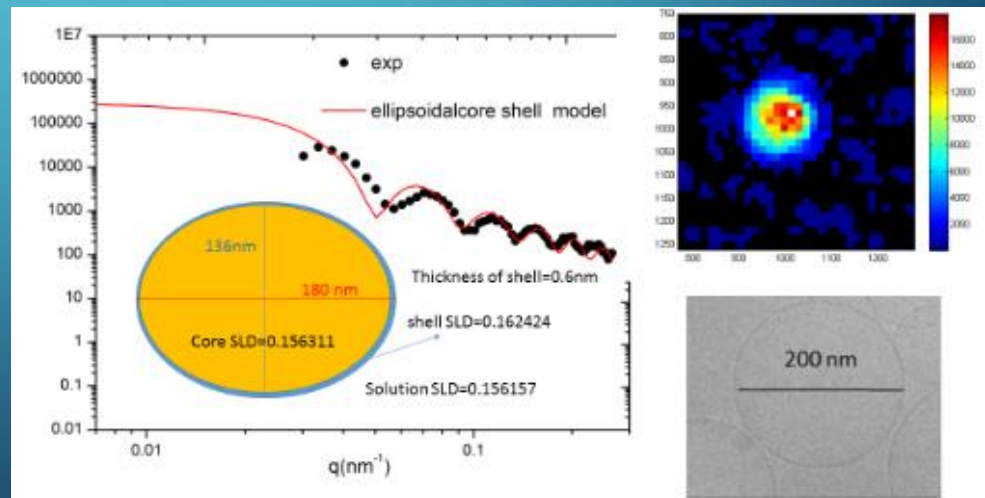
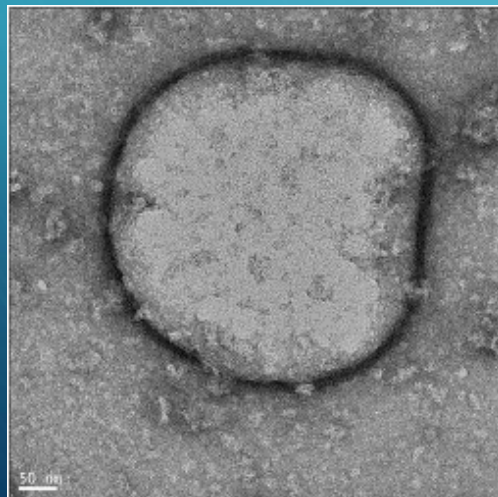


# Coherent diffraction imaging (XCDI)



A single 40 nm AuNP

Two 10 nm AuNP



A single 200 nm liposome

# Tumor Induced Angiogenesis

Cyril Petibois et al. (U. Bordeaux), DY Hong et al. (TSGH),

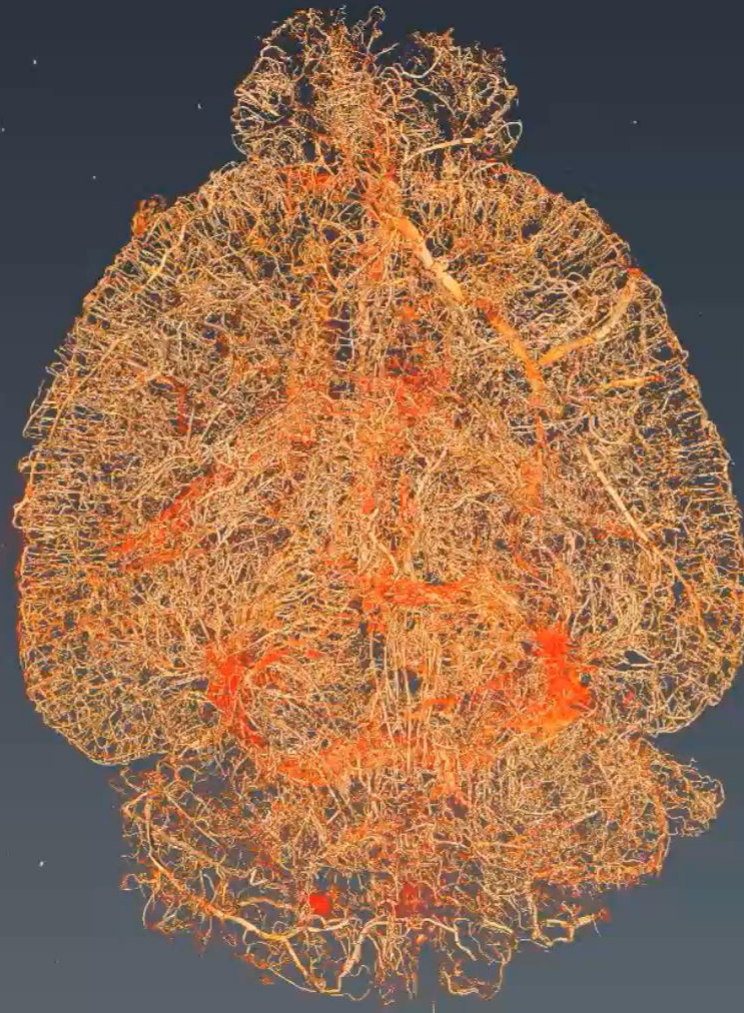
# Large tissue, high resolution and speed

Whole-brain blood vessel reconstruction  
in mouse

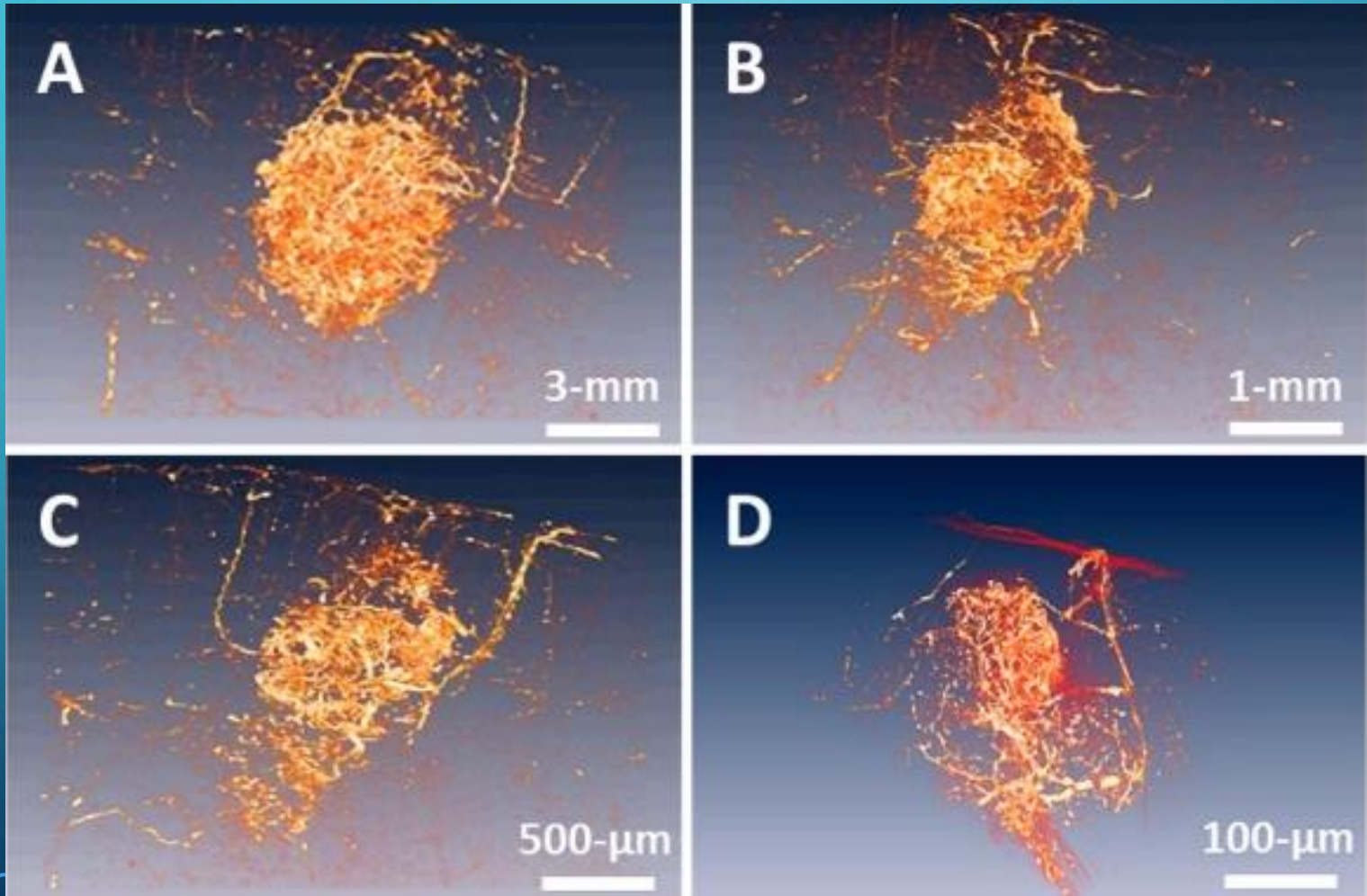
Res. < 0.5  $\mu\text{m}$

Size > 400 mm

Time < 10 min

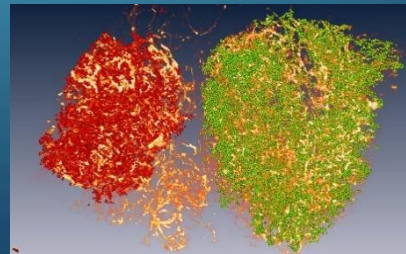
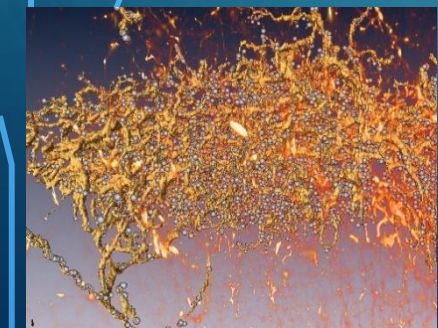
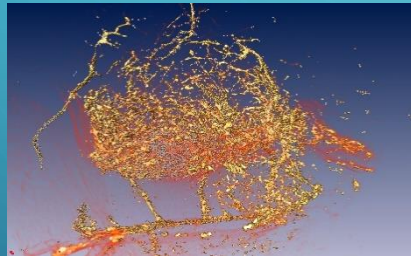
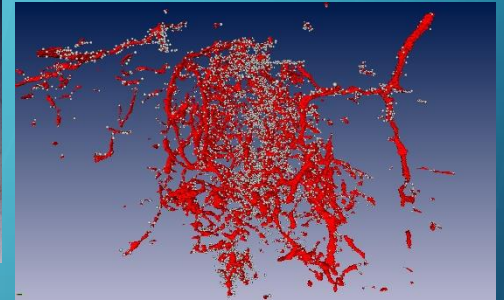
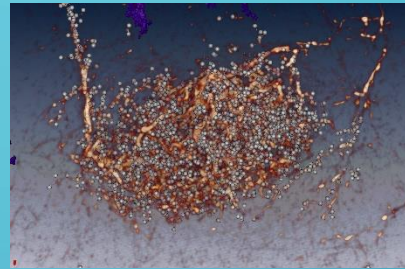
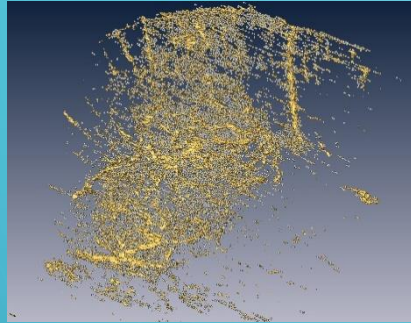


# X-ray imaging of angiogenic microvasculature of glioblastoma

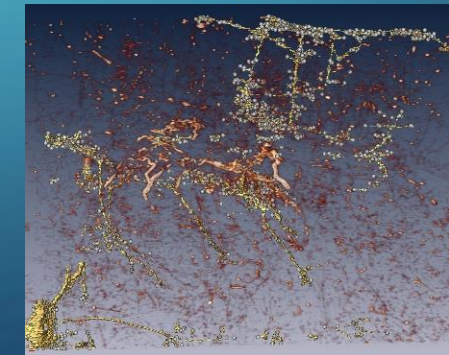
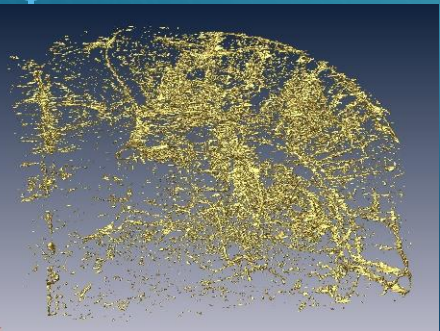
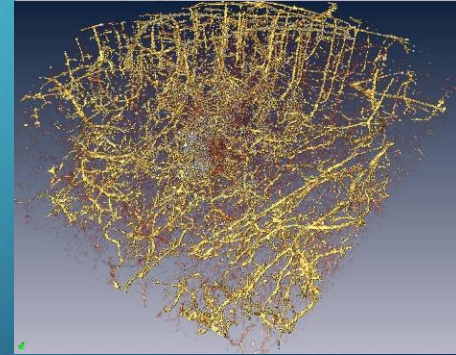
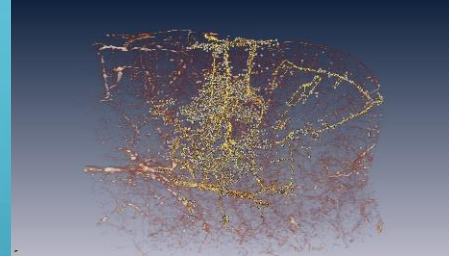
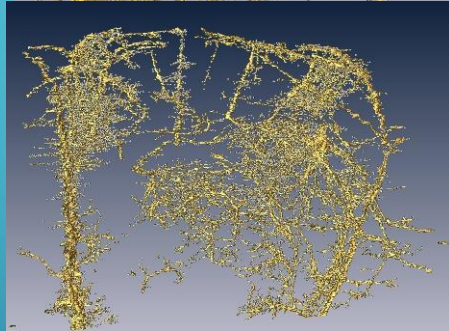
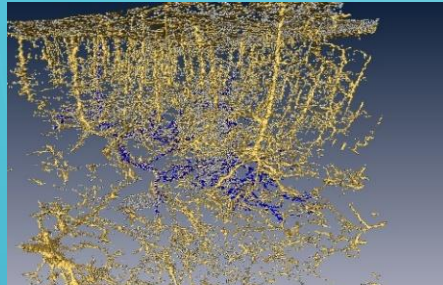




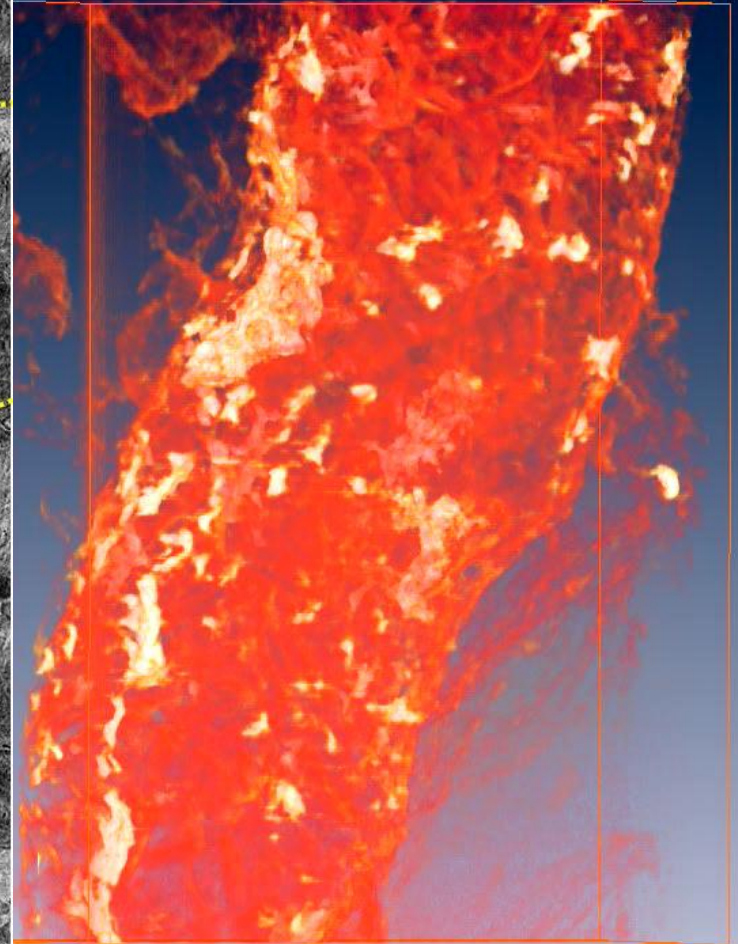
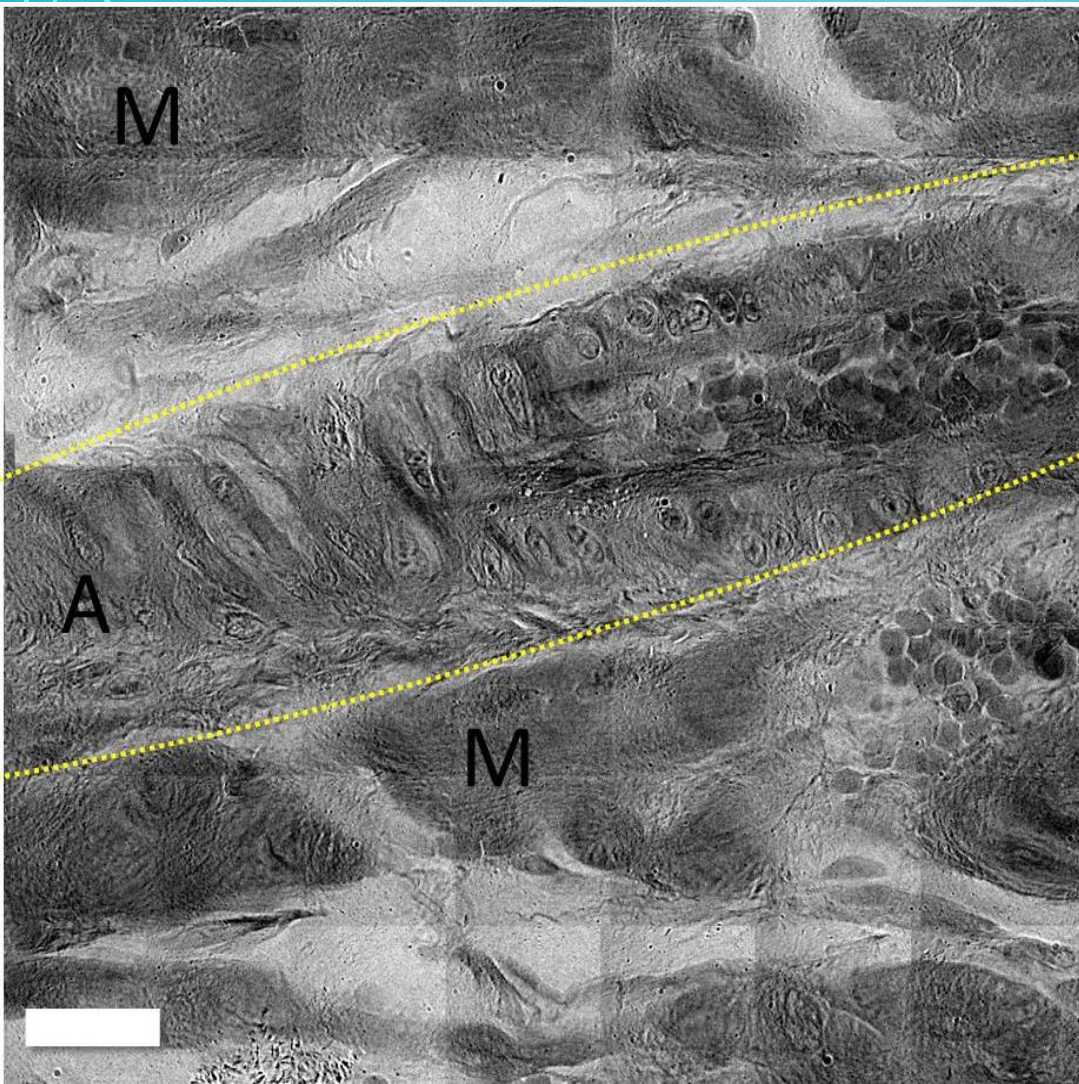
# Solid glioma tumors



# Diffused glioma tumors



# TXM Images of a microvessel





# What do we expect to learn?

- Small differences in microvasculature between different models
- Effect of treatment
- Inhomogeneity in microvasculature with respect to tumor
- Microvasculature with respect to the growth and metastasis of tumor
- Drug development and usage

# There are more problems in brains with neurons (dementia) than vessels (cancer)

- Can X-rays image neurons?
- Can X-rays image blood vessels at the same time?
- Can X-rays study neuro-vascular interaction?
- On dead specimens, of course.



WHY BRAIN?

The  
Economist

JANUARY 6TH - 12TH 2018

The message of protests in Iran

Blue-state Republicans: no right left

Education lessons from Pakistan

The world's worst airports

# The next frontier

When thoughts control machines



# **Mapping the Brain: an Historical Mission for Science and Technology**

- **A fundamental research objective**
- **An effective way to understand and cure brain diseases**
- **Potential for a broad social impact**
- **Important technological byproducts:**
  - **Advanced imaging technologies**
  - **New computational strategies**
  - **Artificial intelligence**

# THE CHALLENGES:



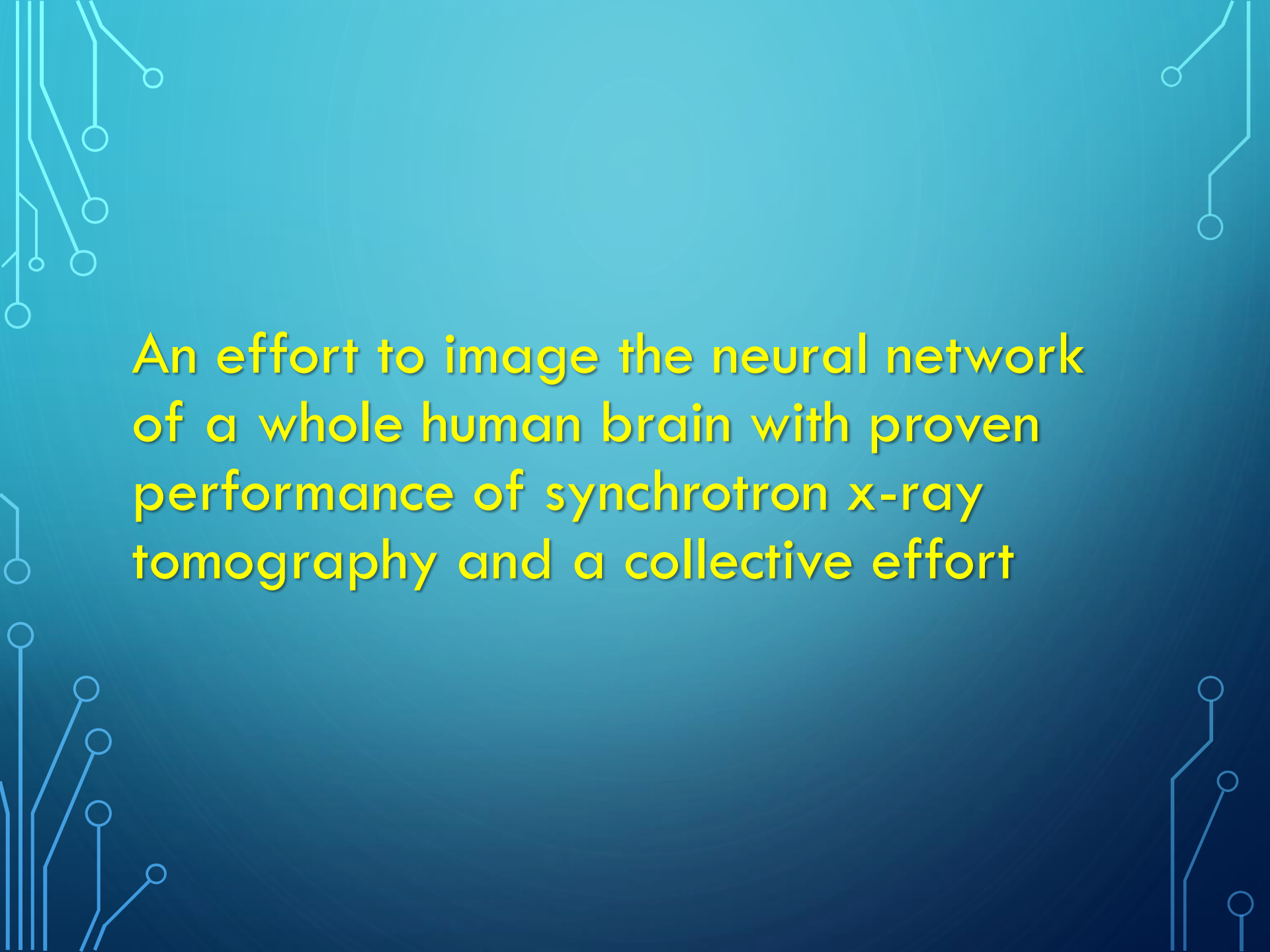
No present technique can image an entire animal brain down to individual synapses with all the required performances:

- High speed and high throughput
- High spatial resolution in 3 D

**...we must do better!**



**Synchrotrons  
for Neuroscience -- an  
Asia-Pacific Strategic  
Enterprise**



An effort to image the neural network  
of a whole human brain with proven  
performance of synchrotron x-ray  
tomography and a collective effort



**SYNAPSE: an international partnership with a milestone research mission**



**First objective:  
mapping the neuron  
connections of an  
entire human brain  
by 2023**

# Mapping the Brain: an Historical Mission for Science and Technology

- A fundamental research objective
- An effective way to understand and cure brain diseases
- Potential for a broad social impact
- Important technological byproducts:
  - Advanced imaging technologies
  - New computational strategies

# IT TAKES DECADES FOR MAPPING ONE HUMAN BRAIN WITH THE PRESENT X-RAY TOMOGRAPHY PERFORMANCES!

**We must accelerate!**

- Parallel image taking at multiple facilities
- Automated specimen sectioning
- Automated specimen mounting and alignment
- High-speed data transfer and sharing for all partners
- New, efficient strategies for reconstruction

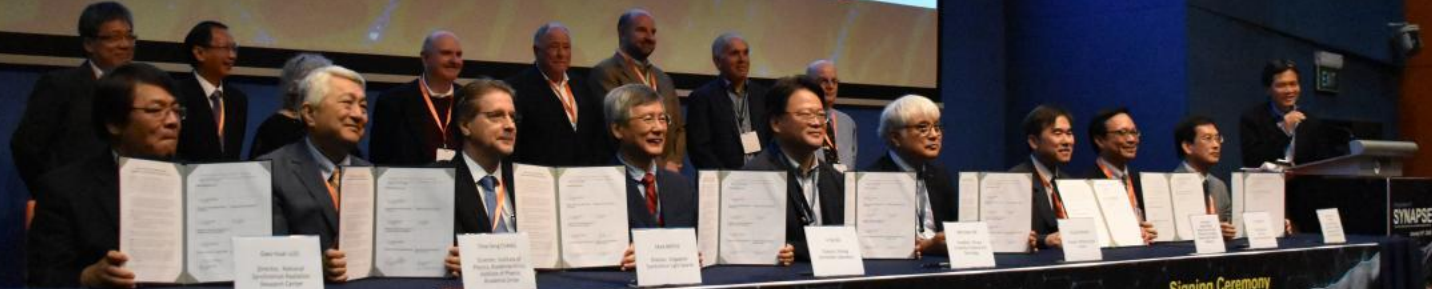
**SYNAPSE is the solution**

# SYNAPSE:

core partners  
with top-level  
synchrotron  
facilities



**In addition to x-ray microtomography facilities, the core partners bring with them a very powerful extended coalition**





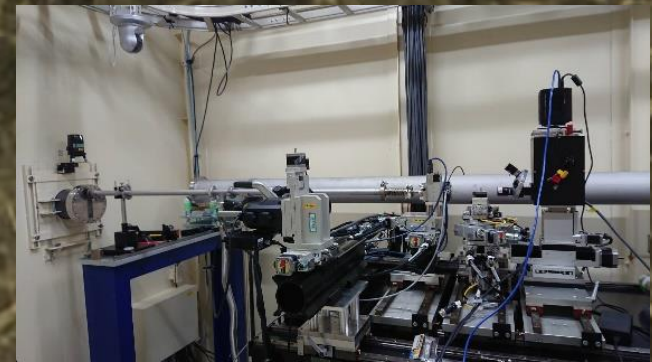
# CONCENTRATE ON 3D IMAGING:

## New radiology techniques combined with other advanced microscopies

- For overall connection mapping: phase-contrast microtomography (0.3  $\mu\text{m}$  resolution in all 3D directions)
- To explore in detail synapses and neuron connections: nanotomography with <10 nm resolution
- To obtain detailed 3D maps of special regions: electron microscopy/continuous sectioning, optical super-resolution microscopy, with nm resolution
- To analyze the whole drosophila brain and large parts of the mouse brain obtaining functional information: Confocal (or Light Sheet) + *FocusClear* imaging
- For single-molecule mapping of drosophila and mouse brains: Super-resolution microscopy with *FocusClear*
- high resolution functional imaging: Raman, IR spectromicroscopy
- Fusing-in low resolution functional imaging, i.e. fMRI
- ...plus other tasks using other frontline techniques



Pohang Light Source II 7C

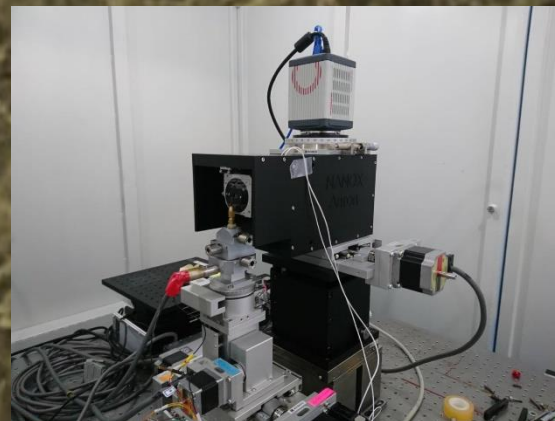


Spring-8 BL32B2



Taiwan Photon Source BL2A

Shanghai Synchrotron  
Radiation Facility BL 09



Singapore Synchrotron  
Light Source PCXT



# SYNAPSE:

core partners  
with top-level  
synchrotron  
facilities

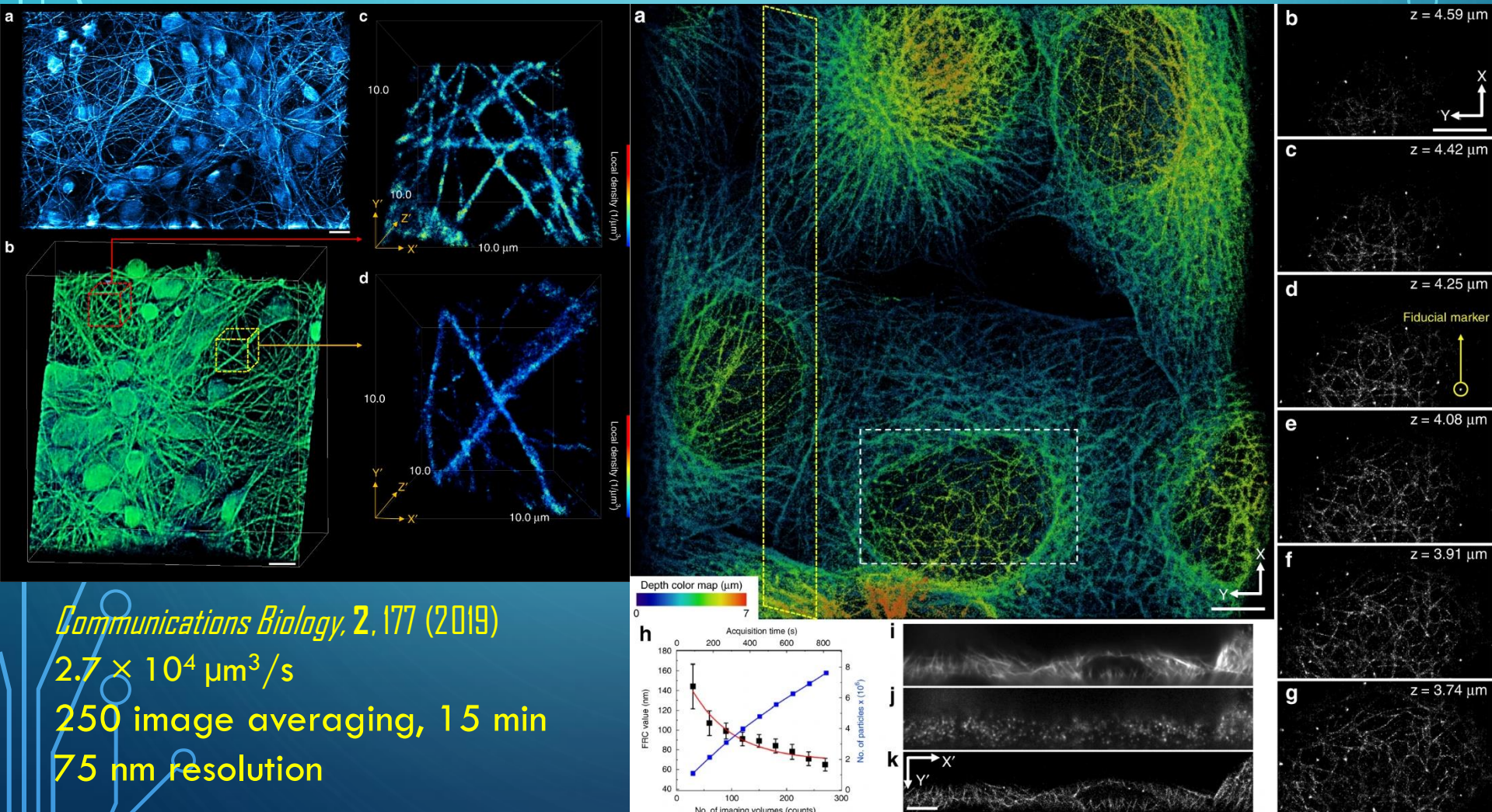


In addition to x-ray microtomography facilities, the core partners bring with them a very powerful extended coalition

# Concentrate on 3D imaging: new radiology techniques combined with other advanced microscopies

- For overall connection mapping: phase-contrast microtomography (0.3  $\mu\text{m}$  resolution in all 3D directions)
- To explore in detail synapses and neuron connections: nanotomography with  $<10$  nm resolution
- To obtain detailed 3D maps of special regions: cryo-electron microscopy with nm resolution
- To analyze the whole drosophila brain and large parts of the mouse brain obtaining functional information: Confocal FocusClear imaging
- For single-molecule mapping of drosophila and mouse brains: Super-resolution microscopy with FocusClear
- ...plus other tasks using other frontline techniques

# Lightsheet Localization super-resolution Microscopy



*Communications Biology*, **2**, 177 (2019)

$2.7 \times 10^4 \mu\text{m}^3/\text{s}$

250 image averaging, 15 min

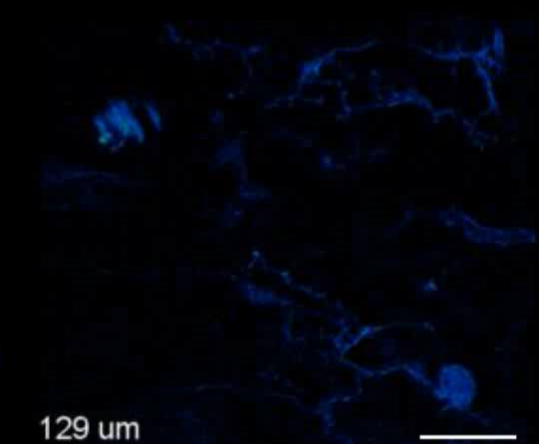
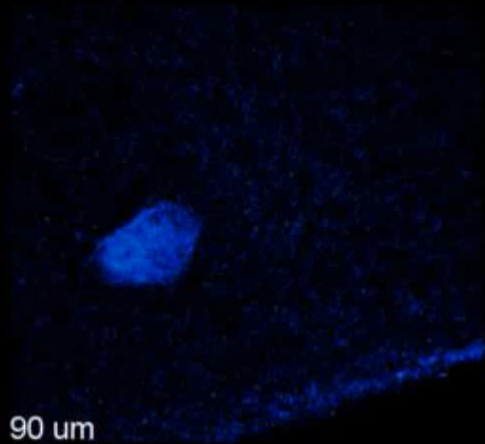
75 nm resolution

# Lattice lightsheet microscopy w/ tissue clearing

LM-CT



LLM-CT



0 um  
Antennal Lobe

90 um  
Lateral Horn

129 um  
Calyx









10 um

ARTICLE

<https://doi.org/10.1038/s41467-019-12715-3>

OPEN

# Rapid single-wavelength lightsheet localization microscopy for clarified tissue

Li-An Chu <sup>1,2,10</sup>, Chieh-Han Lu <sup>2,3,9,10</sup>, Shun-Min Yang<sup>2</sup>, Yen-Ting Liu <sup>3</sup>, Kuan-Lin Feng<sup>4</sup>, Yun-Chi Tsai <sup>3</sup>, Wei-Kun Chang<sup>1</sup>, Wen-Cheng Wang<sup>3</sup>, Shu-Wei Chang <sup>3</sup>, Peilin Chen <sup>3</sup>, Ting-Kuo Lee<sup>2</sup>, Yeu-Kuang Hwu<sup>2</sup>, Ann-Shyn Chiang <sup>1,2,4,5,6,7,8\*</sup> & Bi-Chang Chen <sup>1,3\*</sup>

Optical super-resolution microscopy allows nanoscale imaging of protein molecules in intact biological tissues. However, it is still challenging to perform large volume super-resolution imaging for entire animal organs. Here we develop a single-wavelength Bessel lightsheet method, optimized for refractive-index matching with clarified specimens to overcome the aberrations encountered in imaging thick tissues. Using spontaneous blinking fluorophores to label proteins of interest, we resolve the morphology of most, if not all, dopaminergic neurons in the whole adult brain ( $3.64 \times 10^7 \mu\text{m}^3$ ) of *Drosophila melanogaster* at the nanometer scale with high imaging speed ( $436 \mu\text{m}^3$  per second) for localization. Quantitative single-molecule localization reveals the subcellular distribution of a monoamine transporter protein in the axons of a single, identified serotonergic Dorsal Paired Medial (DPM) neuron. Large datasets are obtained from imaging one brain per day to provide a robust statistical analysis of these

QUESTION AND ANSWER

Open Access

# Q&A: Why use synchrotron x-ray tomography for multi-scale connectome mapping?



Yeukuang Hwu<sup>1\*</sup>, Giorgio Margaritondo<sup>2\*</sup> and Ann-Shyn Chiang<sup>1,3\*</sup>

## Abstract

To understand how information flows and is used in the human brain, we must map neural structures at all levels, providing visualizations similar to those of Google Earth for continents, countries, cities, and streets. Unfortunately, the imaging and processing techniques currently used in connectomics projects cannot achieve complete mapping for the brain of large animals within the timespan of a typical research career. However, feasible improvements in x-ray imaging would change this situation. This Q&A discusses synchrotron x-ray tomography, an exciting new approach for in situ mapping of whole-brain wiring diagrams at multiple levels of spatial resolution.

**In a Presidential Lecture at the 2016 Annual Meeting of Society for Neuroscience, you presented the use of synchrotron x-ray tomography for mapping whole body connectomes in *Drosophila*. Why do we need x-ray imaging for connectomics?**

Recent advances in imaging technologies open the door to mapping a complete wiring diagram of the human brain, a

approach to mapping a small part of the human connectome at least for the foreseeable future.

We sought an alternative to these techniques on deciding that we wanted to map the whole body connectome of *Drosophila* in order to understand the neural basis of its behavior. X-ray microscopy offered advantages in speed and resolution made more apparent and exploitable since its development using modern synchrotron facilities [1, 2]. To meet the challenge of connectome mapping, however, certain performance improvements were required, notably to handle large specimens—the entire animal brain and body—with sufficient image contrast and tunable resolution to detect the fine connections. Overcoming these obstacles took more than a decade, but as a result x-ray imaging is now, in our view, the technique of choice for whole-body connectome mapping.

Many of the advanced x-ray imaging techniques of today are based on synchrotron sources. Synchrotron x-rays offer high brightness with deep-penetration for in situ visualization at a high speed of internal structures within a large tissue—similar to medical computer tomography, but with much better spatial resolution for obtaining multi-level views, ranging from a large brain volume to a single neuron. Alternatively, the in situ

# Brain size

	Volume (mm <sup>3</sup> )	Number of Neurons	brain/body Ratio	Number of Synapse	Density of neurons (/mm <sup>3</sup> )
drosophila	$2 \times 10^{-2}$	$2.5 \times 10^5$ ( $1.3 \times 10^5$ )			$1.25 \times 10^7$
mouse	450	$7.1 \times 10^7$	1:40	$10^{11}$	$1.57 \times 10^5$
marmoset		$6.4 \times 10^8$			
human	$1.2 \times 10^6$	$8.6 \times 10^{10}$	1:50	$10^{14-15}$	$7.2 \times 10^4$

# Big Brain, Big Data

## TECHNOLOGY FEATURE

### BIG BRAIN, BIG DATA

Neuroscientists are starting to share and integrate data — but shifting to a team approach isn't easy.



Diffusion magnetic resonance imaging is just one of many data types that researchers are working out how to handle to bring the brain into focus.

BY ESTHER LANDHUIS

As big brain-mapping initiatives go, Taiwan's might seem small. Scientists there are studying the humble fruit fly, reverse-engineering its brain from images of single neurons. Their efforts have produced 3D maps of brain circuitry in stunning detail.

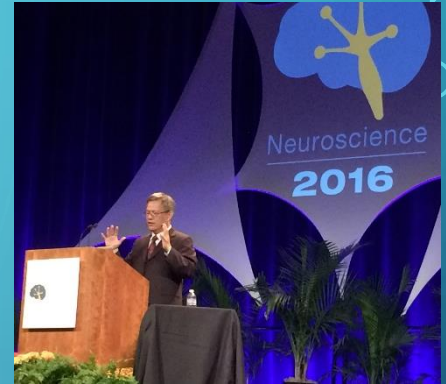
leader Ann-Shyn Chiang, a neuroscientist at the National Tsing Hua University in Hsinchu City, Taiwan — and that's not even half of the nerve cells in the *Drosophila* brain. Using the same protocol to image the 86 billion neurons in the human brain would take an estimated 17 million years, Chiang reported at the meeting. Other technologies are more tractable. In

principal investigator for the Human Connectome Project (HCP), a US-government-funded global consortium that published the brain map. So it goes in the world of neurobiology, where big data is truly, epically big. Despite advances in computing infrastructure and data transmission, neuroscientists continue to grapple with their version of the 'big data' revolution that

BY ESTHER LANDHUIS

As big brain-mapping initiatives go, Taiwan's might seem small. Scientists there are studying the humble fruit fly, reverse-engineering its brain from images of single neurons. Their efforts have produced 3D maps of brain circuitry in stunning detail.

Researchers need only a computer mouse and web browser to home in on individual cells and zoom back out to intertwined networks of nerve bundles. The wiring diagrams look like colourful threads on a tapestry, and they're clear enough to show which cell clusters control specific behaviours. By stimulating a specific neural circuit, researchers can cue a fly to flap its left wing or swing its head from side to side — feats that roused a late-afternoon crowd in November at the annual meeting of the Society for Neuroscience in San Diego, California.



When it comes to the brain, 'complete' can be a moving target. But so, too, is the neuroscience toolset. During his Society for Neuroscience talk, Chiang lamented that it's taken ten years to map half the fly brain. Working with physicists at Taiwan's Academia Sinica, Chiang's team has started to use a technique called synchrotron X-ray tomography to boost data-acquisition speed dramatically.

"It took less than 10 minutes to image a fly brain containing thousands of Golgi-stained single neurons," says Chiang, whose crew is now trying the method in mice and pigs. They plan to integrate confocal and X-ray images on a single platform from which scientists can download data. "With synchrotron X-ray imaging, mapping the human connectome at single-neuron resolution is now more realistic," Chiang says. How easy it will be to meld the maps with other data remains to be seen. ■

Esther Landhuis is a freelance science writer in the San Francisco Bay Area of California.

Nature

<http://www.nature.com/nature/journal/v541/n7638/full/541559a.html#references>

Nature blogs <http://blogs.nature.com/naturejobs/2017/01/26/new-neuroscience-tools-for-team-science-in-big-data-era/>

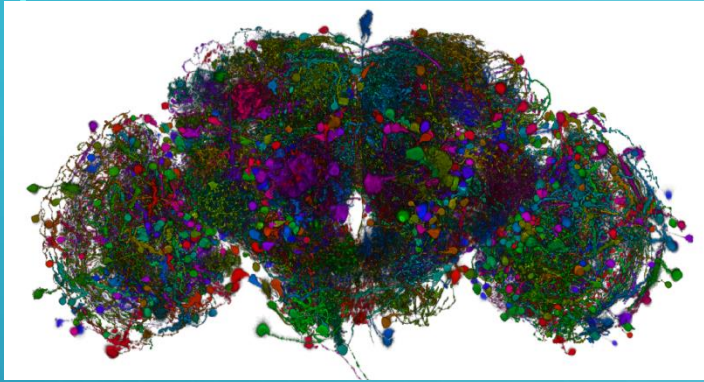
Scientific American <https://www.scientificamerican.com/article/neuroscience-big-brain-big-data/>



# Small Brain, Big Data

## Fluorescence confocal microscopy on connectome mapping

- Drosophila Brain: ~135,000 neurons
- Resolution : 500 nm
- Data amount : ~10 TB
- Time to complete : ~20 year/10 microscopes
- Human brain: 85 billion neurons
- Time to complete : ~17,000,000 年



500  $\mu\text{m}$   $\times$  300  $\mu\text{m}$   $\times$  200  $\mu\text{m}$

## 3D X-ray data

- Single fly brain: ~100GB
- Single mouse brain: ~360TB
- Complete the map with 100 brains: ~36 PetaByte (PB)!
- Human/mouse brain volume: 2500 !

**The New York Times** National Edition  
VOL. CXLV No. 30,284 NEW YORK, TUESDAY, DECEMBER 14, 2010 \$2.00

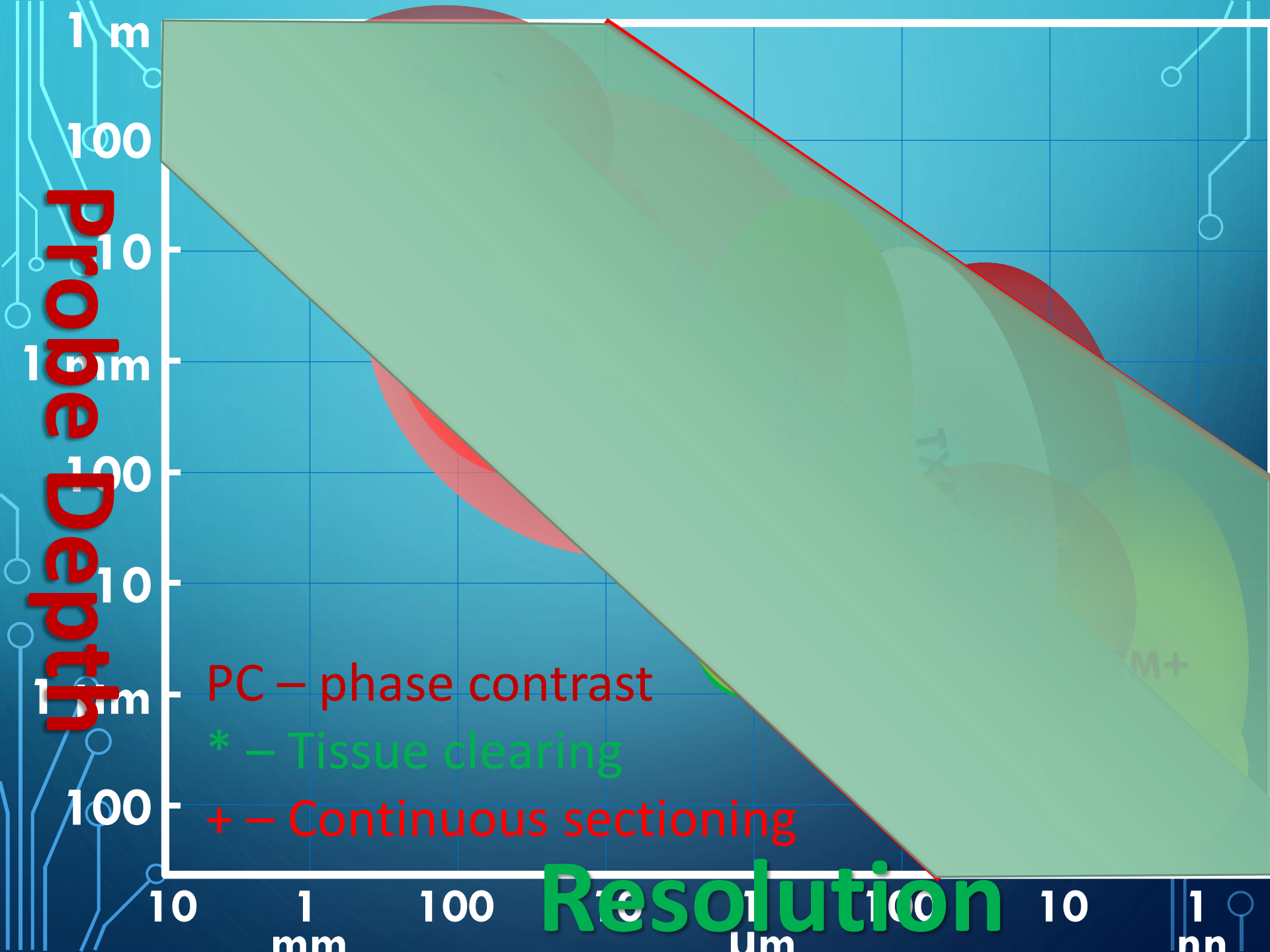
**Decoding the Human Brain, With Help From a Fly**

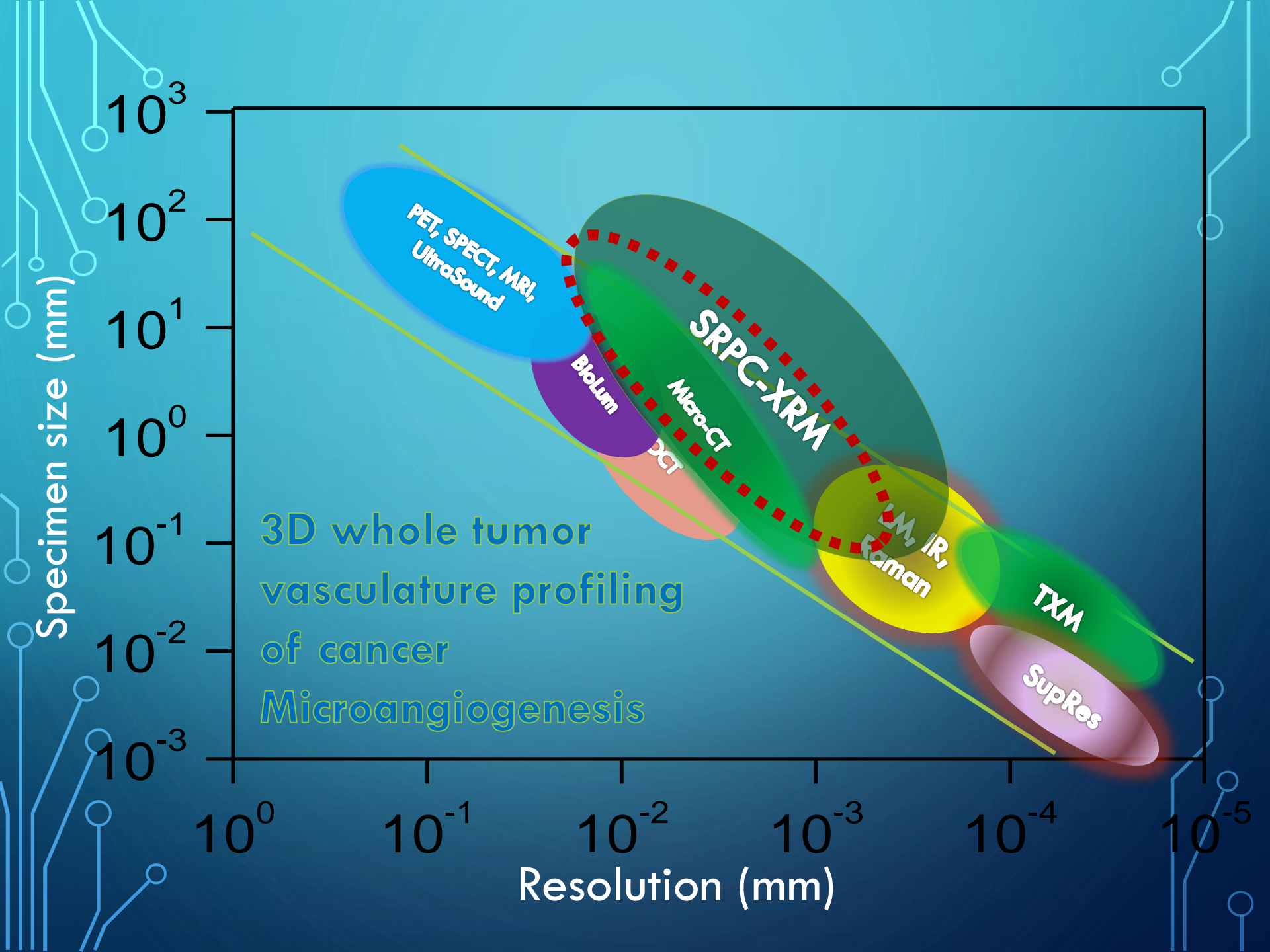
**New Atlas of the Fruit Fly Brain**

**Decoding the**

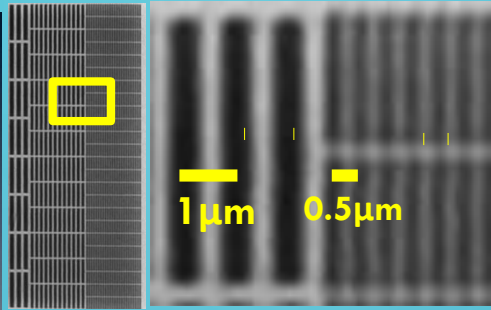
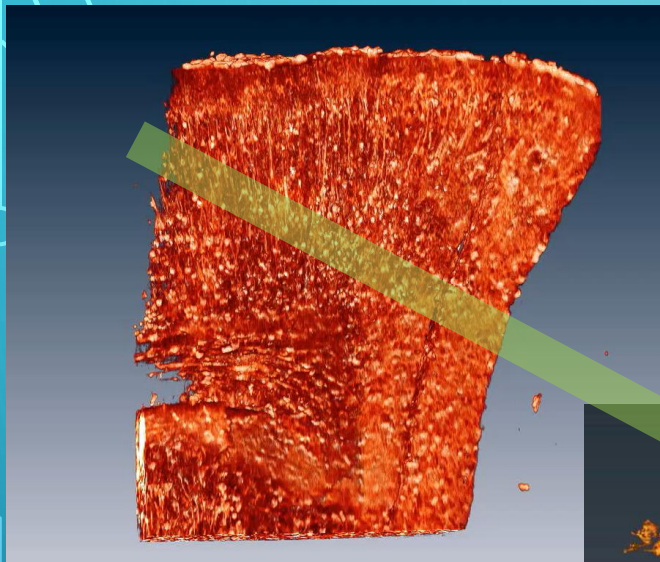
By **NICHOLAS WADE**

Taiwanese researchers have managed to bar code some 16,000 of the 100,000 neurons in a fruit fly's brain and to reconstruct the brain's wiring map.

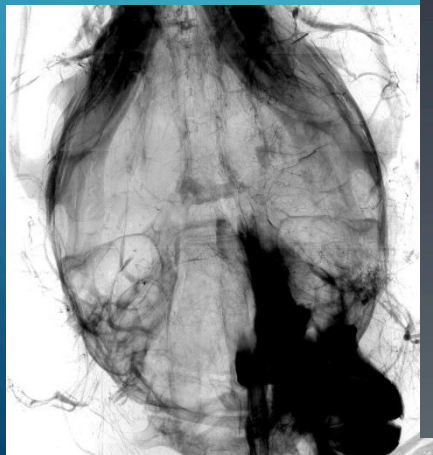
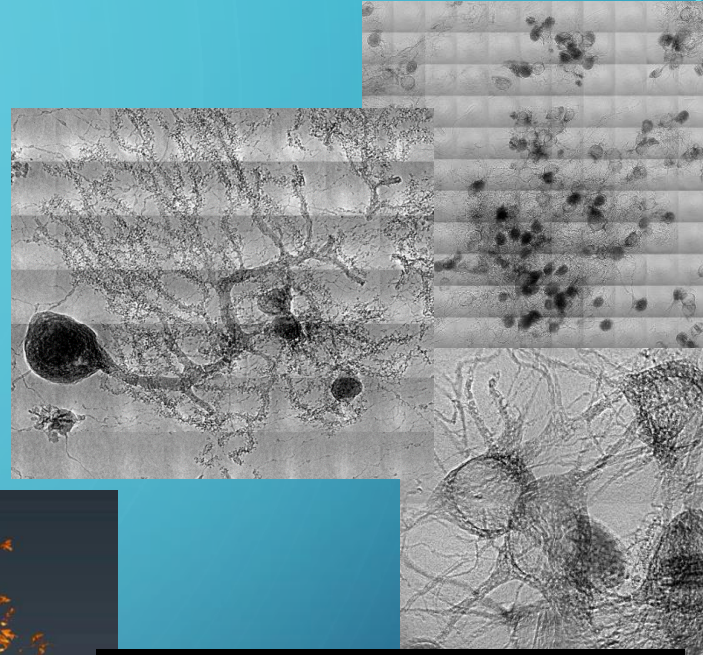




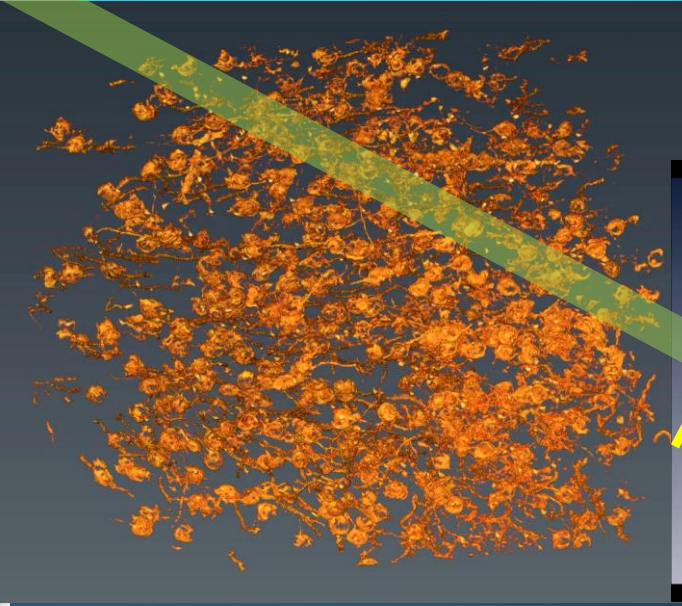
# X-ray imaging — From Animals, Organs, Neurons to Synapses



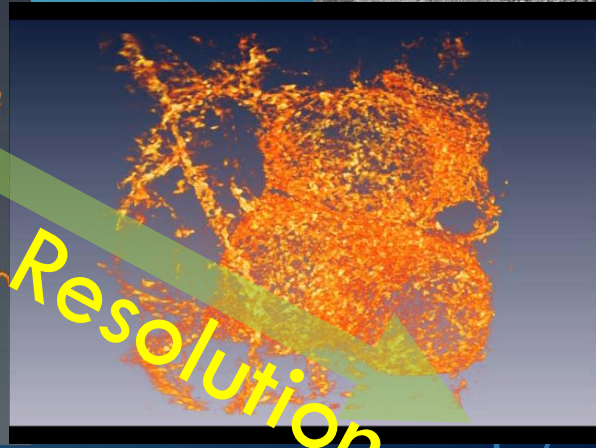
< 0.5 μm resolution (μCT)



400 nm



Mouse brain



Resolution

20 nm



3D imaging with AXON

A

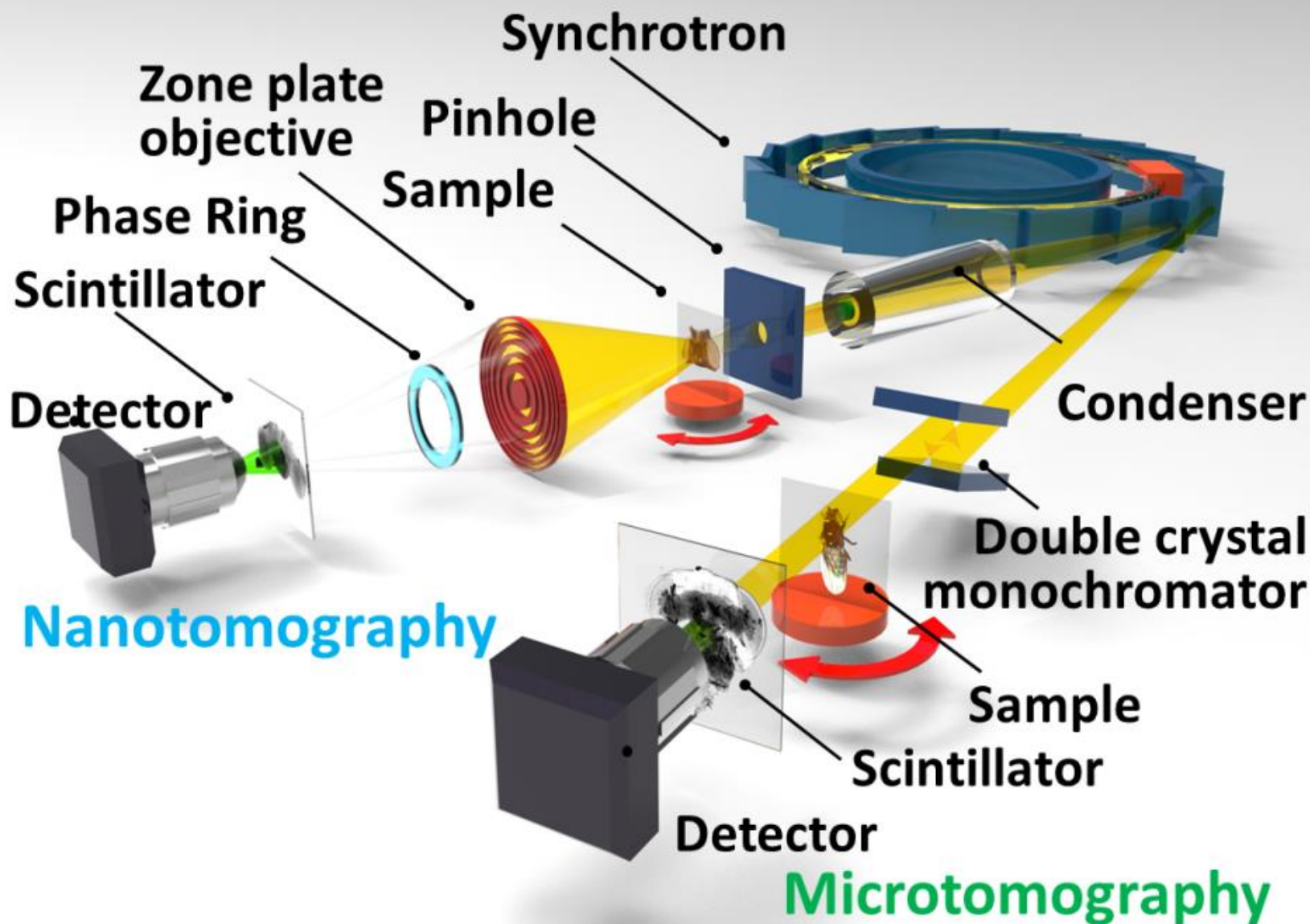
X-ray

O

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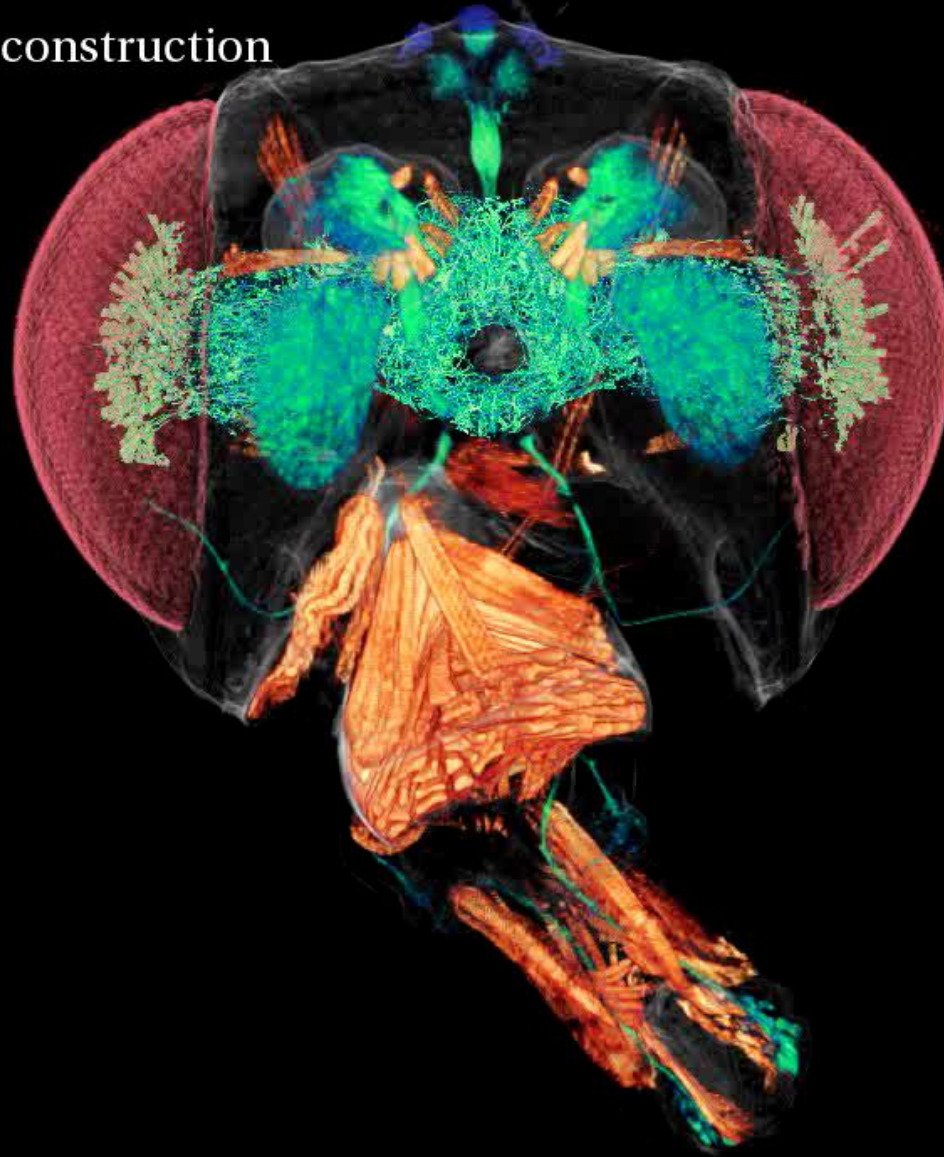
# AXON Implementation: TLS, APS, PLS-II, TPS..



# Whole brain imaging without sectioning



X-ray Tomographic Reconstruction



# Merging Different Brains

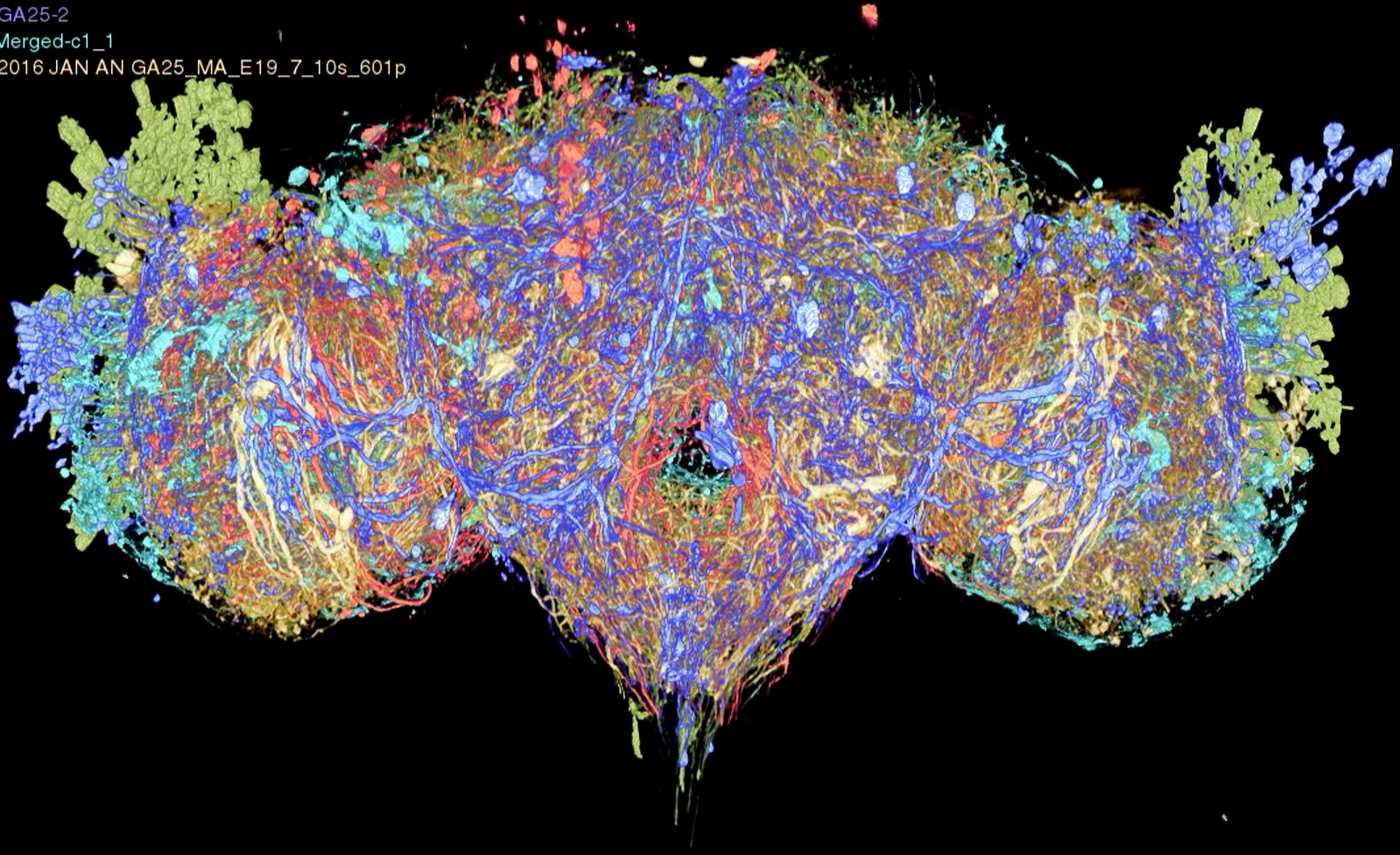
160318-GA25W\_16Mar\_4

20151202-GA25-1

20151202-GA25-2

2016SEP-Merged-c1\_1

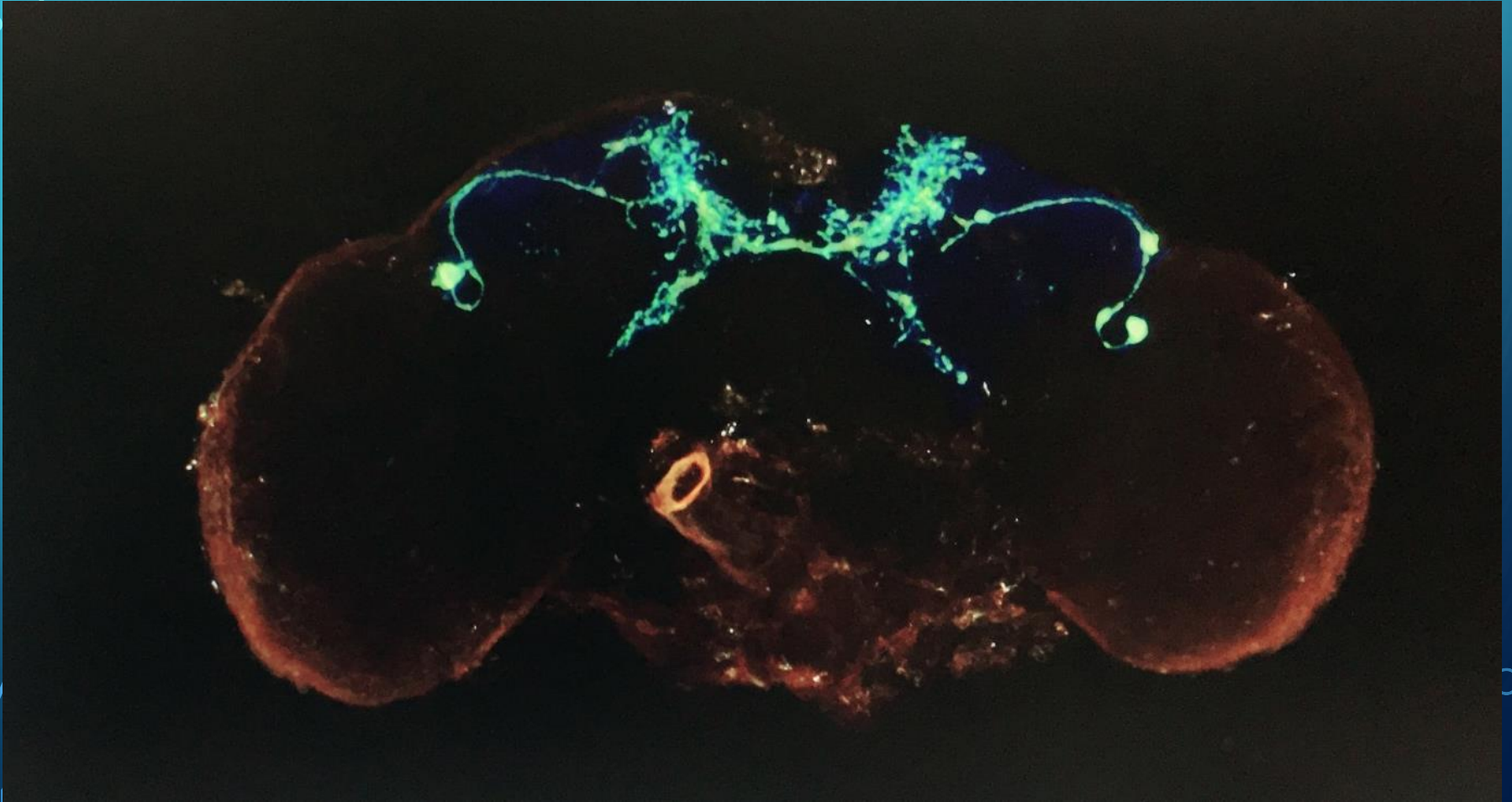
20160308-2016 JAN AN GA25\_MA\_E19\_7\_10s\_601p





# DAL neurons

enzmet/gold toning/gold enhance



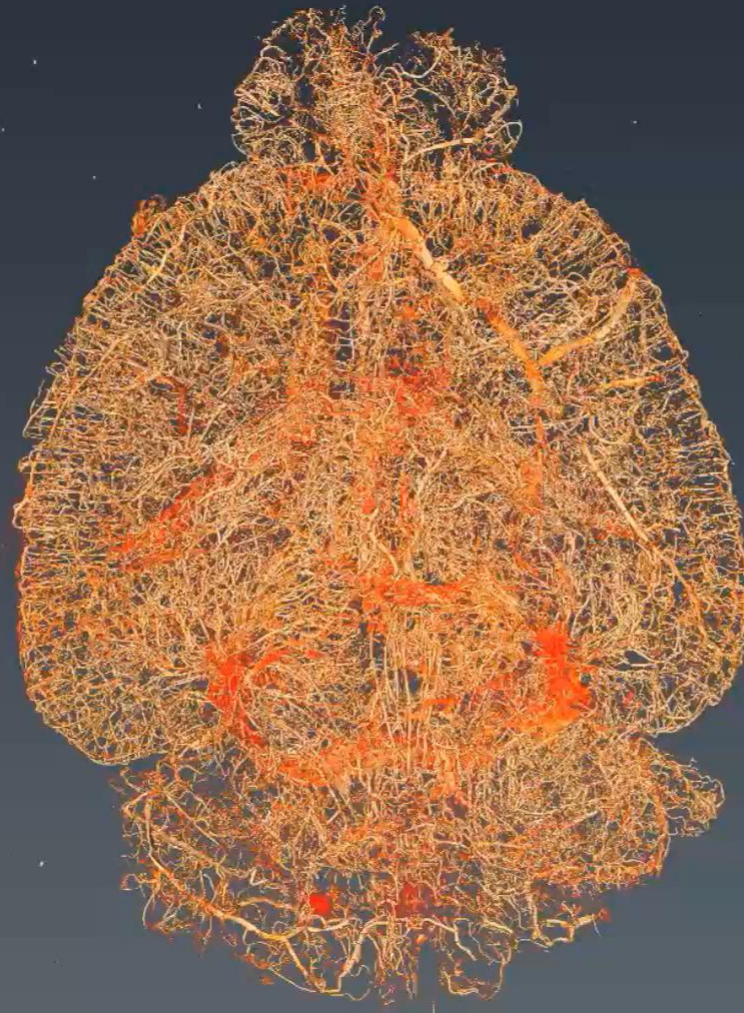
# Large tissue, high resolution and speed

Whole-brain blood vessel reconstruction  
in mouse

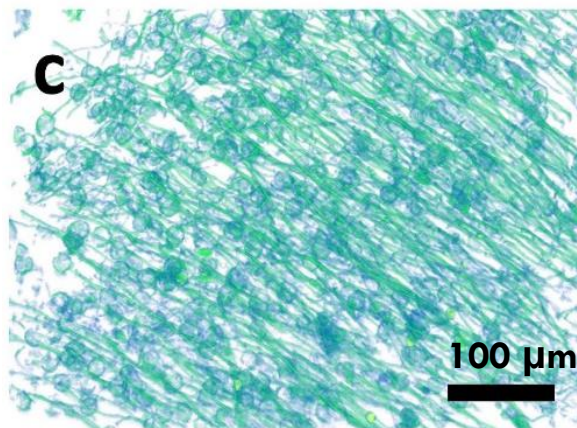
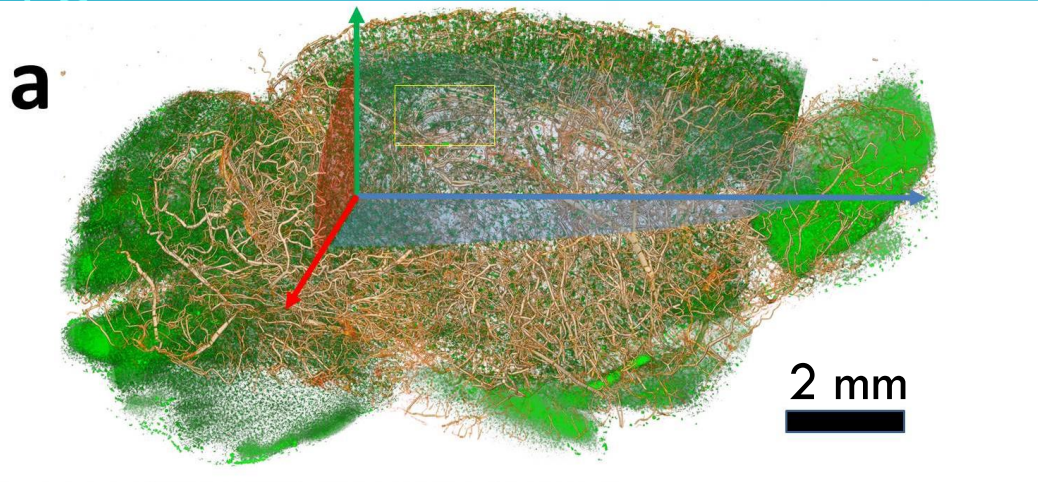
Res. < 0.5  $\mu\text{m}$

Size > 400 mm

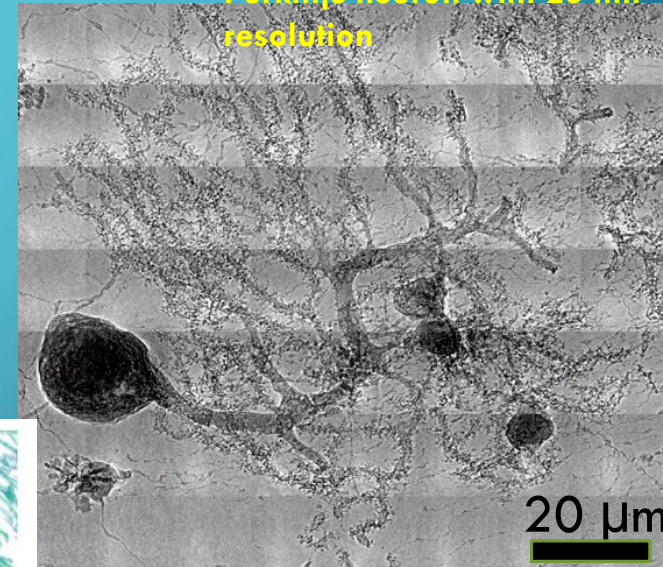
Time < 10 min



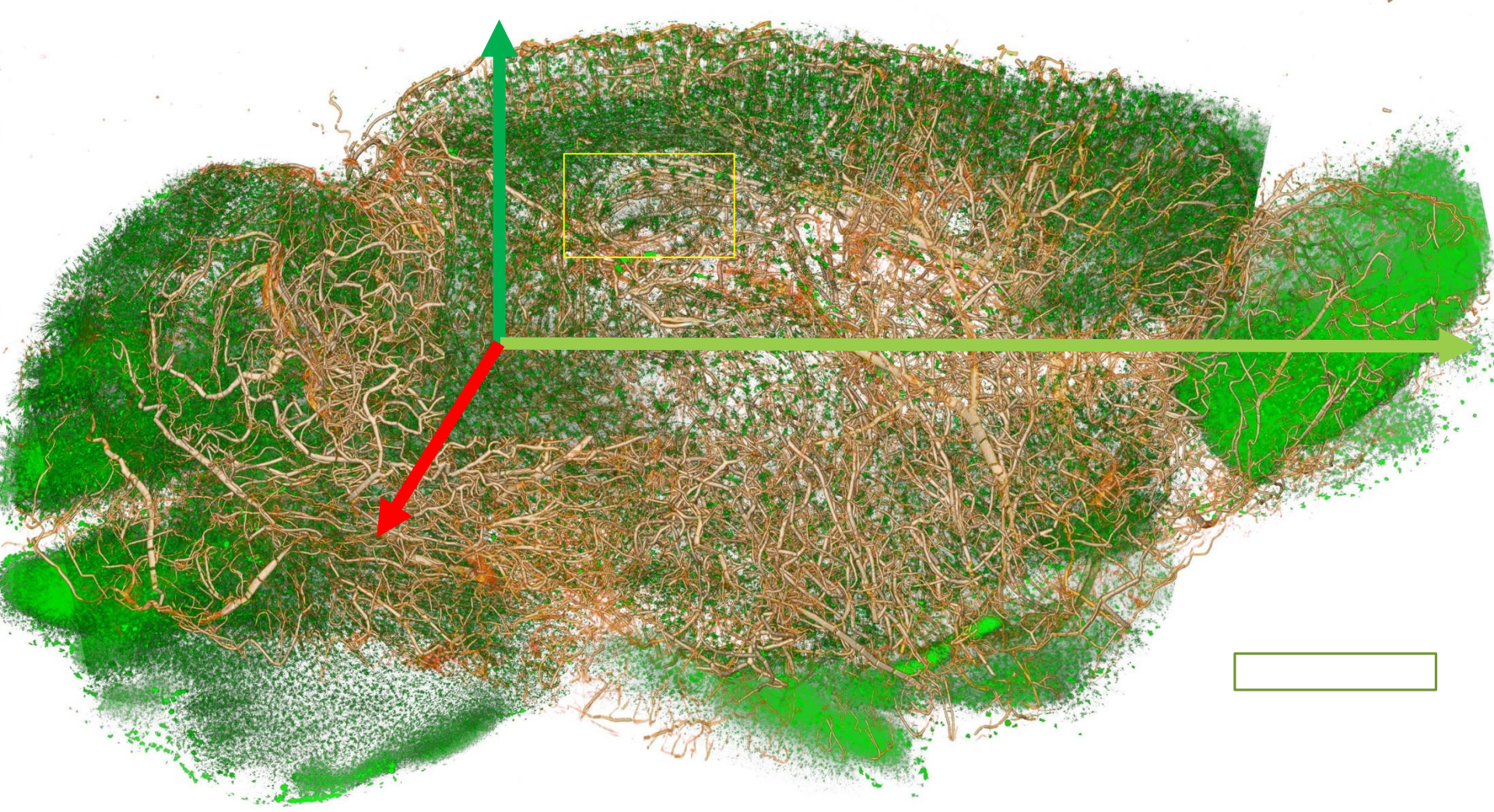
# AXON on Mouse Brains

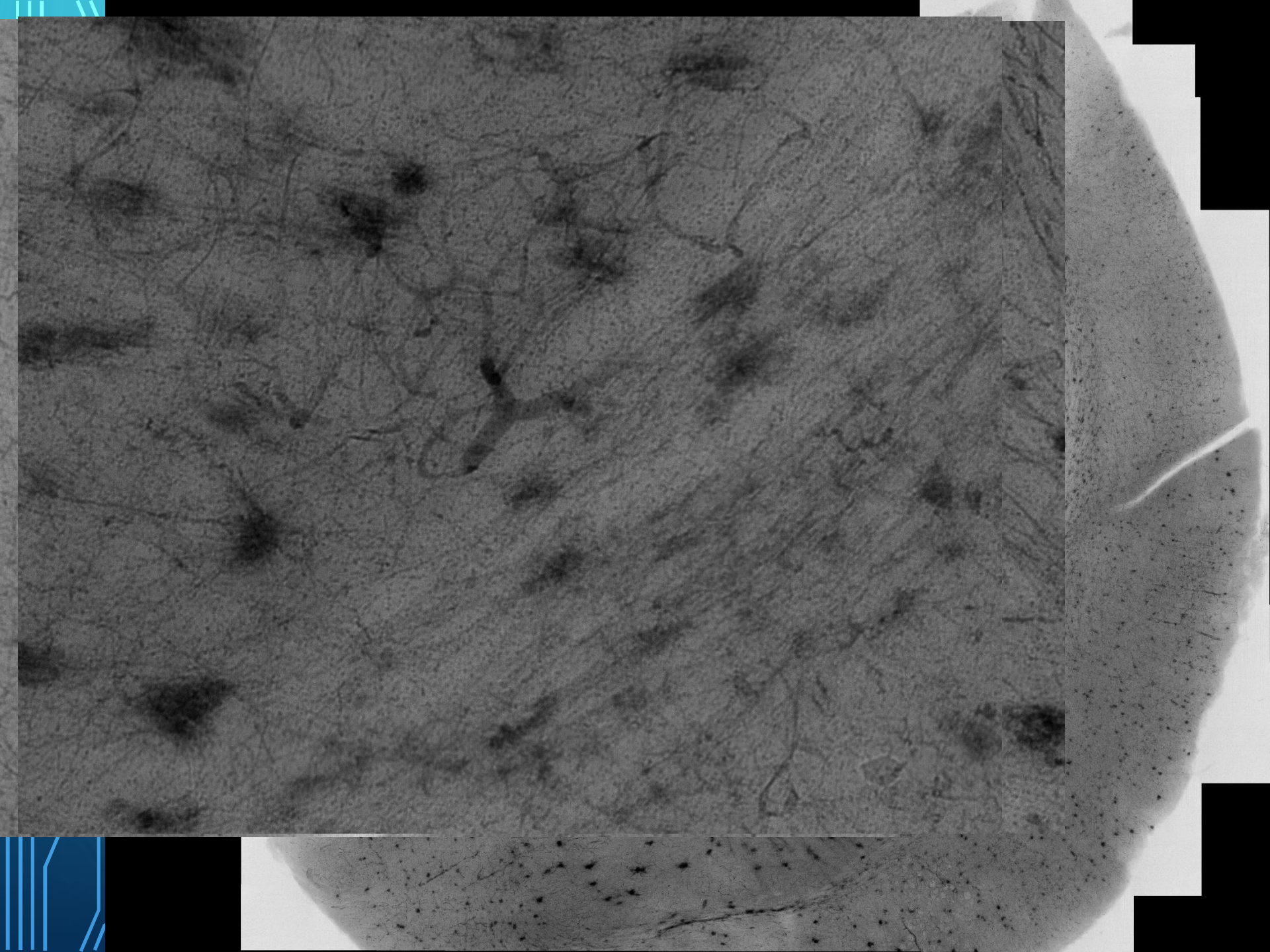


X-ray imaging of a mouse Purkinje neuron with 20 nm resolution

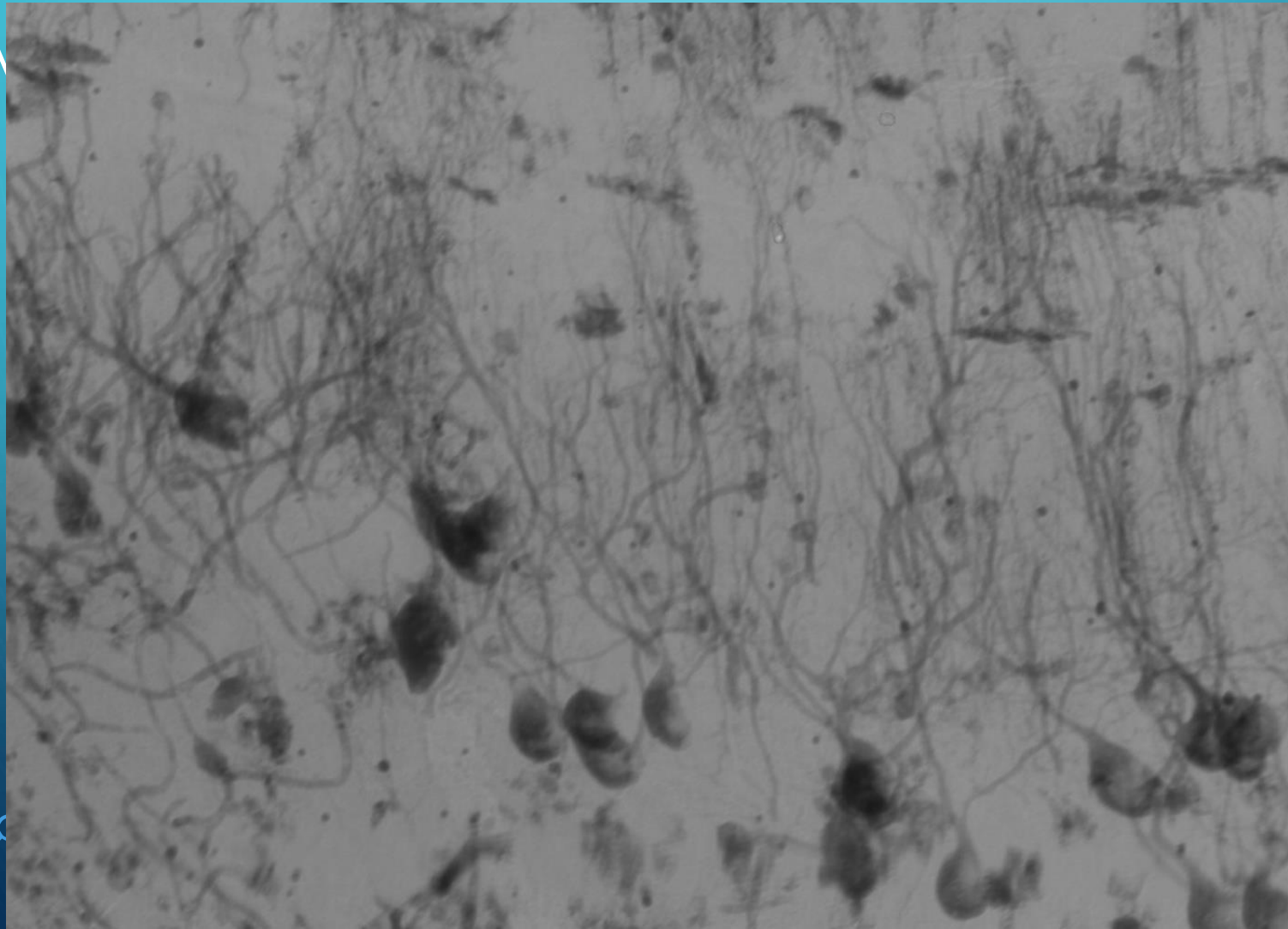


X-ray imaging of a whole mouse with 0.5  $\mu\text{m}$  resolution. Tomography image shows blood vessels (golden) and neurons (green)

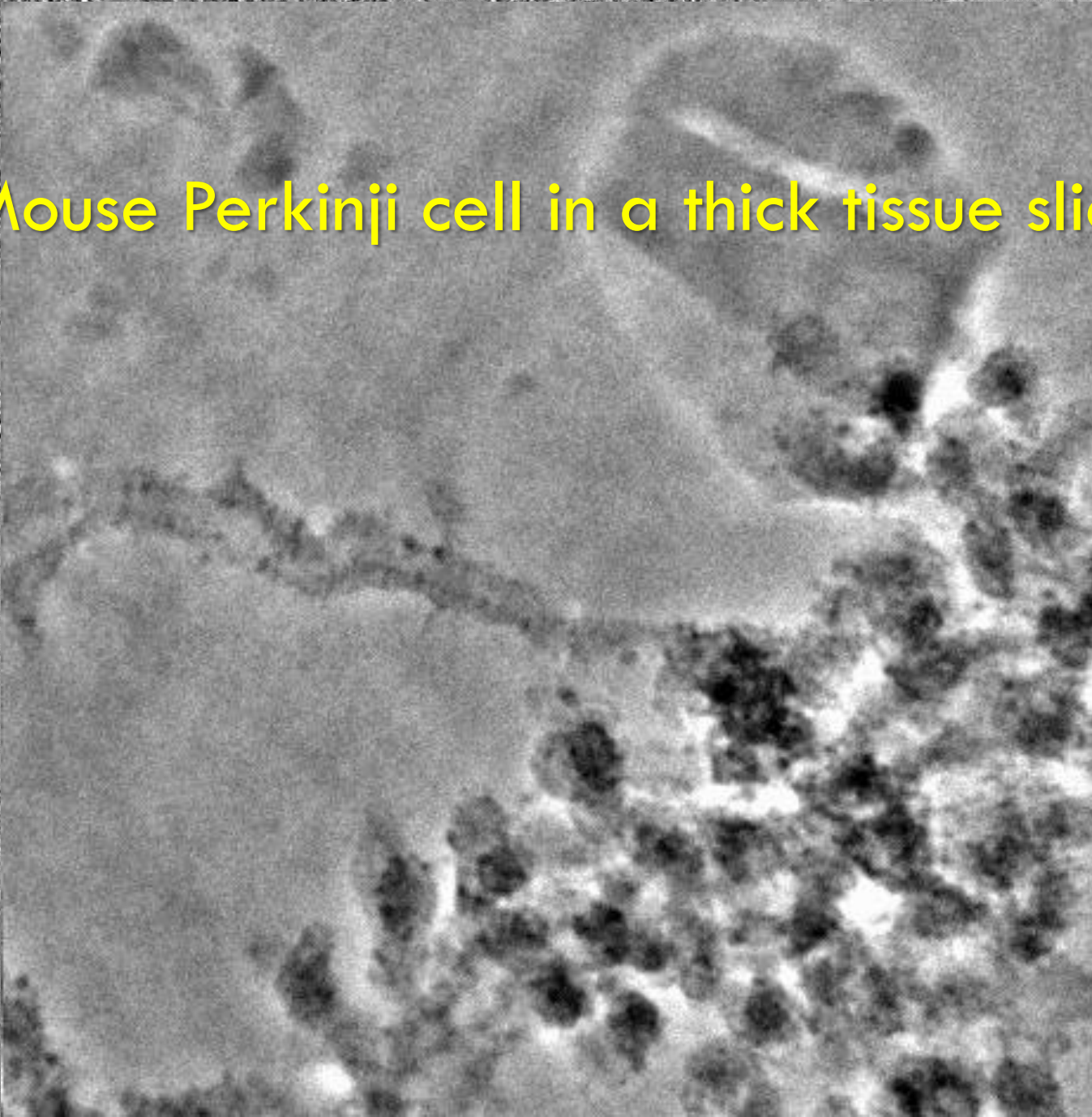




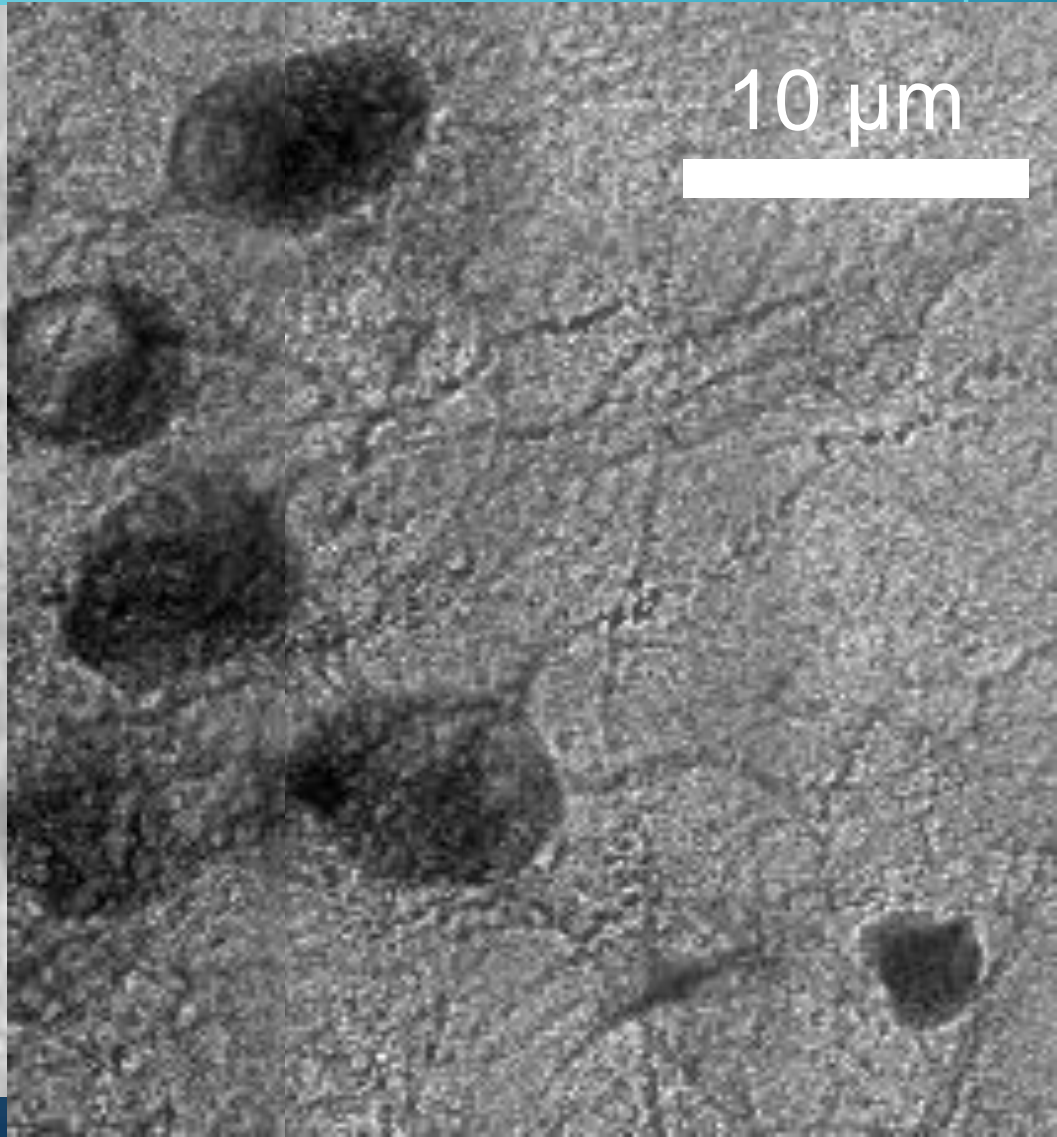
# Recent progress – whole mouse brain imaging



Mouse Perkinji cell in a thick tissue slice

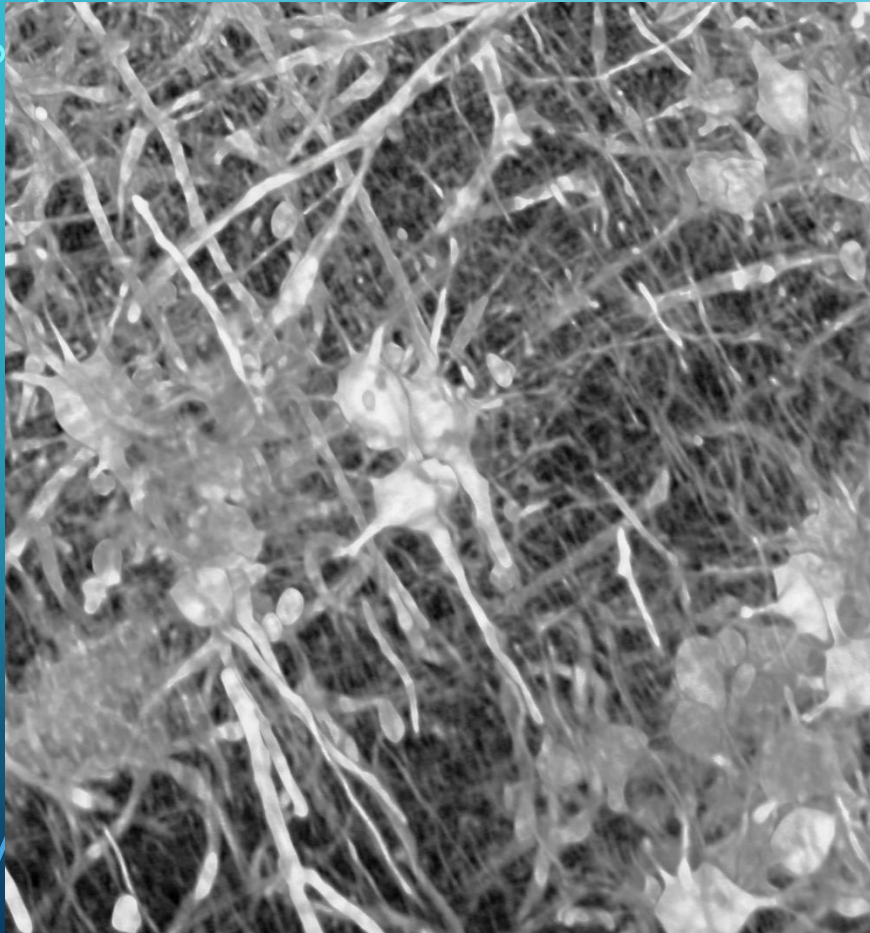


# Same specimen, different resolution





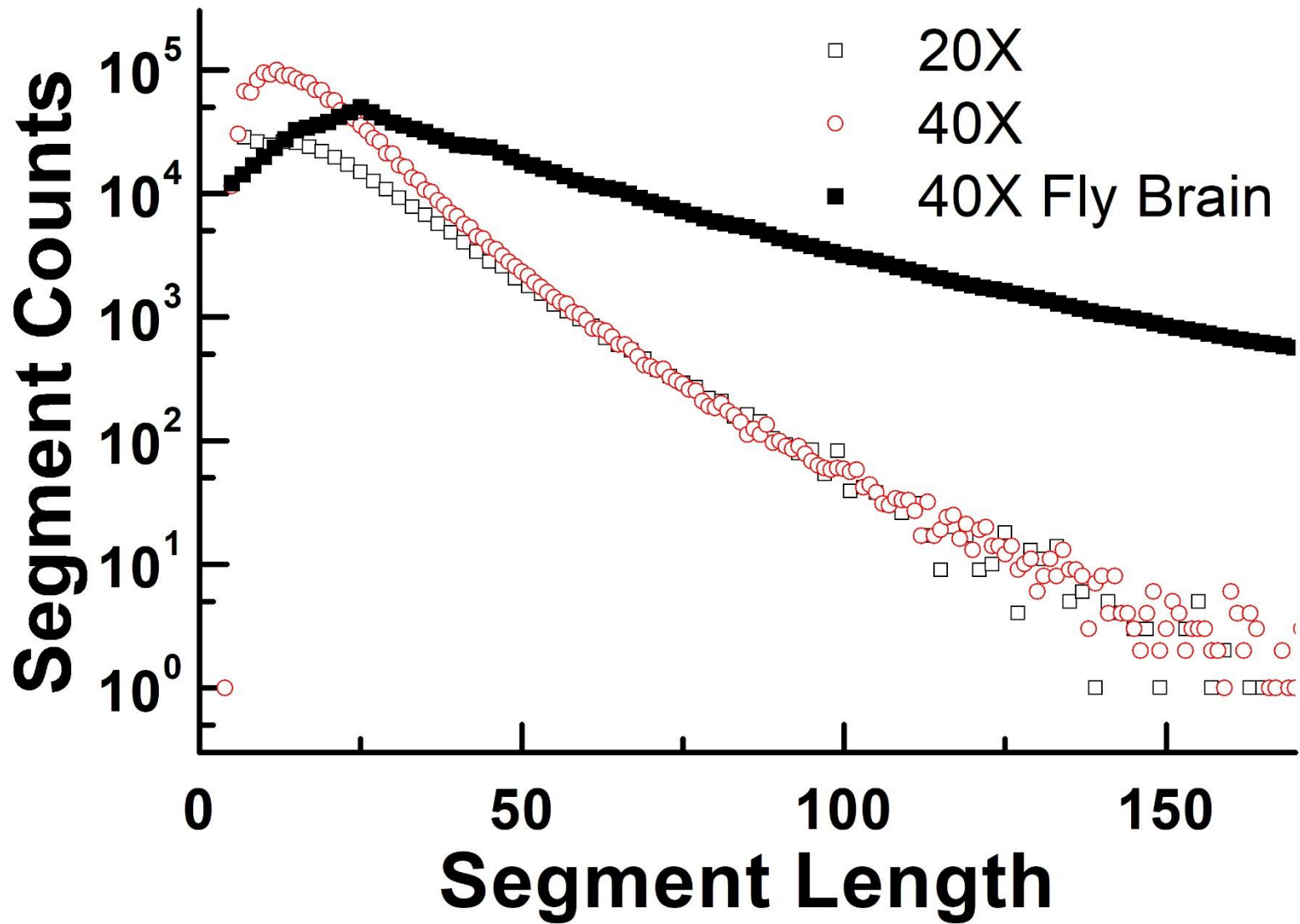
# Same specimen, different resolution



20x, NA = 0.75, resolution  $\sim 0.5 \mu\text{m}$

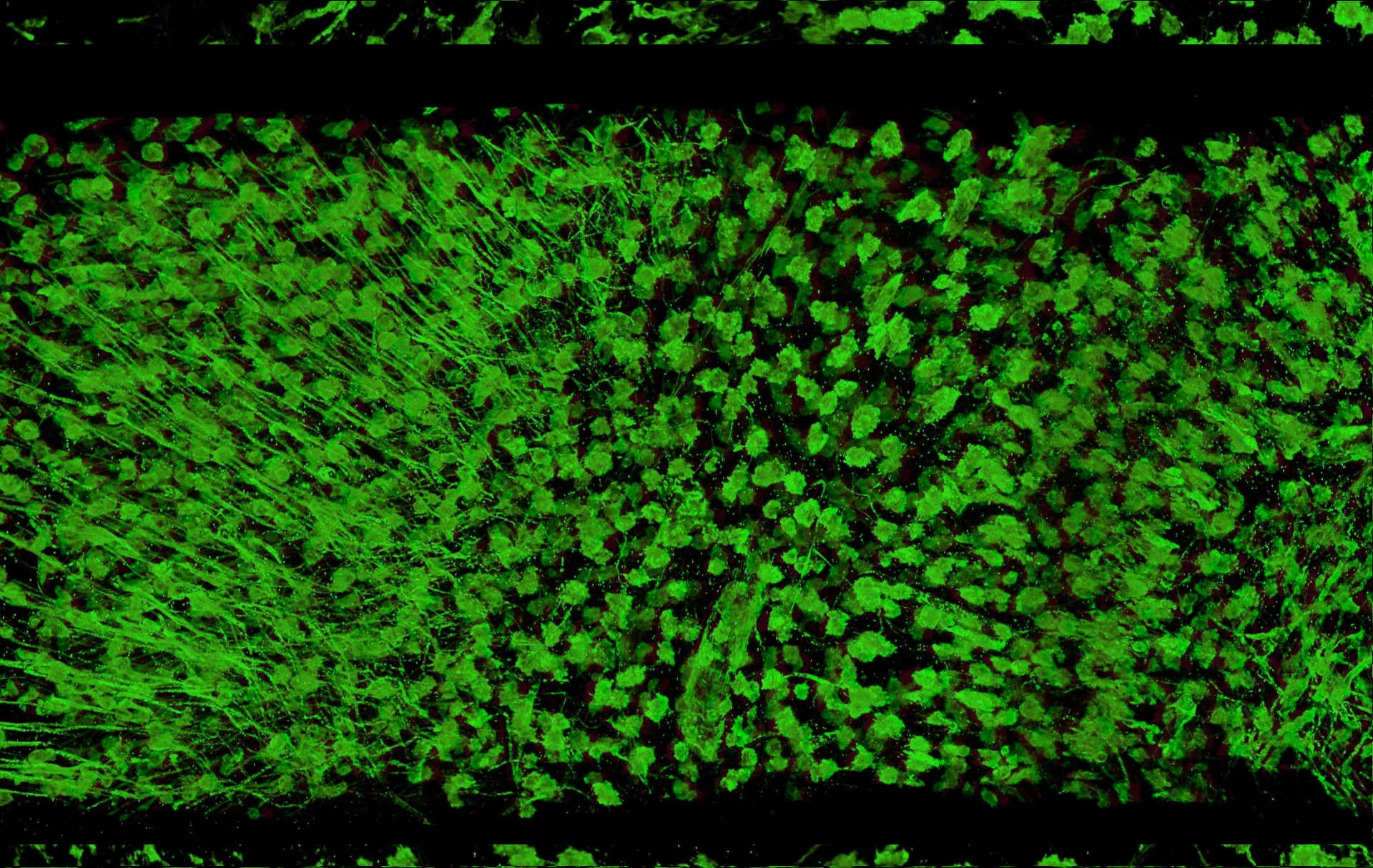


40x, NA = 0.95, resolution  $\sim 0.3 \mu\text{m}$

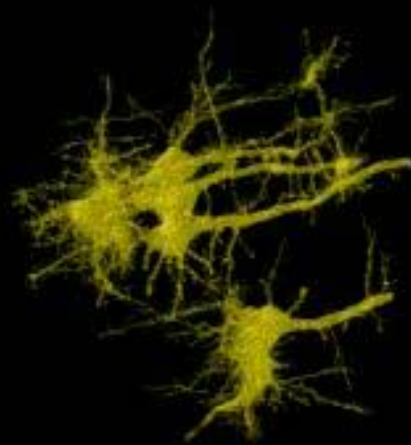




And speed..



# Mouse brain, $(0.8 \mu\text{m})^3$

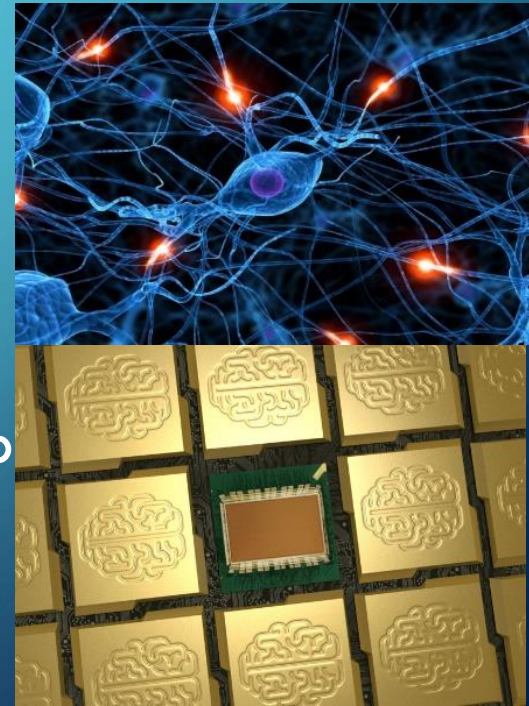


# Technology impacts — Beyond connectome

- X-ray imaging: phase contrast radiology and tomography, nanotomography
- Super-resolution imaging
- 3D continuous sectioning
- 3D spectromicroscopy
- 3D pathology
- Big Data analysis
- Neuromorphic computing & AI

# Brain inspired computation, neurosynaptic computing, neuromorphism and artificial brains

- Power consumption
  - 86 billion neurons, ~1000 synapses per neuron
  - >trillion transistor, >10 GW
  - Human brain 20 W!
- Simulate Brain from neuron (HBP)
- Neurosynaptic computer using neuromorphic chip
- Artificial Intelligence



# Flybrain Simulators

- 130,000 neurons
- Build with real connectome map
- With input and output terminals for simulations
- Functional information included in the simulations
- Understand how fruit fly brain function (with a super computer)
- Better algorithm and computer architecture for AI
- Simulate and understand brain diseases, and the cure



# Next steps:

- Sub-10 nm imaging
- Multi-modality imaging
- X-ray molecular/functional imaging
- X-FEL applications?
- Laser wavefront accelerators?

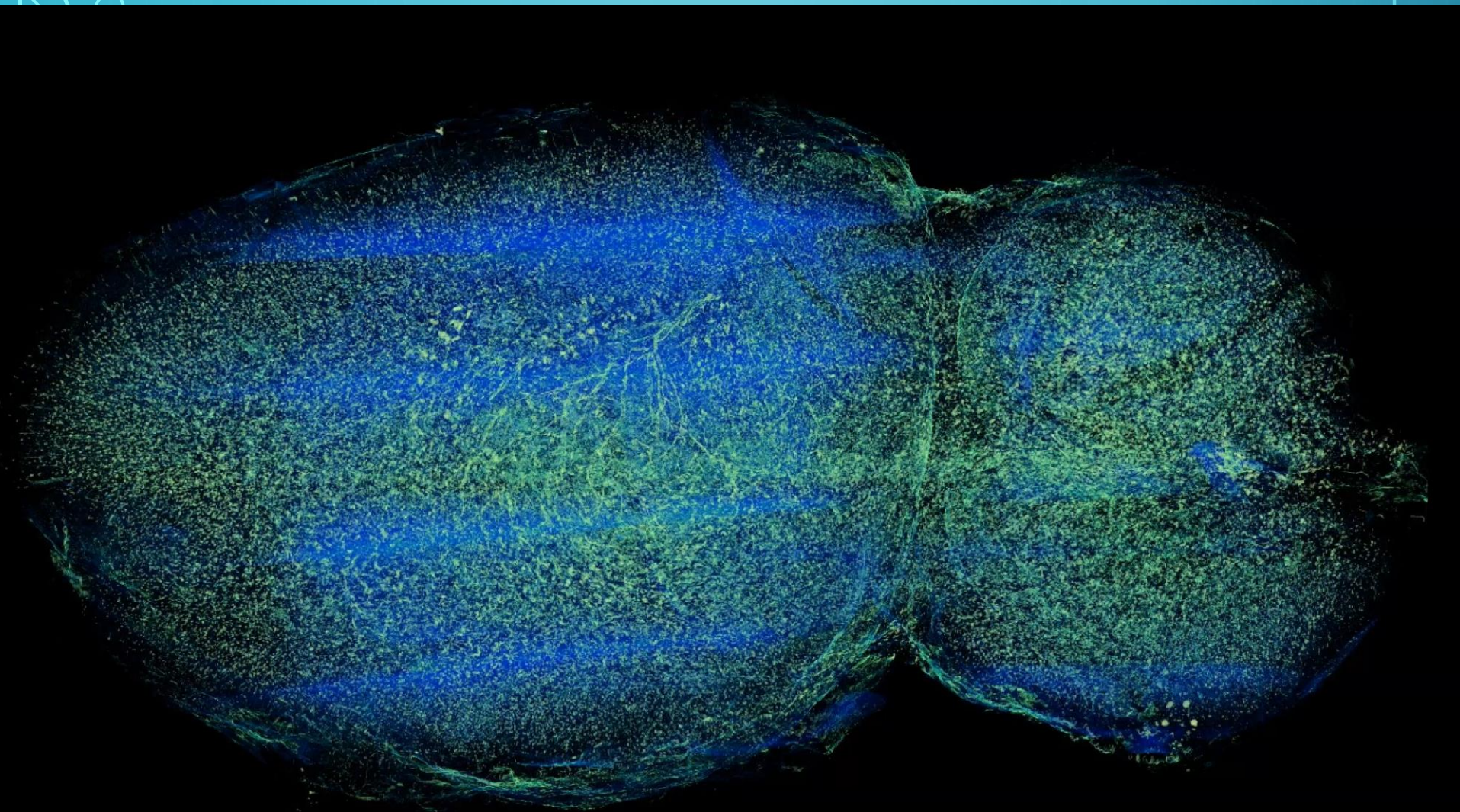
# Toward Sub-10 nm

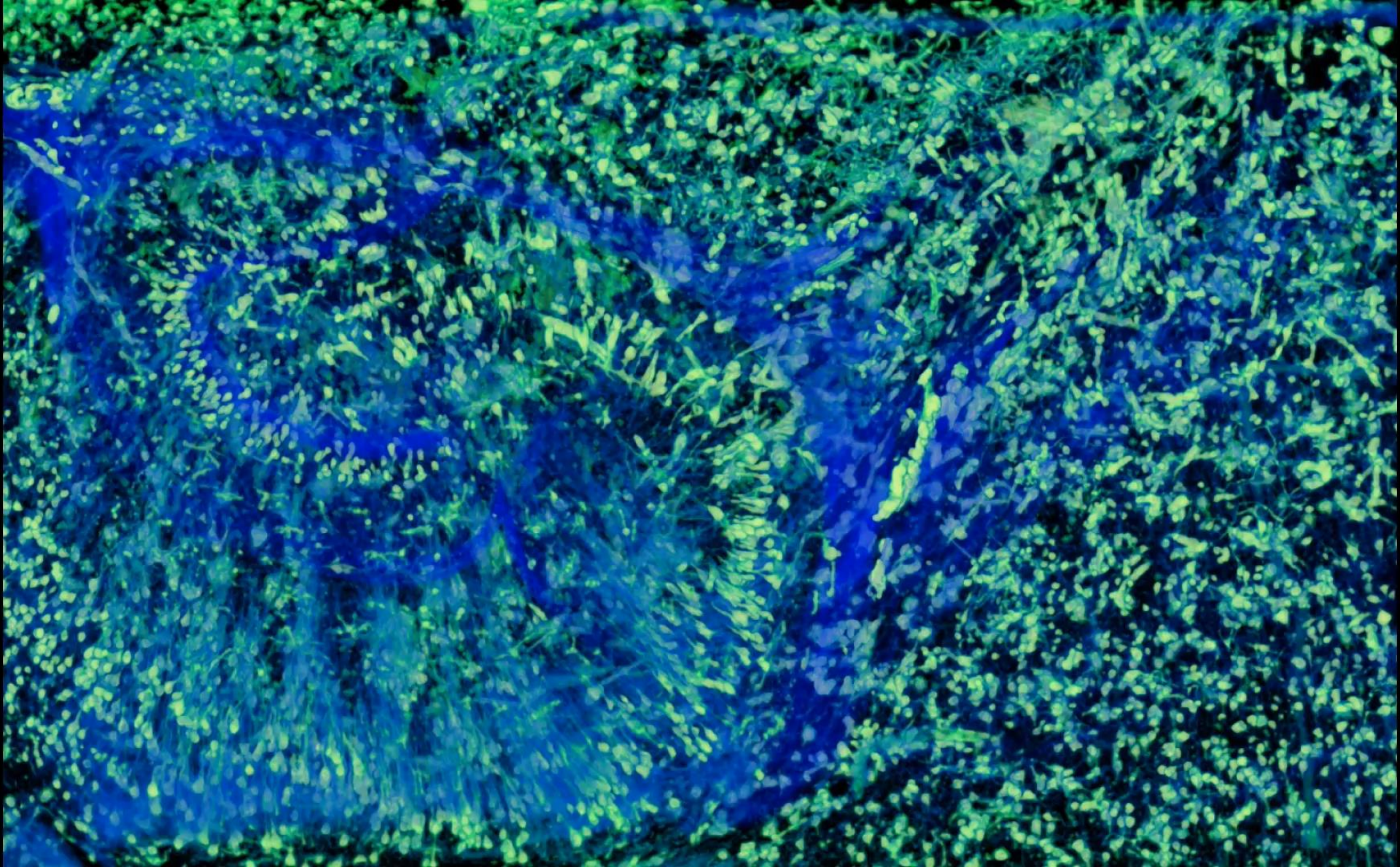
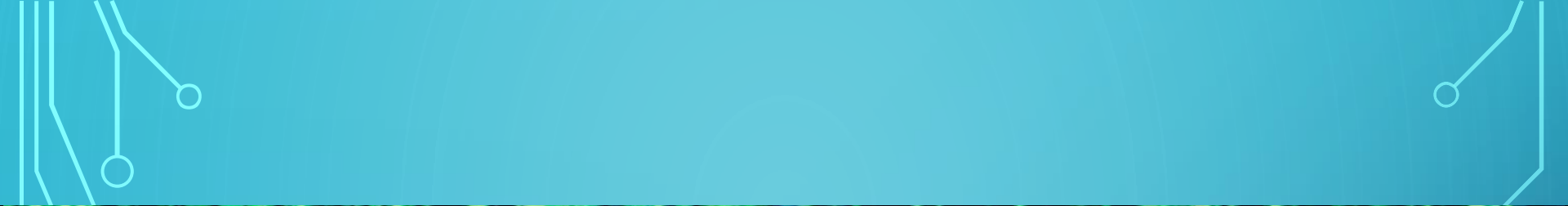
- Must be an international effort
- Mobilization of resources and user base
- Focus on novel configurations and applications
- A consortium of beamlines at NSRRC, SPring-8, PLS-II, BSRF (Beijing), SSRF (Shanghai), SSLS, NSLS-II (Singapore)
- ...

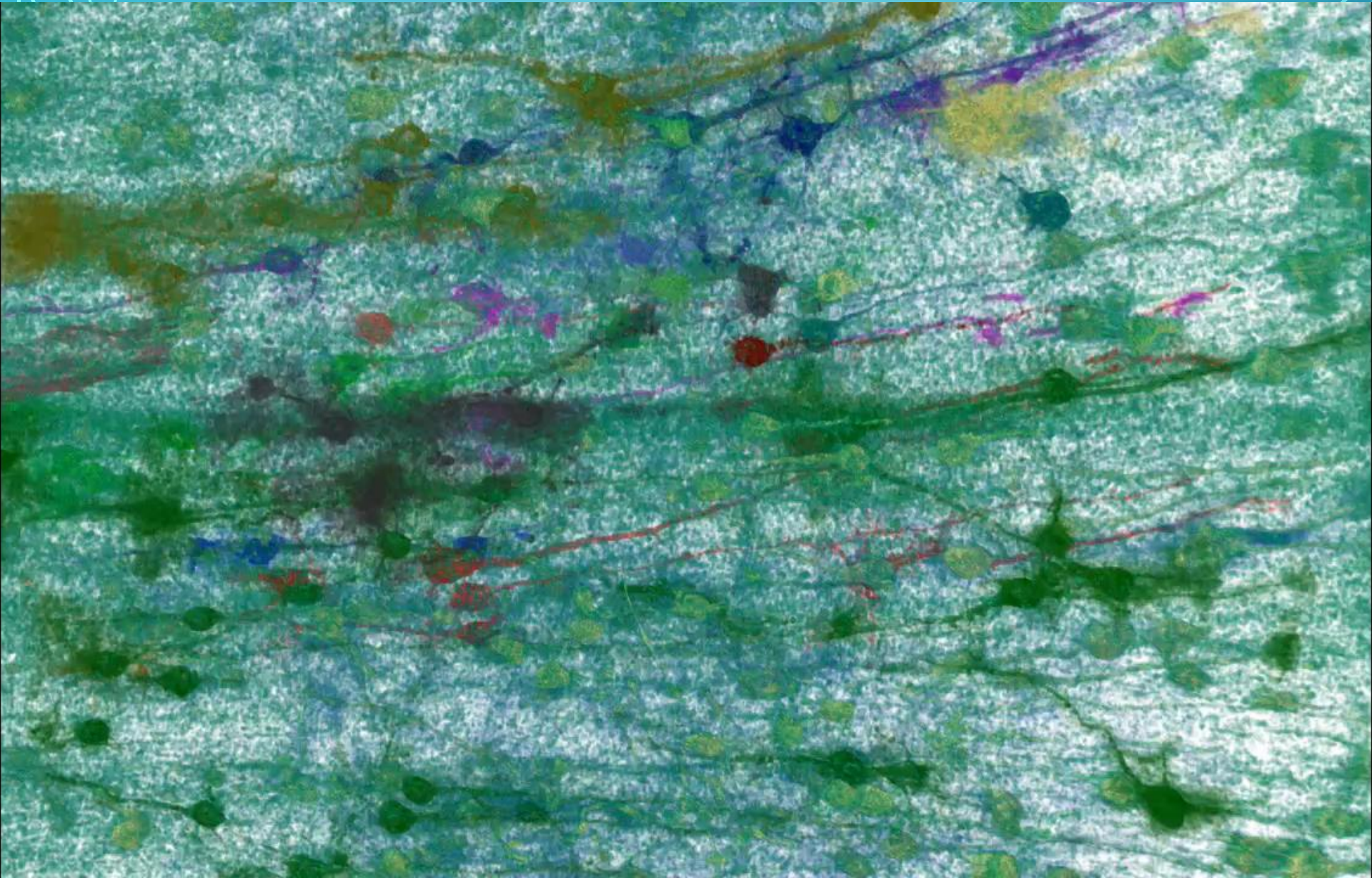
# Objectives – to map in 3 years:

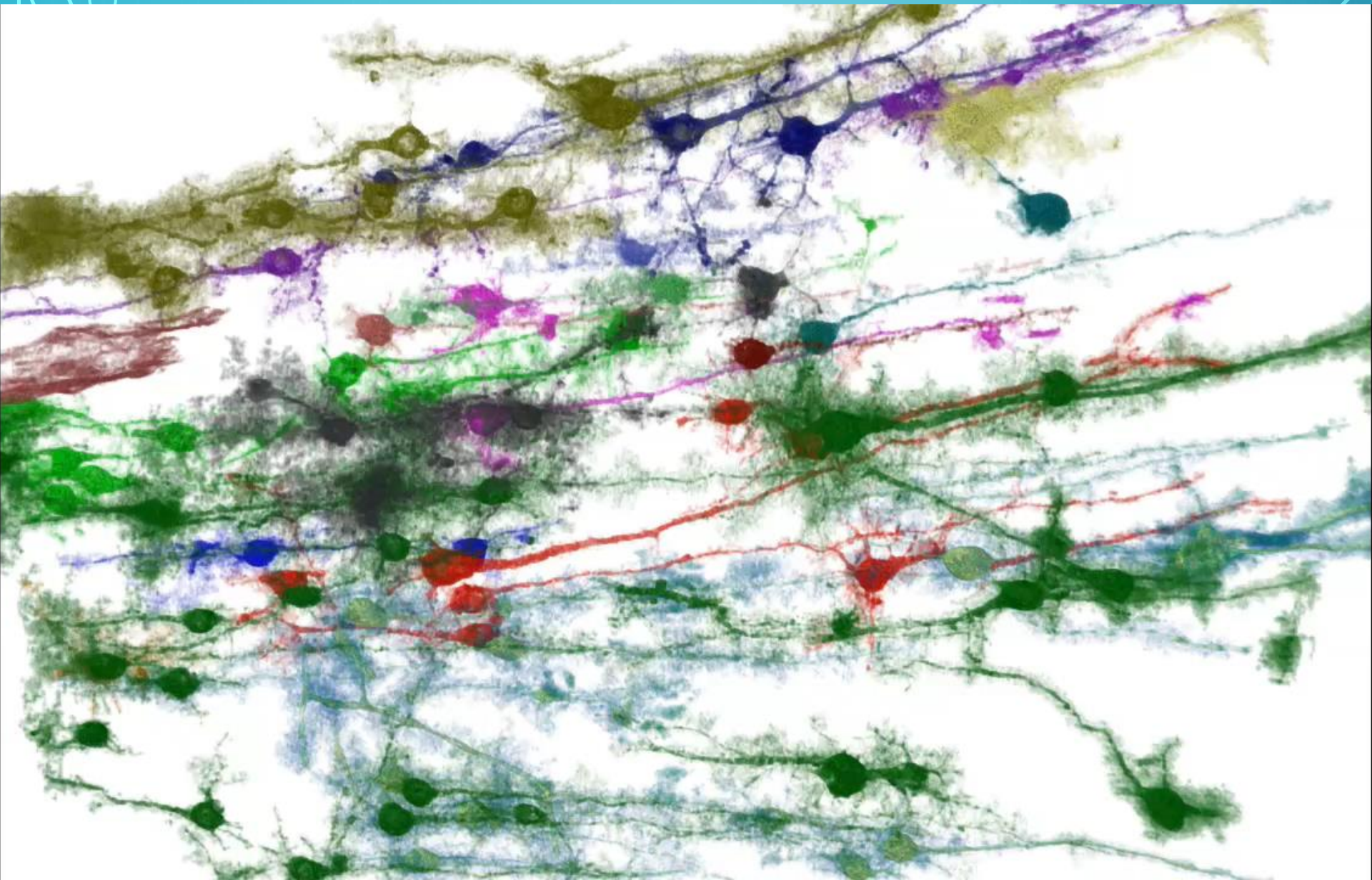
- One whole human brain
- 200 whole mouse brains
- Develop the microtomography technique for >100x improvement in speed to enable complete human connectome mapping

# Whole brain w/ 1 $\mu\text{m}$ resolution





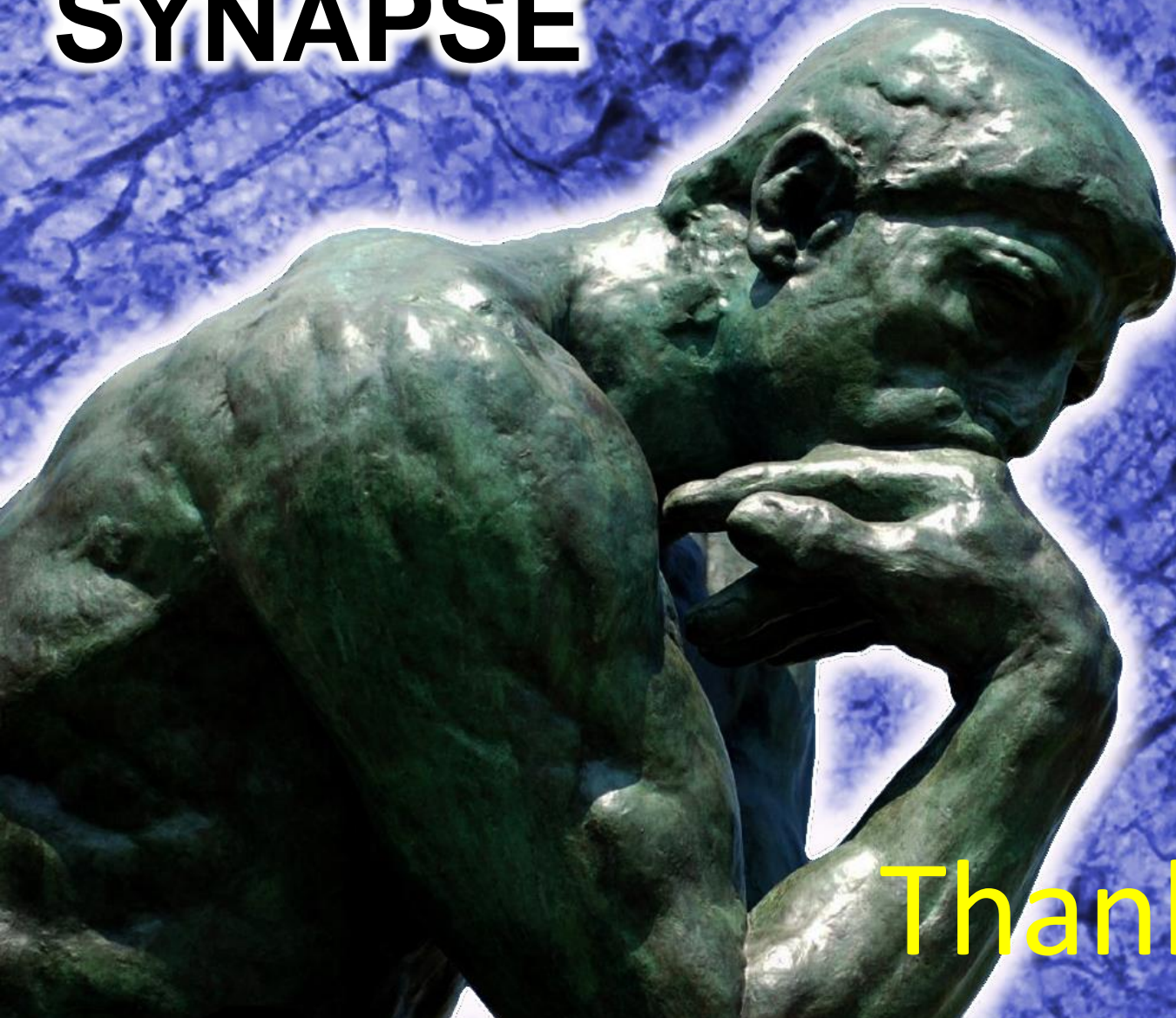








# SYNAPSE



Thank you!!

**2023: groundbreaking progress in human brain knowledge brought by the Asia-Pacific countries**

# Go fast and go far with friends!

(from China, France, Germany, Japan, Korea, Singapore, Switzerland, US, ..)

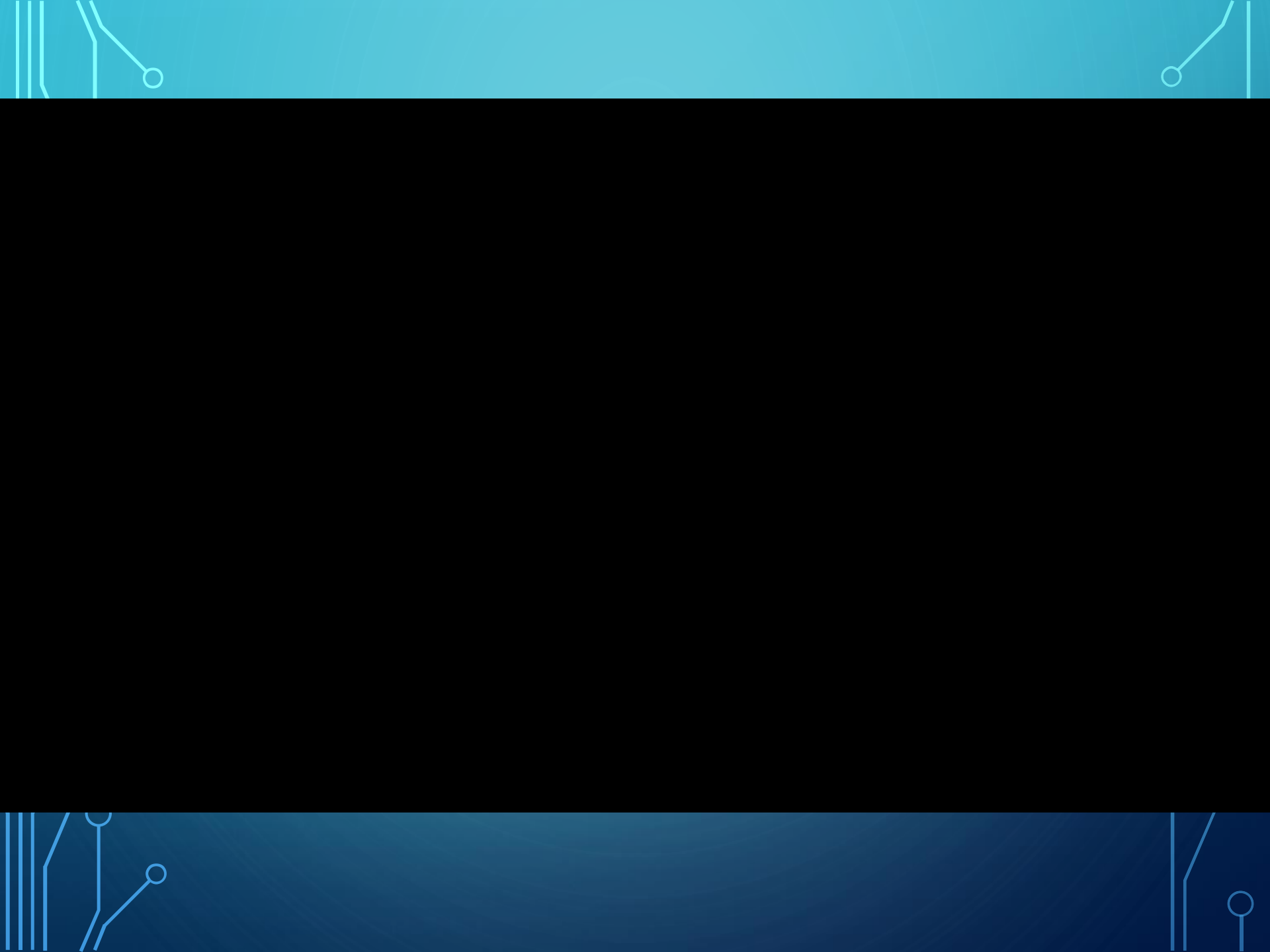


**TO THE MIND ~~FRONTIER~~**



Thanks for the funding support from:

- Ministry of Science & Technology
- National Program for Nanoscience and Nanotechnology
- Academia Sinica Thematic Projects
- ANR-MOST, INSERM-MOST
- RIKEN-MOST
- USAF-MOST





The SYNAPSE strategy:  
data acquisition  
and management

# Data size:

- Our experience from the tomography of a  $0.5^3 \text{ mm}^3$  volume containing one drosophila brain:
  - Size of each raw projection image: 32 MB
  - Tomography set (1000 projection images): 32 GB
  - Reconstructed images for volume rendering:  $\approx 128 \text{ GB}$
- Scaling to one mouse brain:  $[450/(0.5)^3] \times 128 \text{ GigaByte} \approx 460 \text{ TB}$
- Limited staining rate ( $\sim 5\%$ )  $\rightarrow$  100 mouse brains required for complete map:  $\sim 46 \text{ PB}$
- Scaling to one human brain (2,700 times the mouse volume):  $460 \text{ TB} \times 2,700 \approx 1,240 \text{ PB}$

# Total image-taking times at current speed:

- ~5 min for a  $(0.5)^3 \text{ mm}^3$  volume:

For one mouse brain:  $(450/0.5^3) \times 5 \text{ min}$   
 $\approx 1.8 \times 10^4 \text{ minutes} \approx 12.5 \text{ days}$

For 100 mouse brains: 1250 days  $\approx 3.4$  years

- ~15 min for a  $1 \text{ mm}^3$  volume (with a 4K x 4K detector):

For one human brain:  $1.2 \times 10^6 \times 15 \text{ min}$   
 $\approx 1.8 \times 10^7 \text{ minutes} \approx 34 \text{ years}$

For 100 human brains: 3,400 years

# We must increase the throughput. Without new technologies, by:

- Increasing the number of synchrotron beamlines used in parallel, coordinated within SYNAPSE
- Reducing the number of projection images for each tomography, from 1000 to 100.

With new algorithms

With artificial intelligence

It becomes thus realistic to map one human brain in 4 years – the first SYNAPSE objective

# Further throughput increases possible with new technologies:

- Staining rate increase from 5% to 30% (demonstrated but not optimized). The number of brains for a complete map could decrease to <20.
- Faster imaging: Higher x-ray flux (10x), higher detector sensitivity (5x) and better algorithms (to further reduce the number of projections) – overall, the image acquisition speed could be increased by two orders of magnitude.

The image taking time for full human brain mapping could be reduced to a few years



# Image taking is not all: data managing is a big challenge!

- Tomography reconstruction for drosophila: currently, 10 times slower than image acquisition
- To perform one automatic segmentation and tracing within a comparable time, it takes now the largest computing facility in Taiwan, NCHC
- Commercial graphic workstations cannot handle the rendering and visualization of the reconstructed data set
- Another big challenge: creating and managing the database
- Further complication: adding functional information to the map

# Crucial data handling tasks:

- Image morphing, warping and fusion
- Correlation identification
- Tracing, segmentation
- Database architecture and optimization
- Database access
- Visualization

# Adding functional information:

- Using the connectome as the skeleton/grid.
- Adding local high-resolution 3D information on synapse connections with:
  - X-ray tomography with 10 nm resolution.
  - Super-resolution 3D fluorescence microscopy (with tissue clearing)
  - Super-resolution vibrational spectromicroscopy (Nano-IR and Nano-Raman)
  - Continuous sectioning with electron and optical microscopy
- Chemical information in large region:
  - Functional MRI, PET & SPECT imaging (with 100  $\mu\text{m}$  resolution)
  - 3D IR (with  $<3 \mu\text{m}$  resolution)
- Simulations based on the structural and functional information

# Other technological developments targeted by SYNAPSE:

- Algorithms and related software
- Computer hardware: machine-learning graphic station
- Automation in specimen handling
- Automation in image acquisition
- Artificial Intelligence for automation of image processing
- Image taking hardware: specimen manipulation, detectors and high-brightness X-ray sources

# Why us?

## We are uniquely qualified!

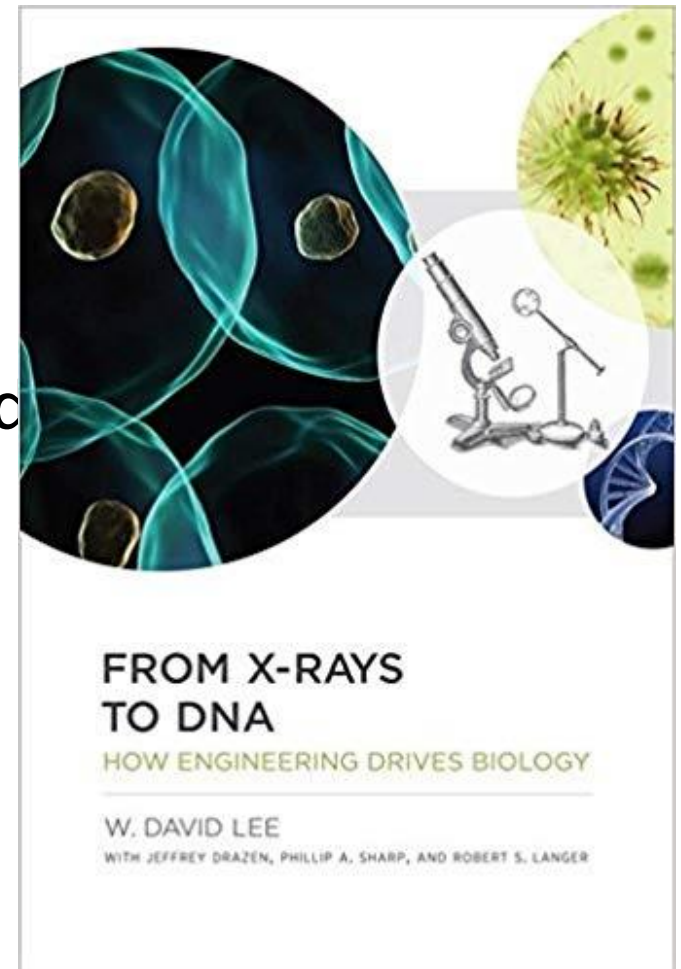
- We pioneered the special techniques required for the SYNAPSE initiative, including:
  - X-ray phase-contrast tomography
  - World-record x-ray microscopy
  - Deep tissue super-resolution microscopy
- We have a solid record of relevant previous accomplishments and all the required know-how for the core and support techniques
- The new partnership is built on long-standing and very successful collaborations

# Sizes of Brains:

	Volume (mm <sup>3</sup> )	Number of neurons	Brain/body ratio	Number of synapses	Density of neurons (mm <sup>-3</sup> )
drosophil a	$2 \times 10^{-2}$	$2.5 \times 10^5$ ( $1.3 \times 10^5$ )			$1.25 \times 10^7$
mouse	450	$7.1 \times 10^7$	1:40	$10^{11}$	$1.57 \times 10^5$
marmoset		$6.4 \times 10^8$			
human	$1.2 \times 10^6$	$8.6 \times 10^{10}$	1:50	$10^{14}$ - $10^{15}$	$7.2 \times 10^4$

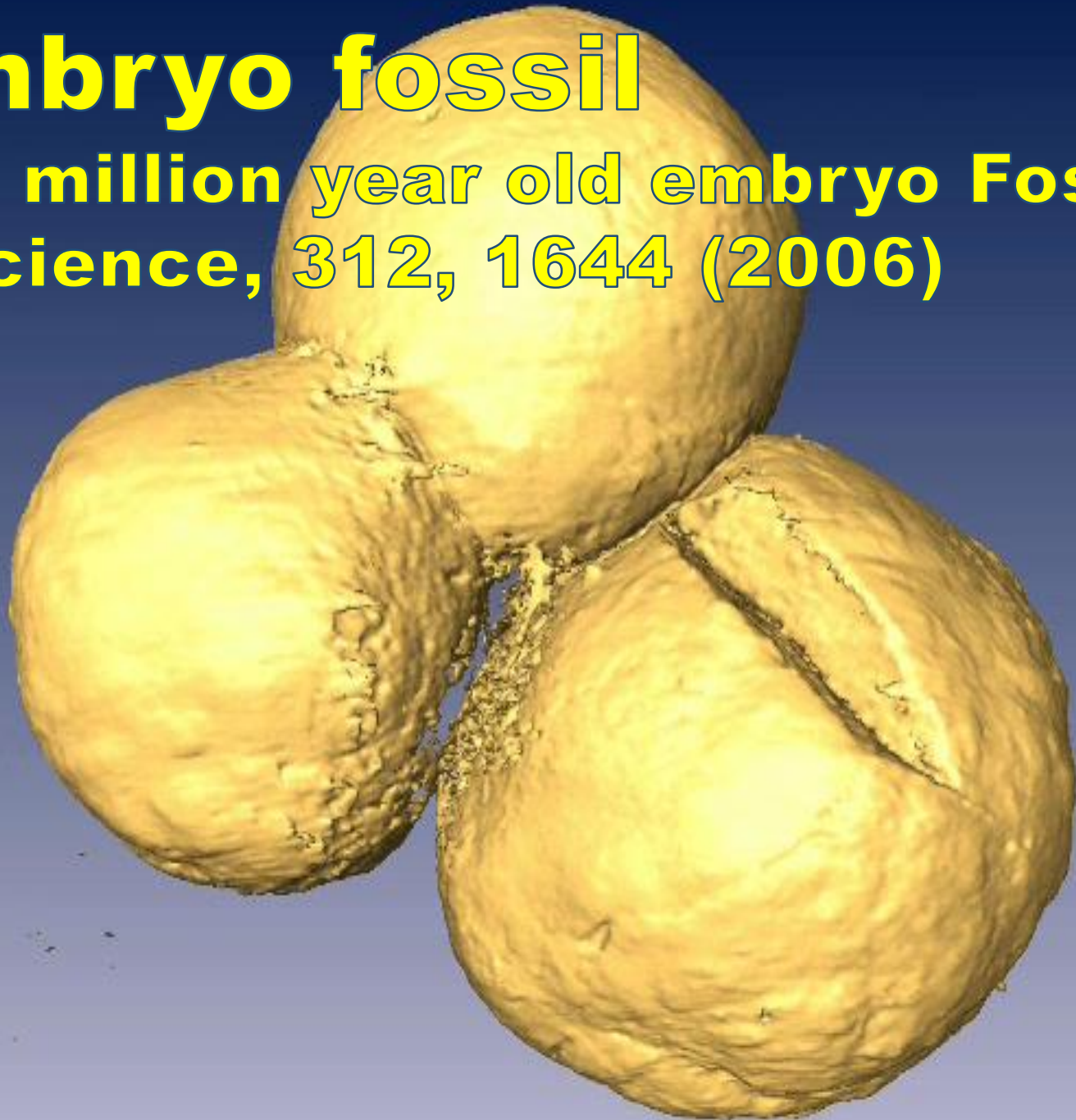
# Physics opens the new era of biology and medicine, again and again.

- Microscopy
- Spectroscopy
- X-rays
- Protein crystallography
- NMR, SPECT, PET, CT, ultrasound
- Mass spectrometry
- Nuclear Medicine
- Digital Camera
- ...



# Embryo fossil

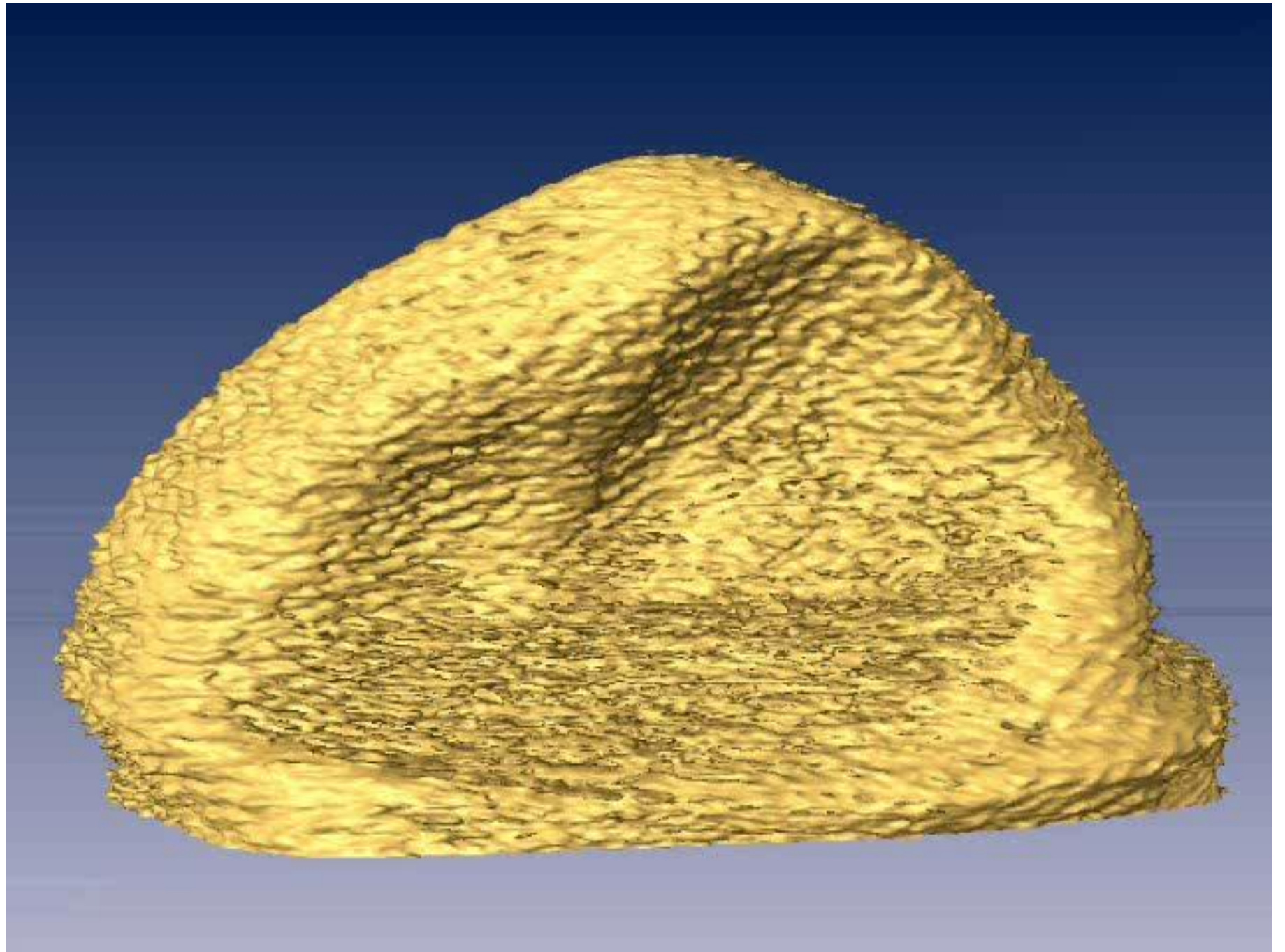
540 million year old embryo Fossil  
-- Science, 312, 1644 (2006)

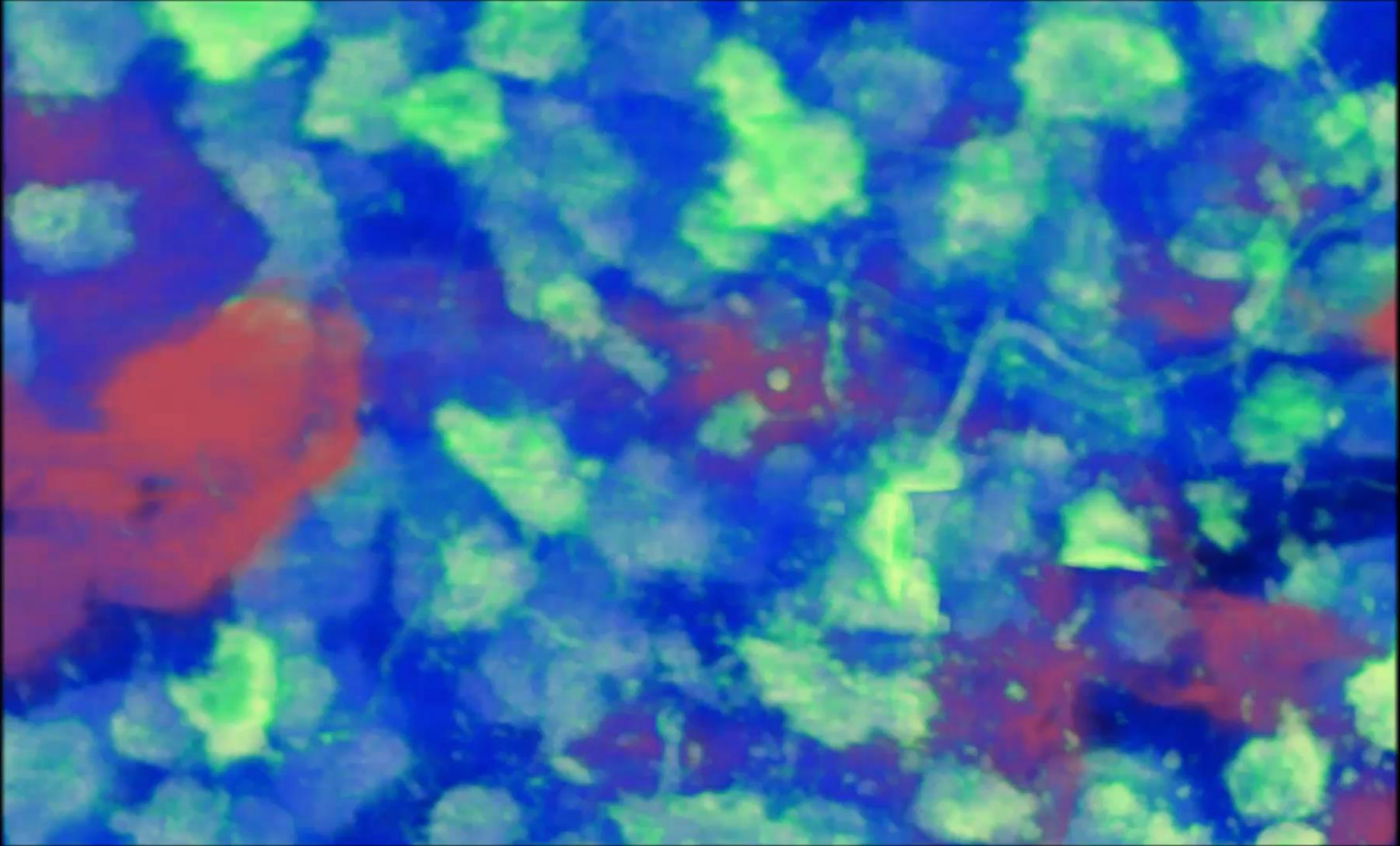




540 million year old embryo fossil





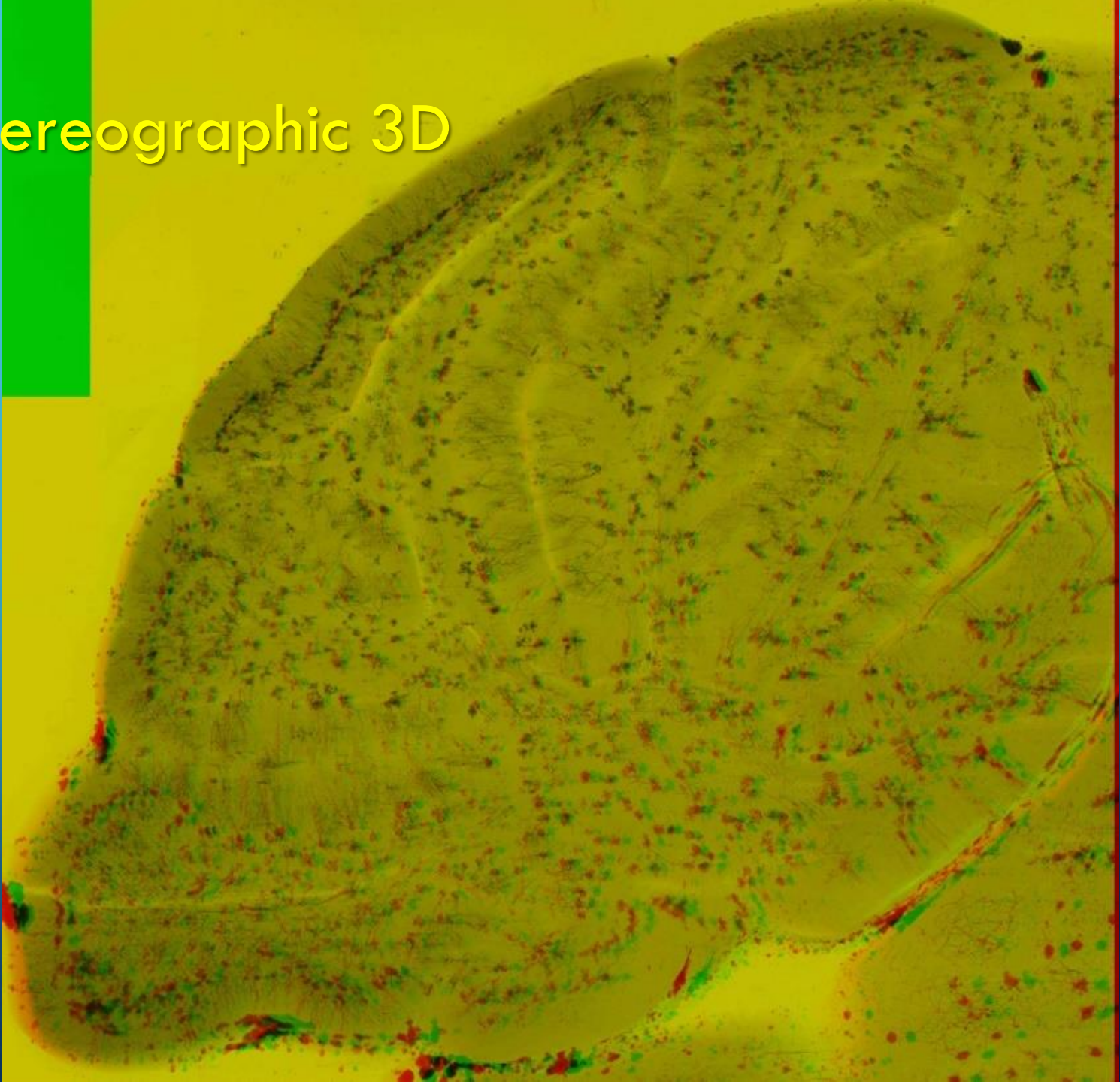


# Concentrate on imaging

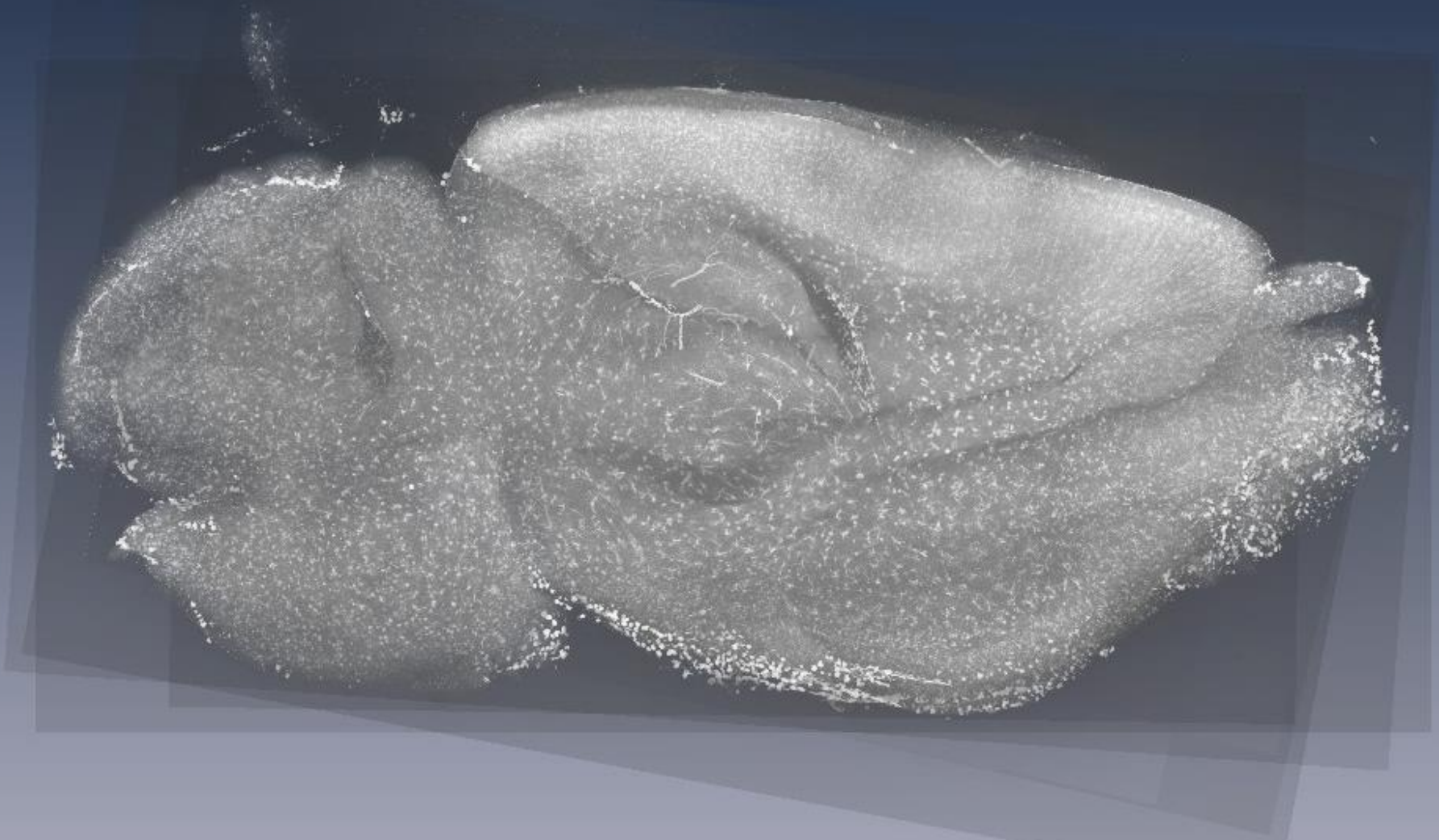
## Structure and Network:

- X-ray phase contrast microtomography, 0.3  $\mu\text{m}$  resolution in all 3D directions: to provide the wiring map
- X-ray nanotomography: 10 nm to provide the structure and location information of targeted synapse and neuron connections
- Confocal + *FocusClear*<sup>®</sup>: Full drosophila brain and regions of mouse brain with functional information
- Cryo-EM: small region 3D map down to nm resolution.
- Super-Res + *FocusClear*<sup>®</sup>: Full drosophila brain and small region of mouse brain but concentrate on single molecule detection

# Stereographic 3D



# Volumetric view



# ONE STEP FURTHER...

Chian Ming Low, Eng Soon Tok, Alvin Teo, Tin-wee Tan (National University of Singapore)

# Zhong Zhong & Hua Hau (中中與华华)





# Monkey Hippocampus (*Macaca Fascicularis*)

## Hippocampal Formation

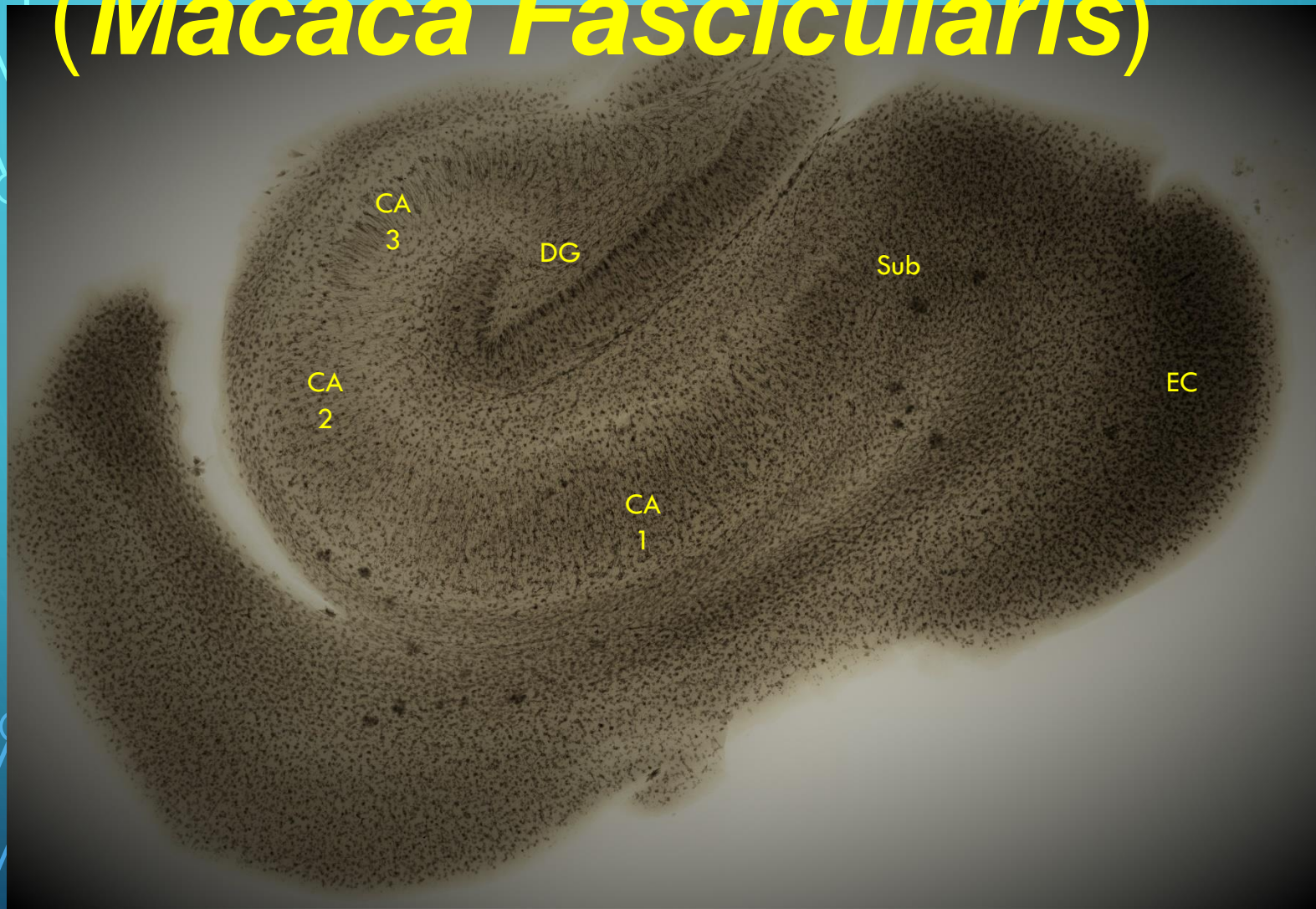
DG – Dentate Gyrus

CA – Cornu Ammonis

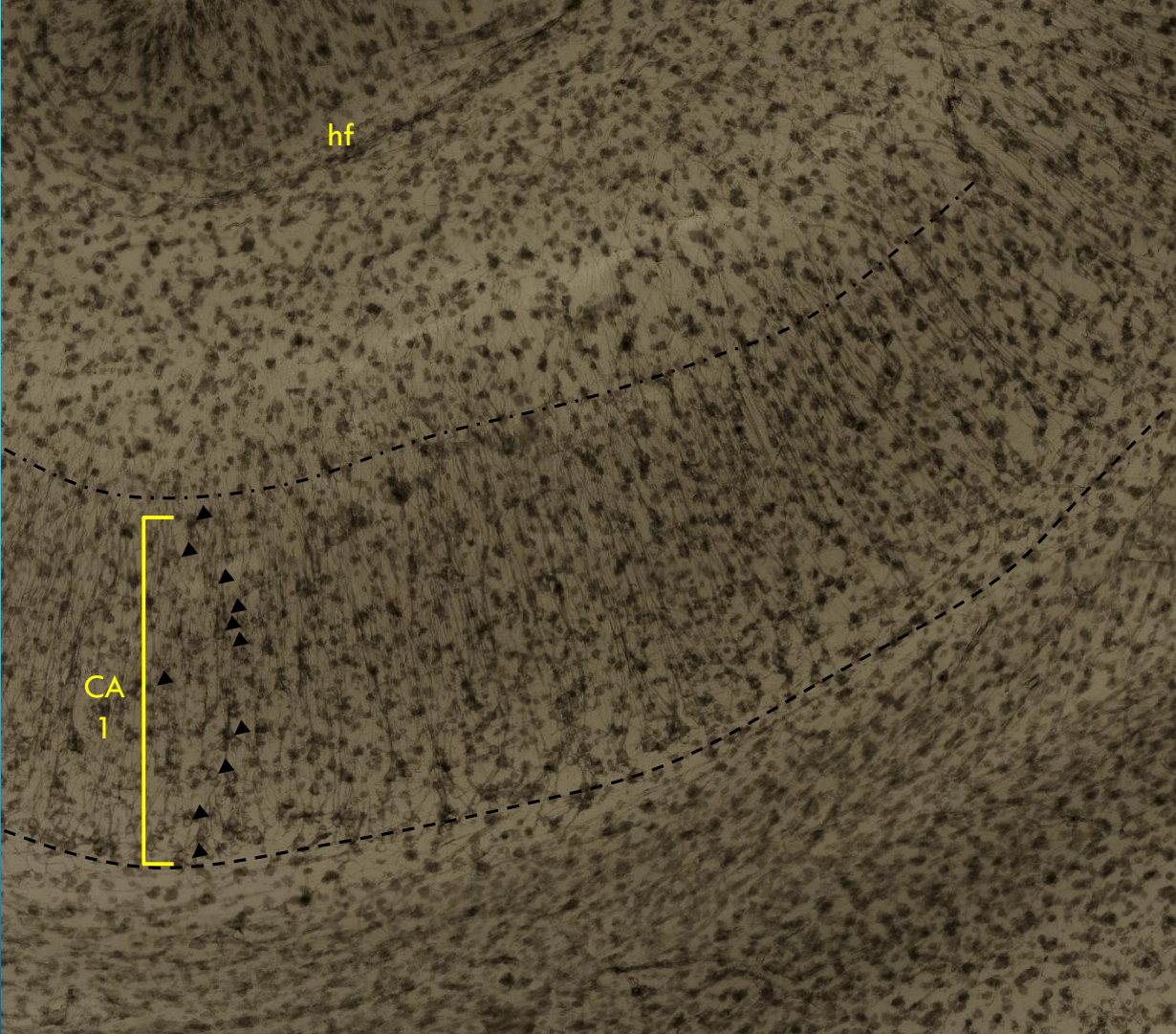
Sub – Subiculum

EC – Entorhinal  
Cortex

Note: hippocampal  
formation consists of  
these regions – DG,  
CA3, CA2, CA1,  
Subiculum &  
Entorhinal  
cortex...you can also  
call it hippocampus (a  
more 'general' term)

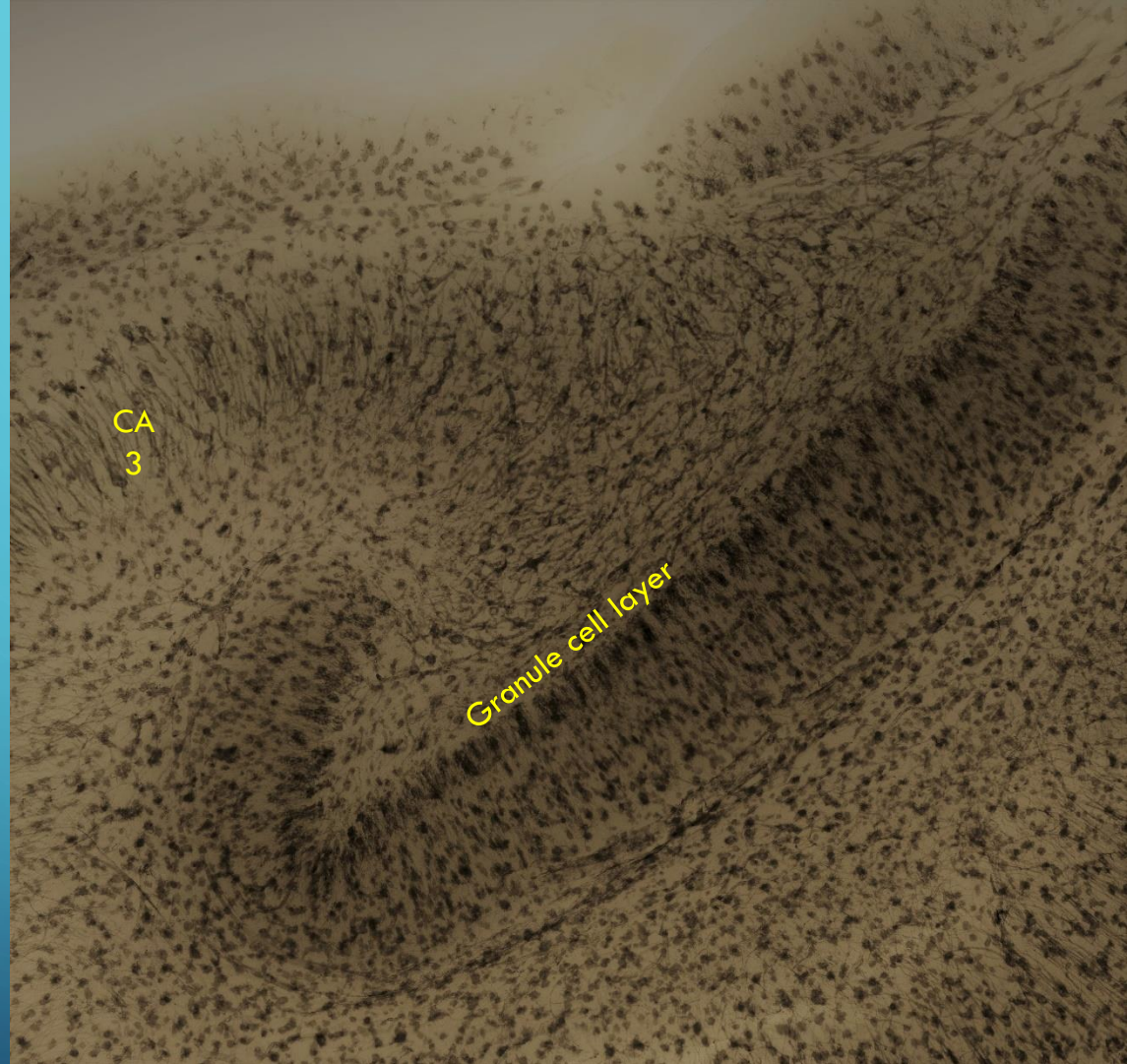


CA1	Rat	Monkey
# Divisions	1-2	7
# Cell Thick	5	10-15
Boundary between PCL-SR	Clear	Less clear
Boundary between CA1-Subiculum	Clear	Less clear
Entorhinal Cortex – laminar organization	Less clear (e.g. Layers V-VI) 2 divisions	Clear (e.g. Layers V-VI) 7 divisions

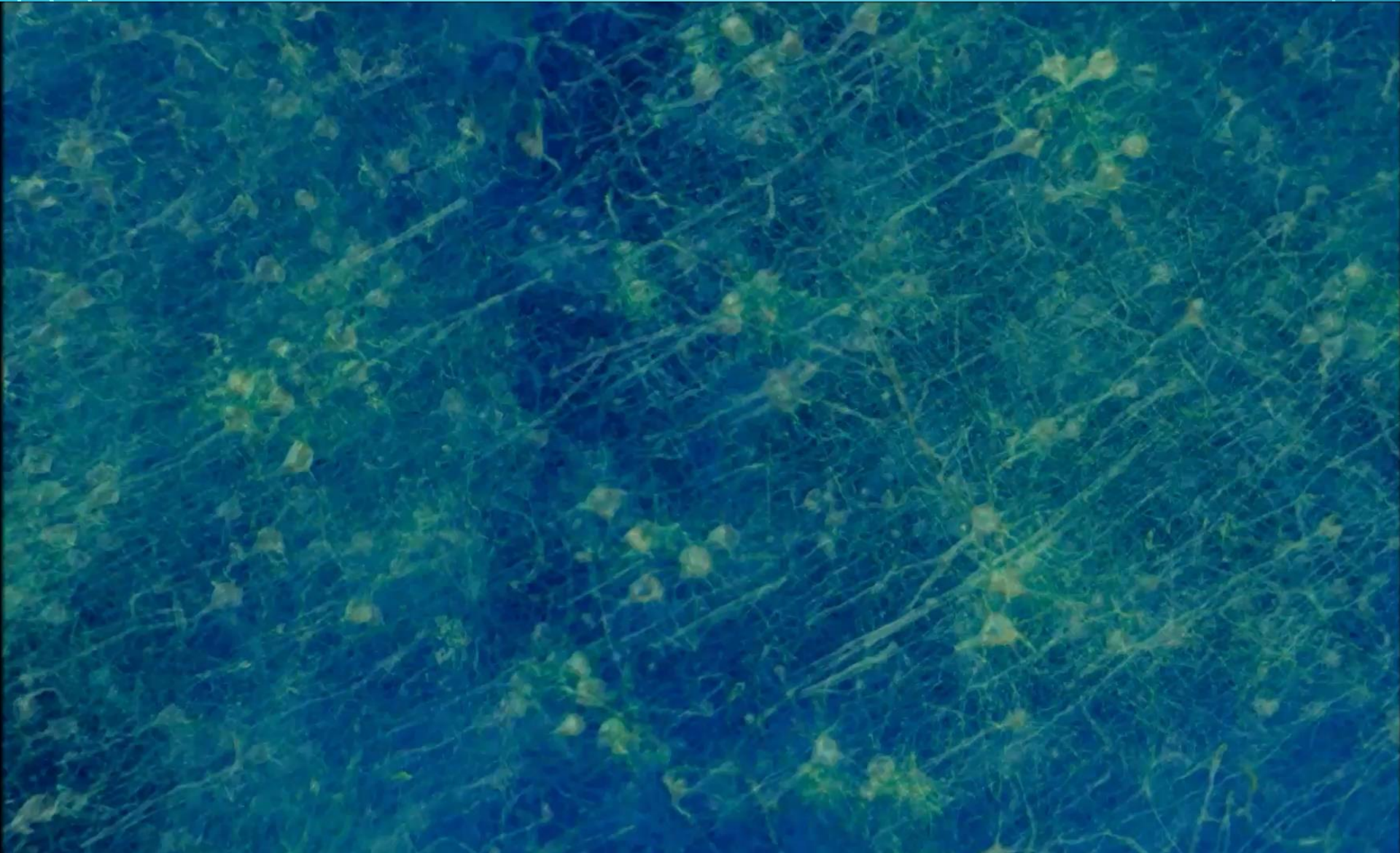


hf – hippocampal fissure (a 'line' that DG is separated from CA1)  
 Arrowhead – neuron soma

# Dentate Gyrus



# Mouse brain 2x2x20 mm<sup>3</sup>



# Besides Image Taking:

- Full biological characterization
- Specimen preparation: novel labeling procedures
- Coordinated use of different imaging facilities at partner countries, to reach the required throughput
- Standardization and mutual validation for all participating facilities
- Advanced image processing with new computer techniques
- Full data access for all partners and eventually to public

# The core partners bring with them a very powerful extended coalition:

- Through NUS/SSLs: research teams from university (NUS, NTU, NYP), public institutions (NCSS)
- Through SARI/SSRF: research teams from universities (Shanghai Jiao Tong U., Shanghai U.), public institutions (SINAP, IoN)
- Through POSTECH/PAL: research teams from universities (POSTECH, KAIST), medical institutions (ASAN, Samsung, SNU, Yonsei)
- Through RIKEN/Spring8: public institutions (RIKEN QBiC-BDR, BSI)
- Through ANSTO: research teams from universities (U. Sydney, U. Wollongong), public institutions (Australian Synchrotron, Brain & Mind Center)
- Through AS: research teams from universities (NTU, NTHU, NCTU, NCKU), public institutions (ITRI, NHRI, NHPC), medical institutions (NTUH, Chang Gung H, Mackay H, TSGH, VGH, Chinese Medical UH, Taipei, Medical UH, Kaohsiung MU, Cheng Kung UH) and from the private sector (TTY Pharma, Delta Electronics)