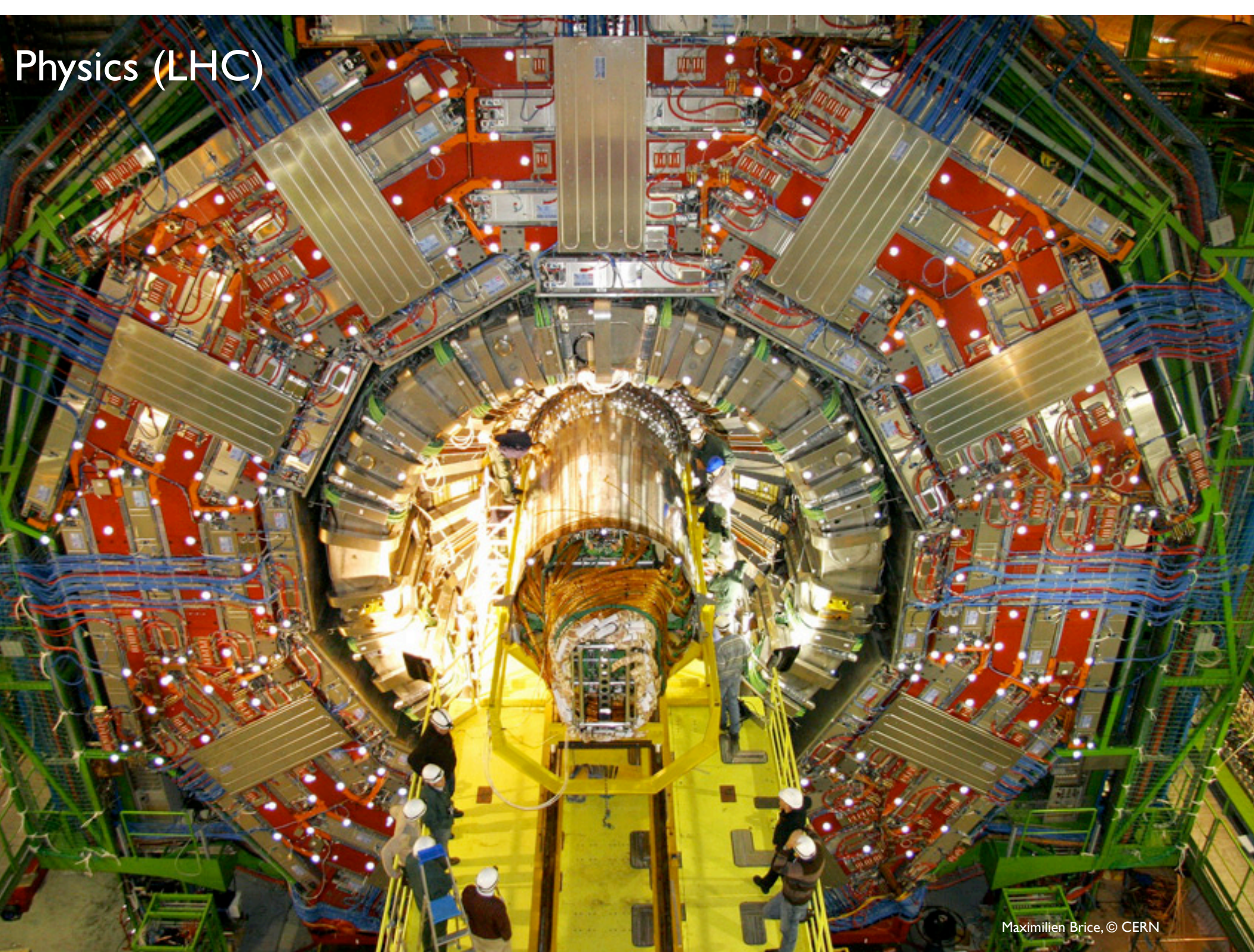


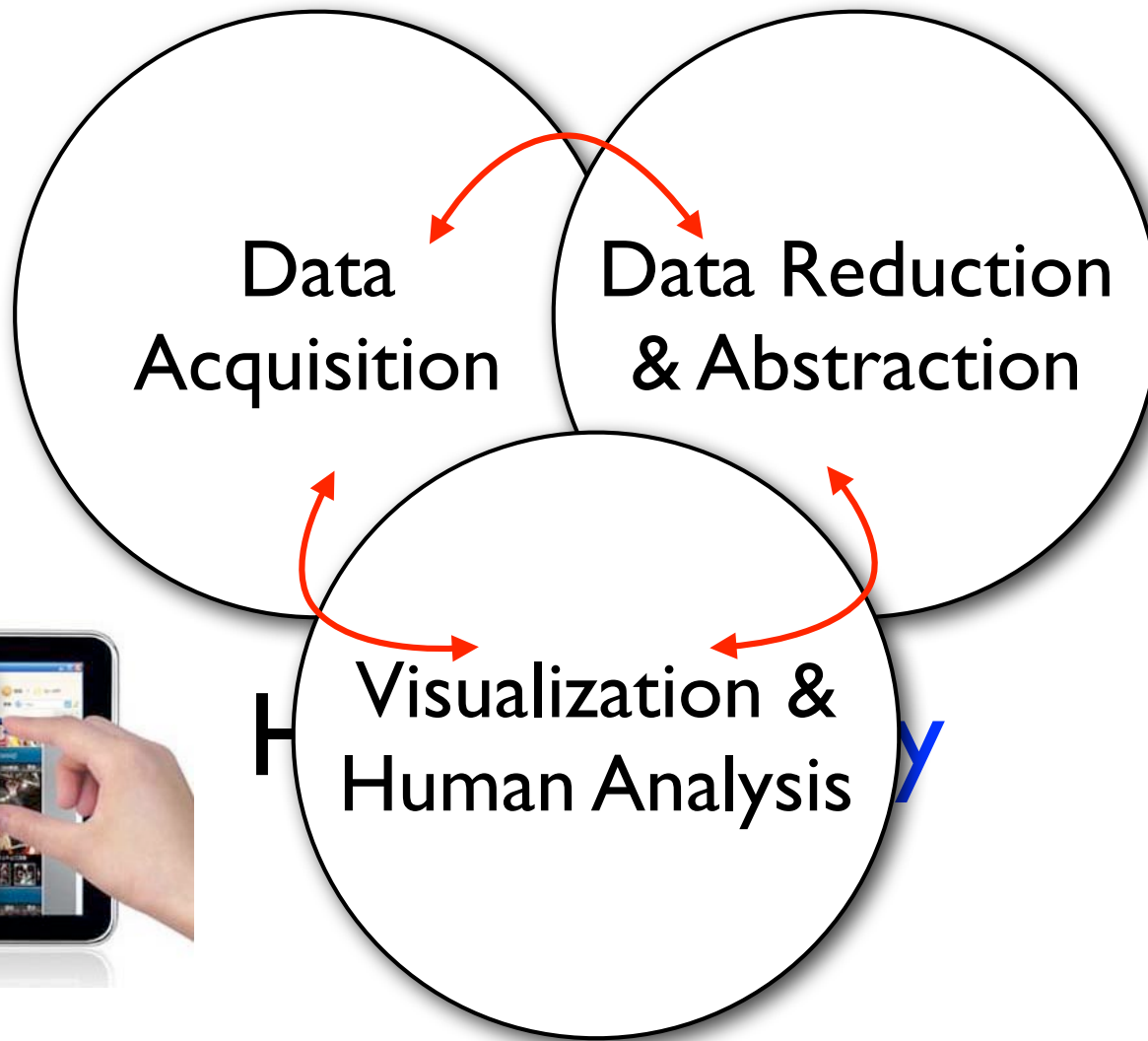
Visual Computing in Connectomics

Hanspeter Pfister, Harvard University
pfister@seas.harvard.edu

Physics (LHC)



Challenges



Visual Computing

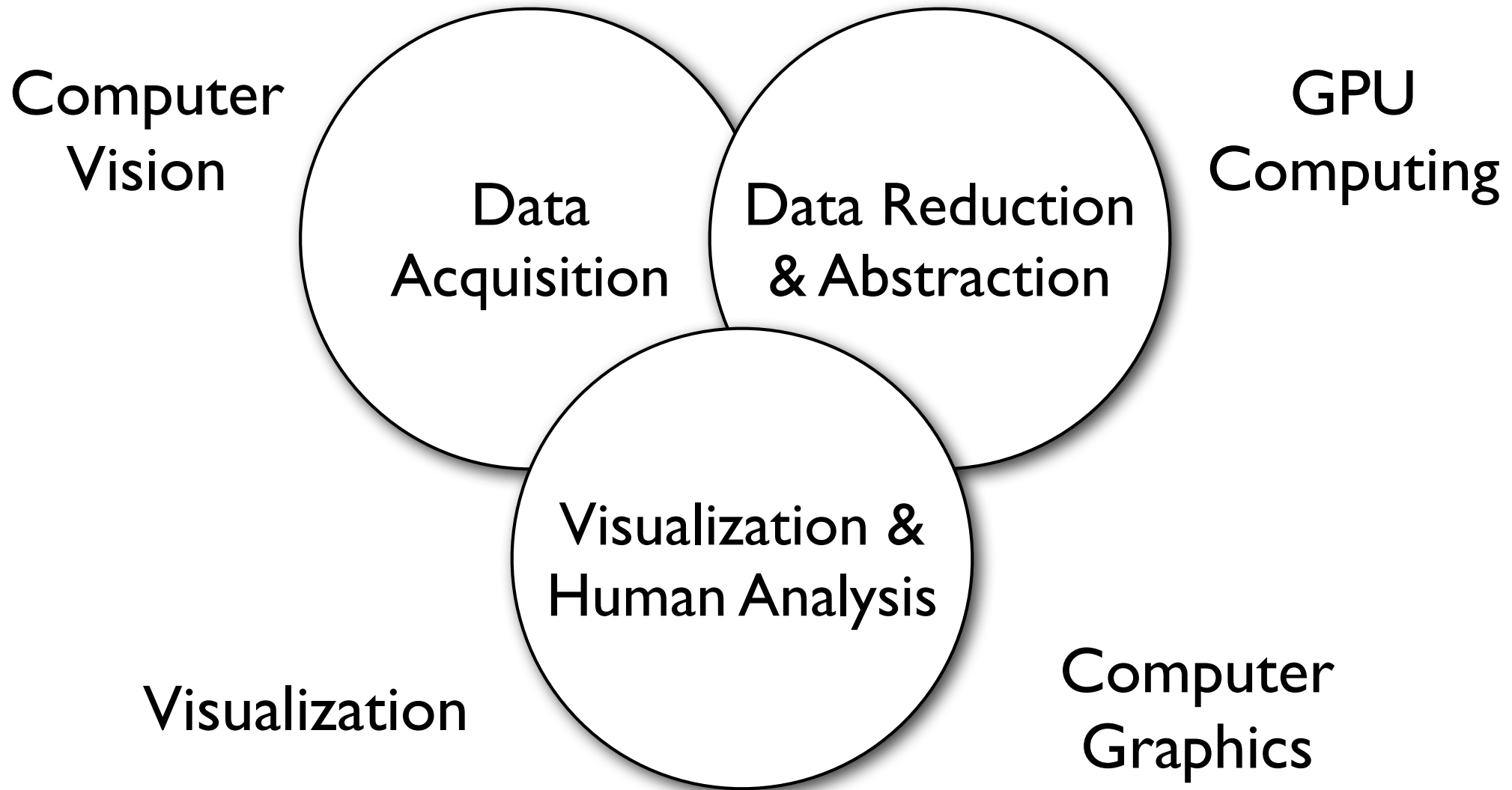
Data
Acquisition

Data Reduction
& Abstraction

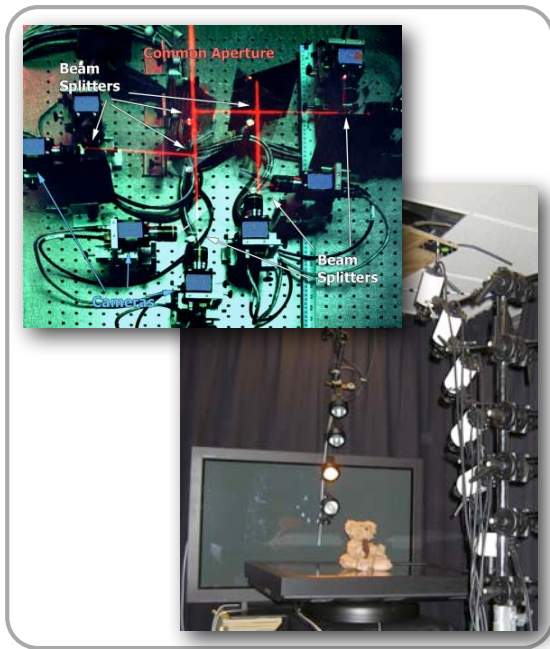
Visualization &
Human Analysis



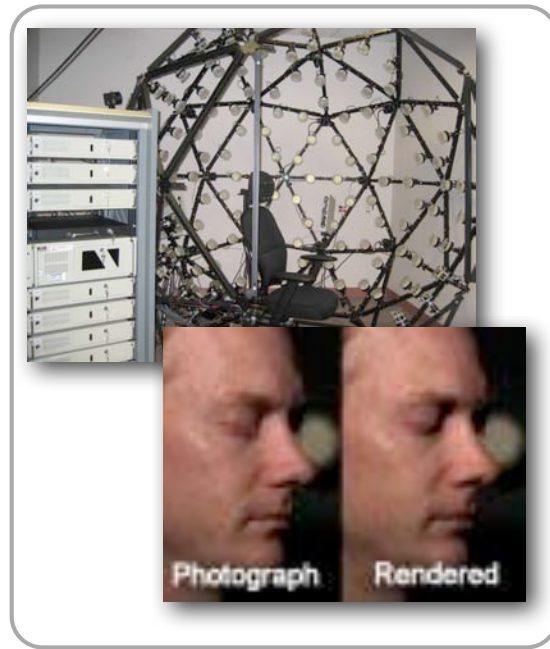
Visual Computing



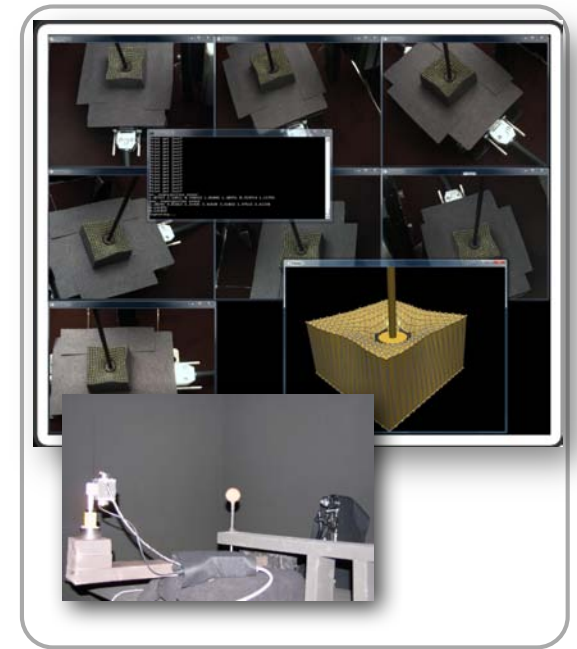
Data Acquisition



Computational
Photography



Face Scanning

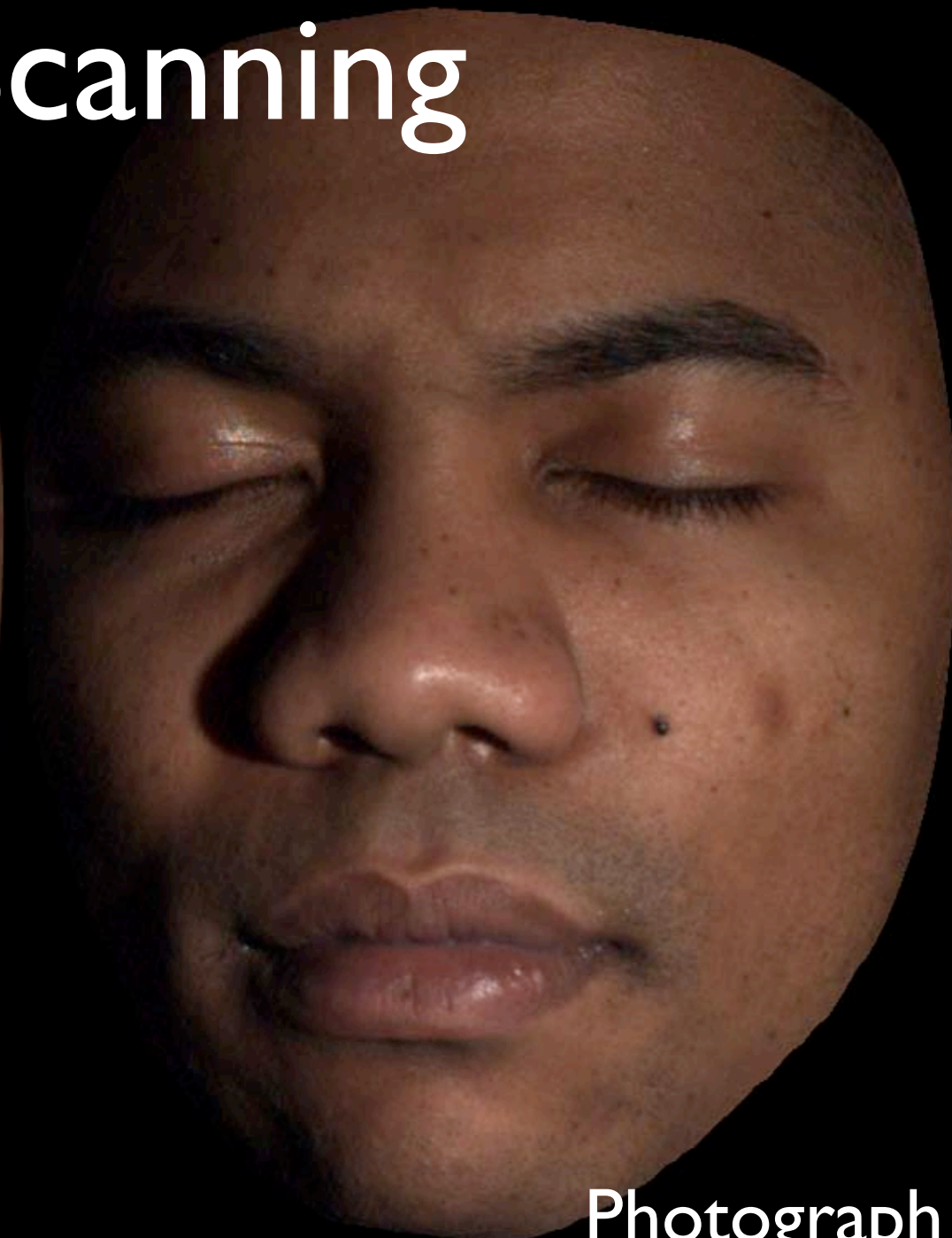


Material
Measurements

Face Scanning



Our Model



Photograph

Face Scanning Dome



Face Database

- ~500 subjects
- ~12 TB of data
- Statistical models of facial appearance

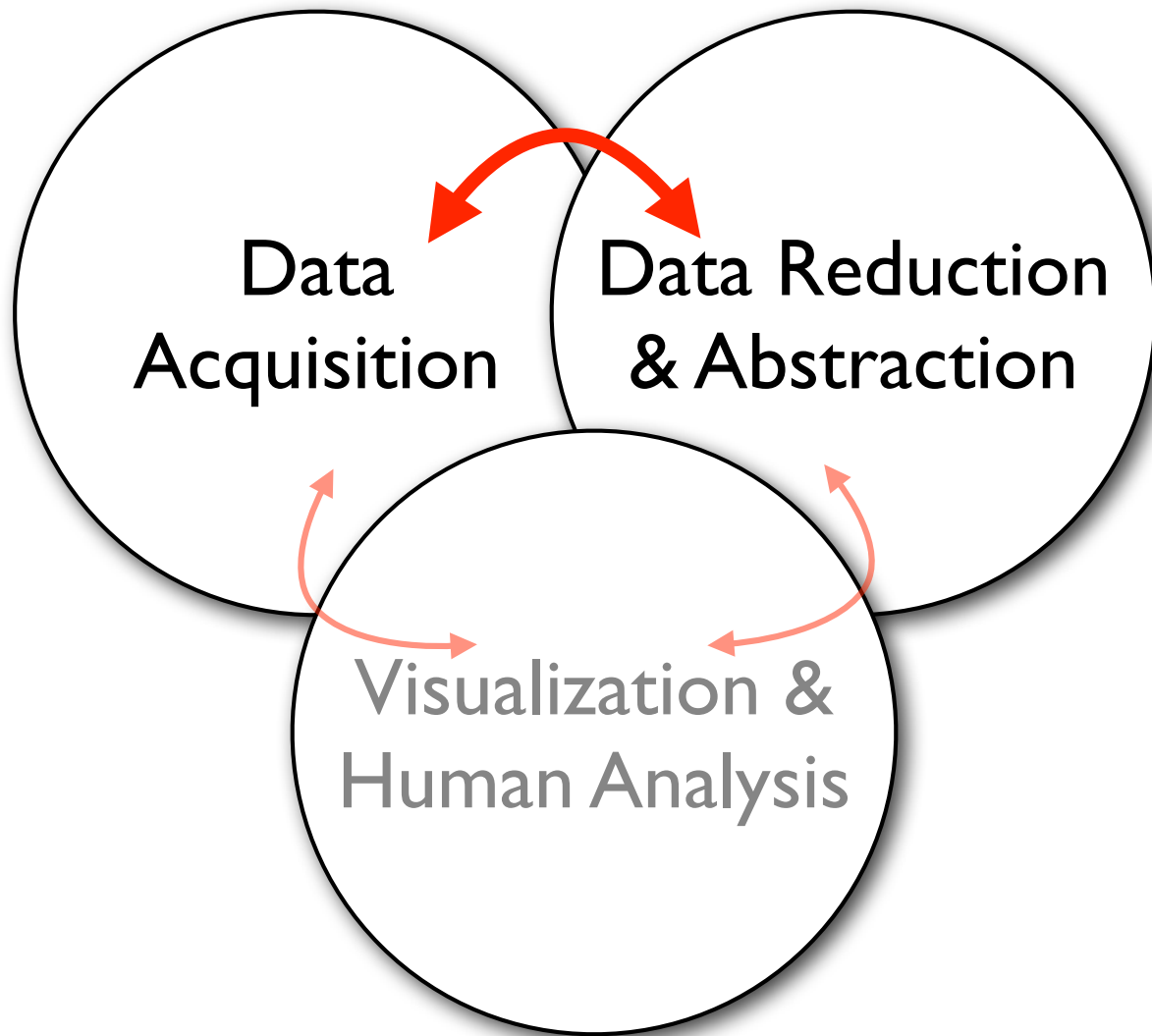


[Weyrich et al., SIGGRAPH 2006]



[Golovinskiy et al., SIGGRAPH 2006]

Visual Computing



Data Reduction & Abstraction

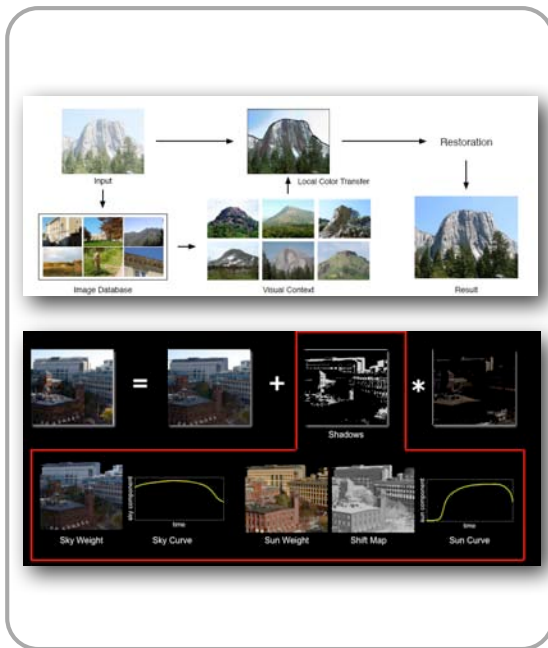
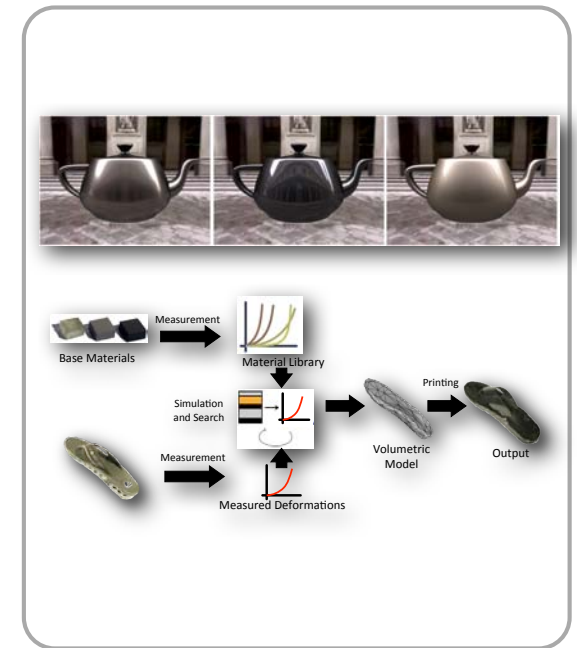


Image & Video
Analysis



Face Models



Material Models

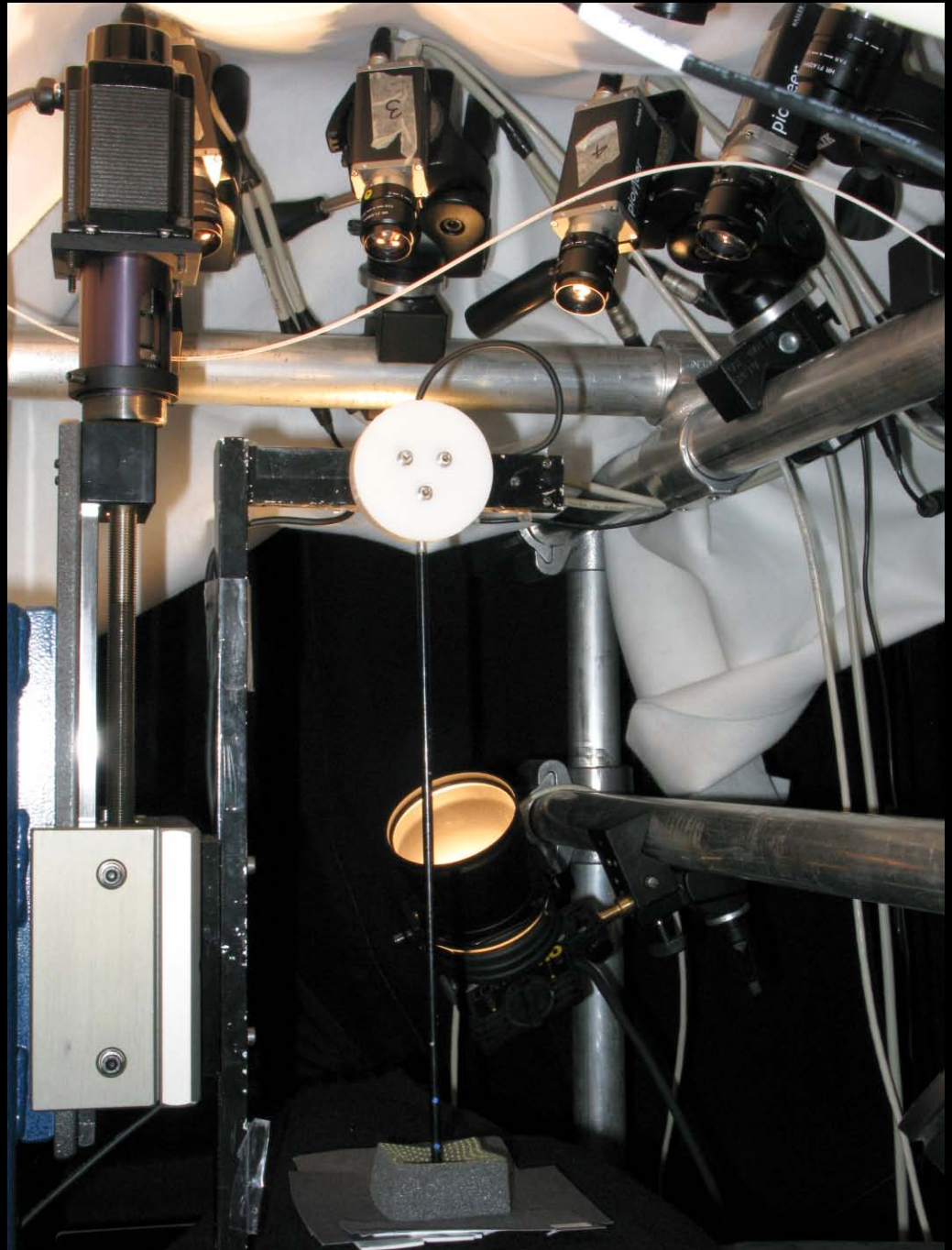
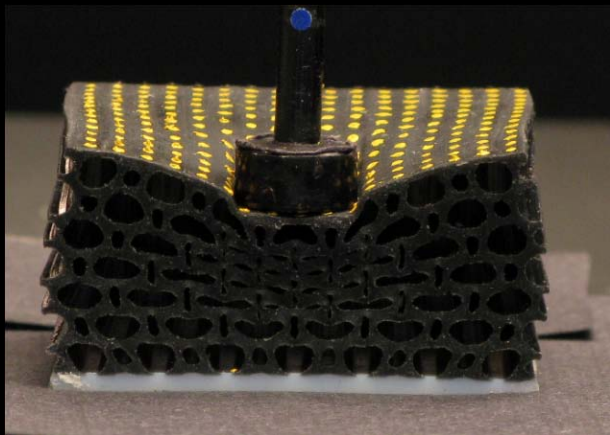
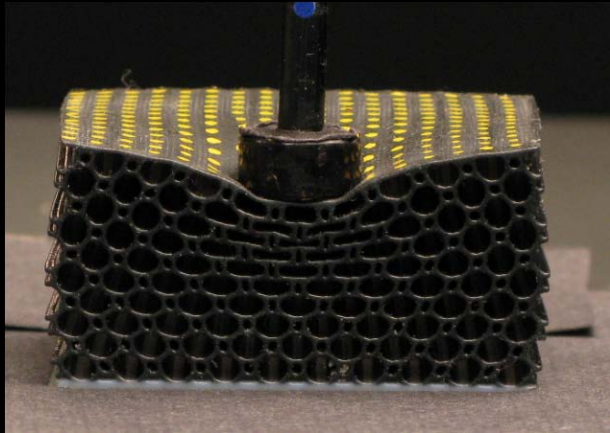
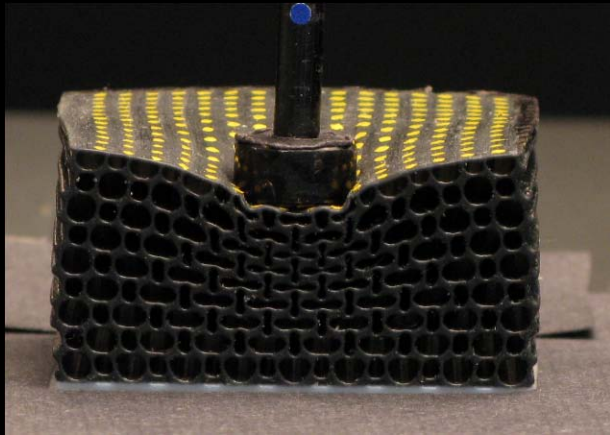
Material Design & Fabrication

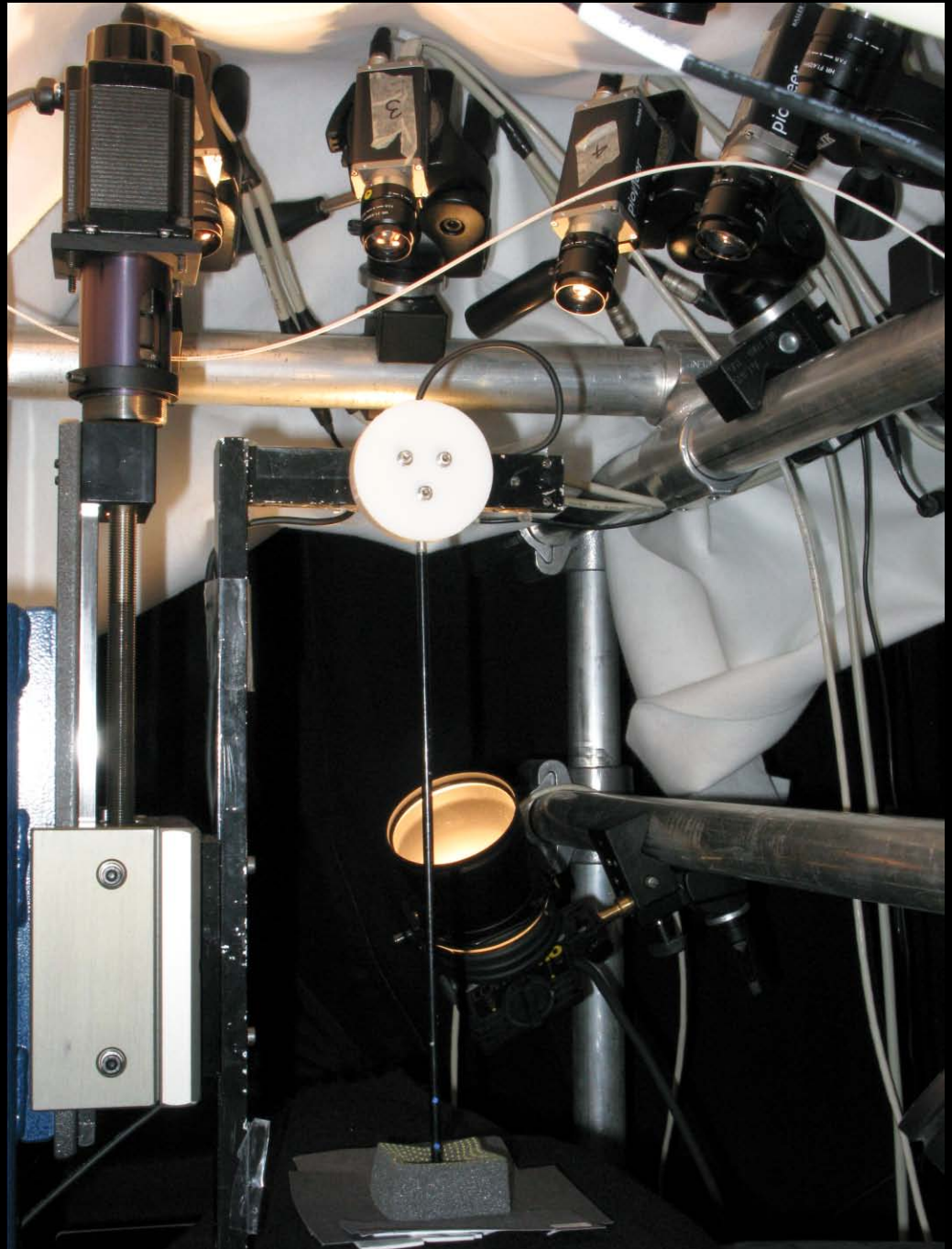
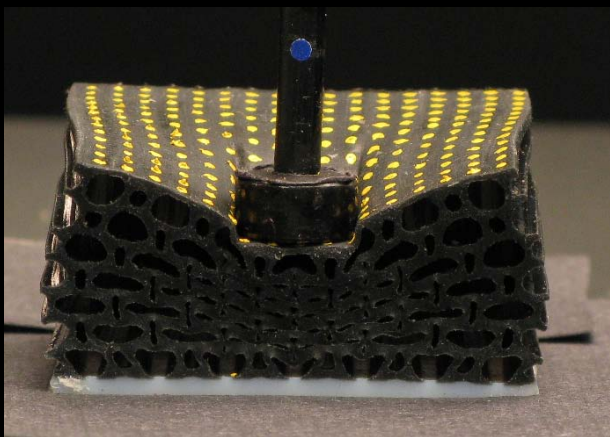
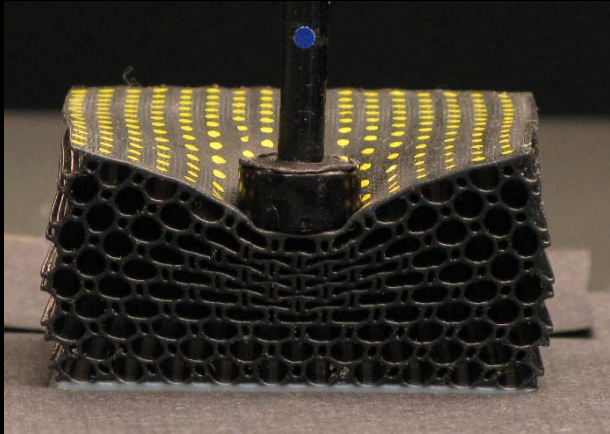
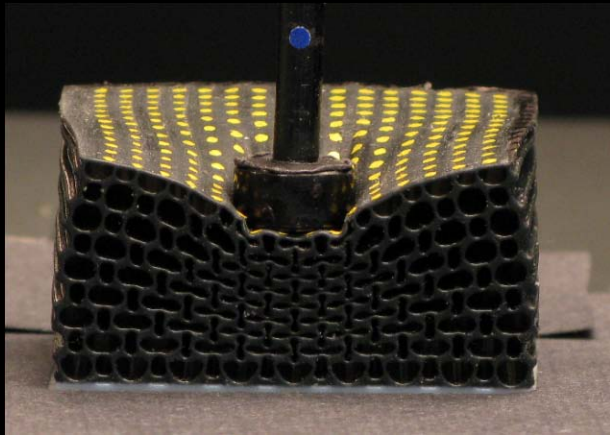


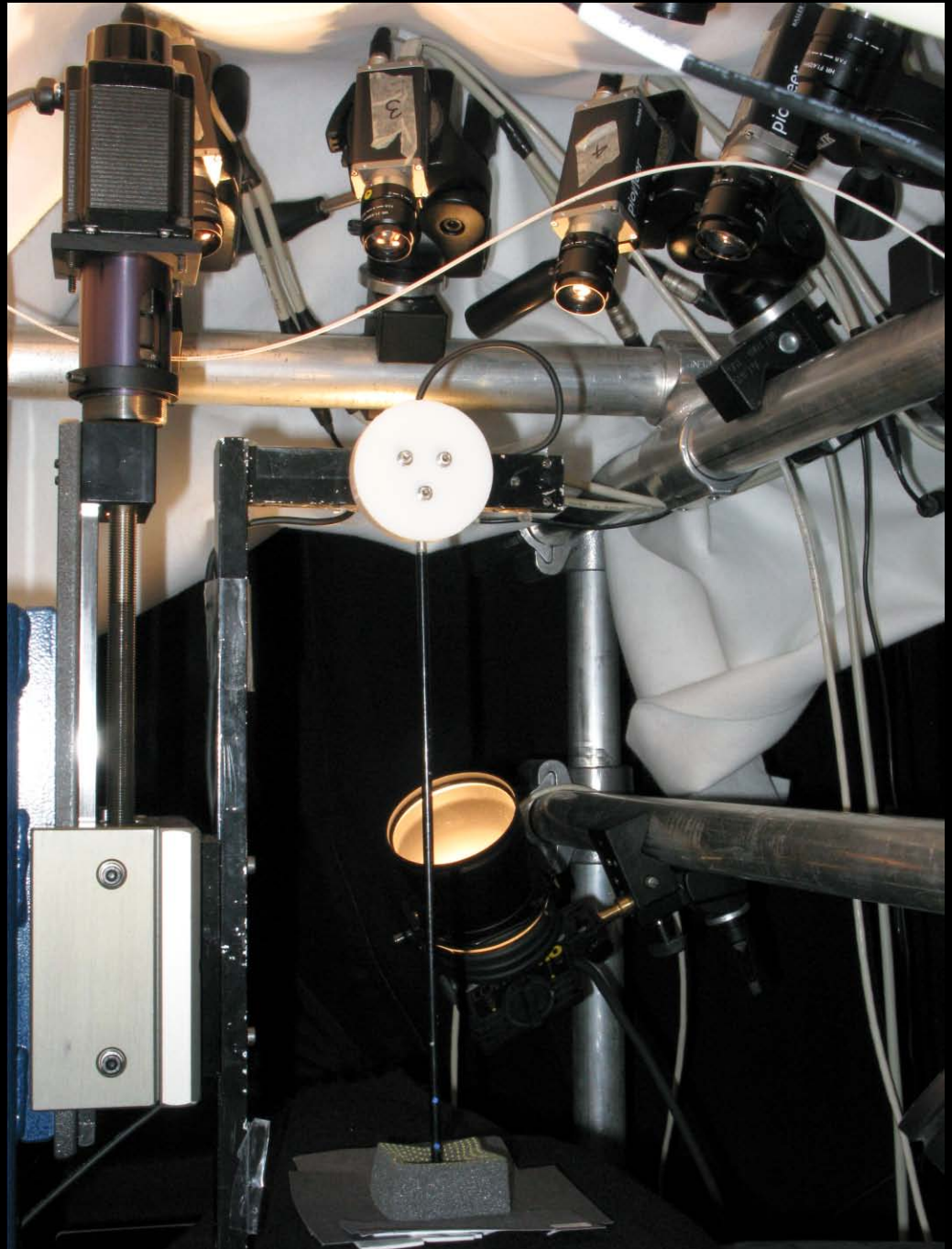
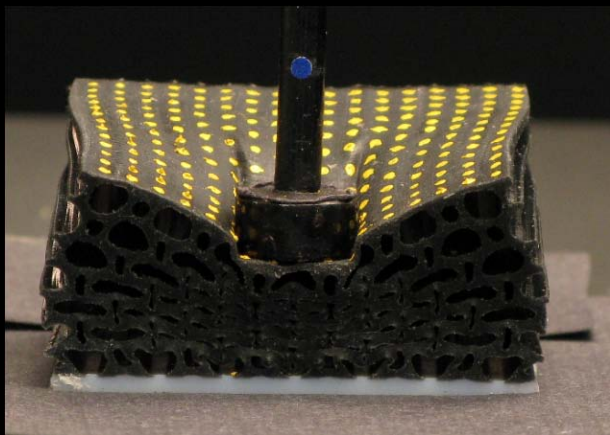
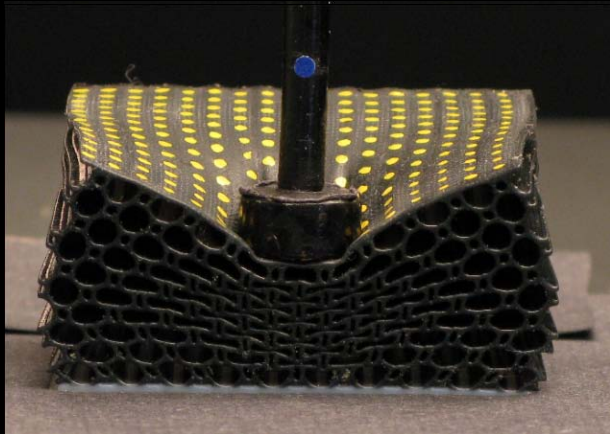
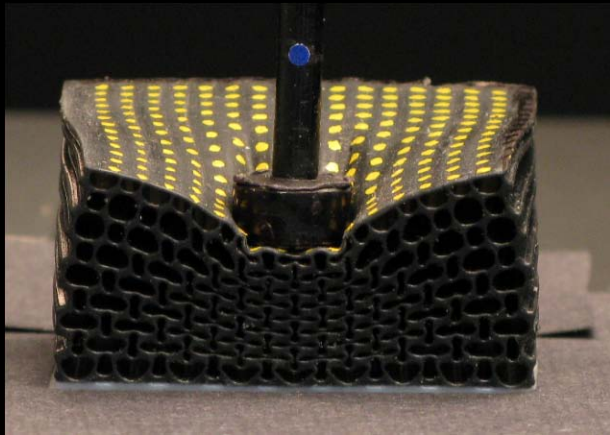
**Real
Object**

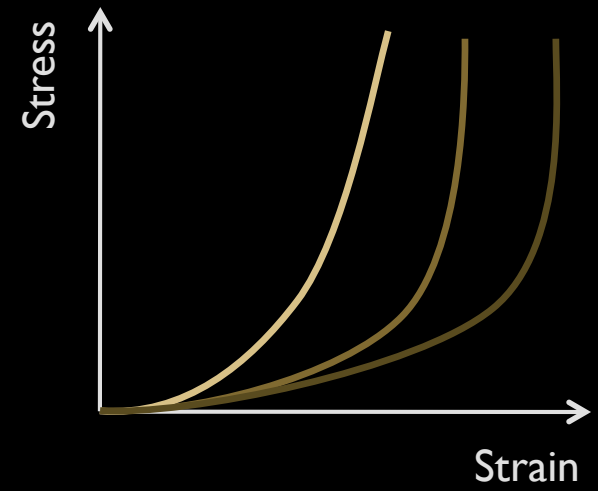
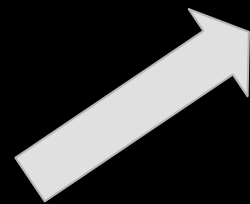
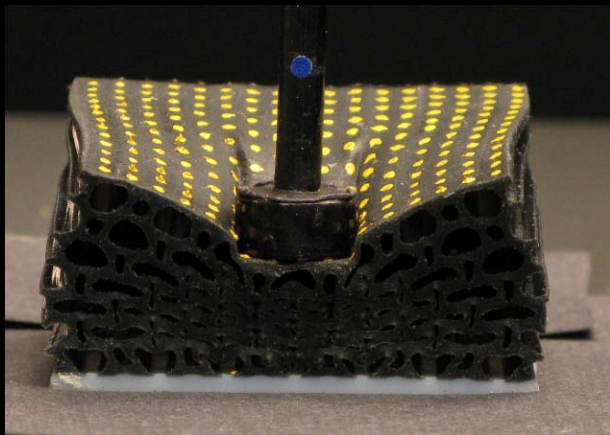
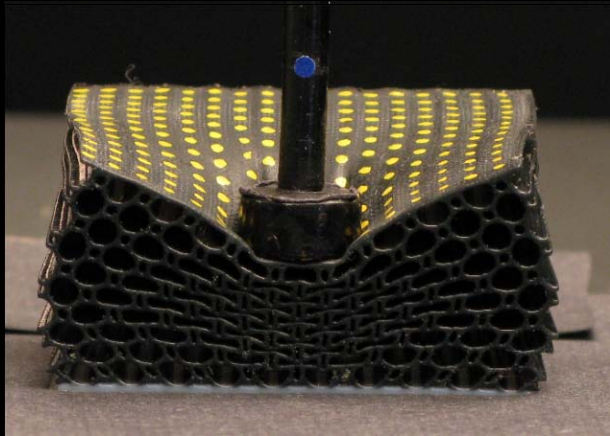
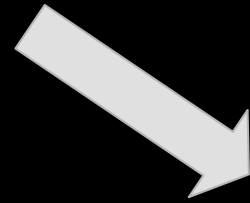
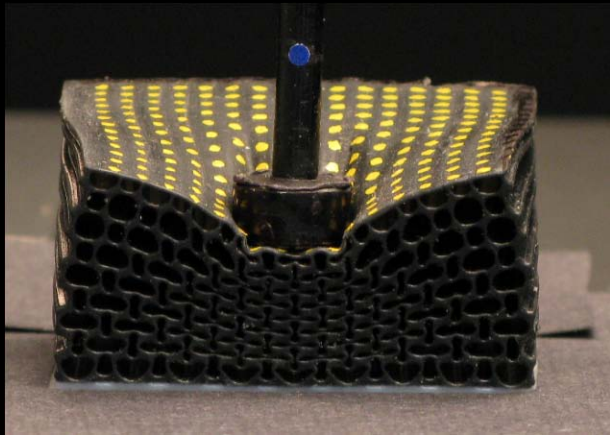


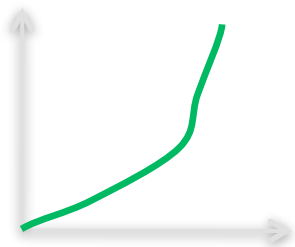
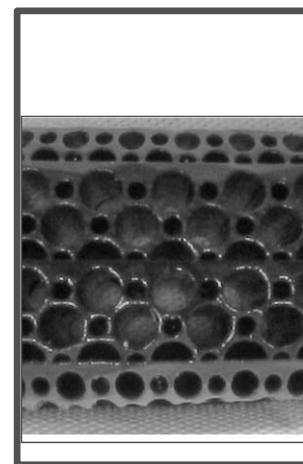
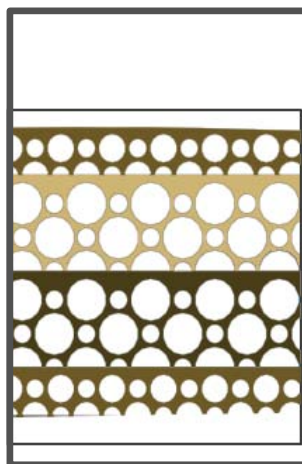
Reproduction







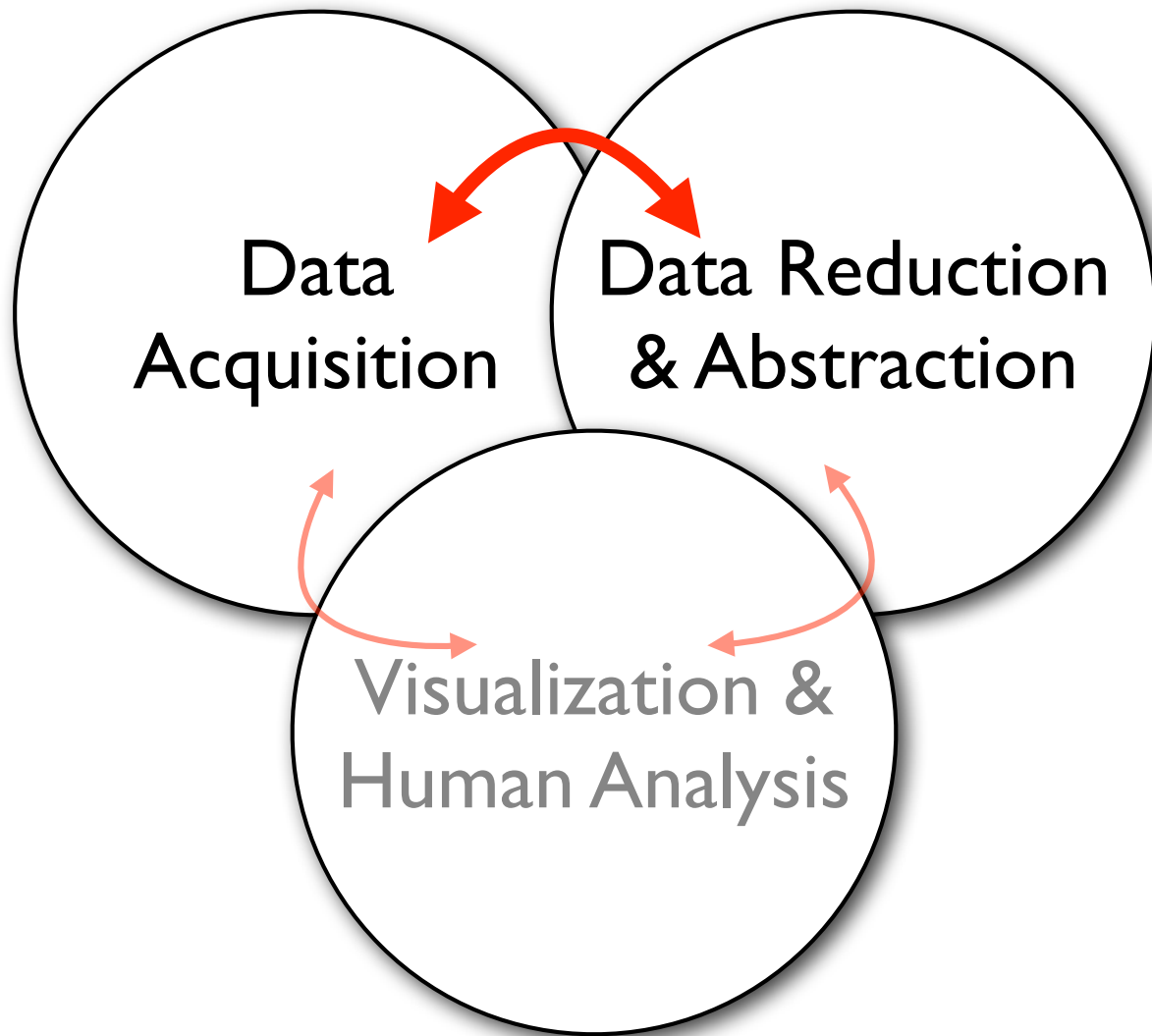




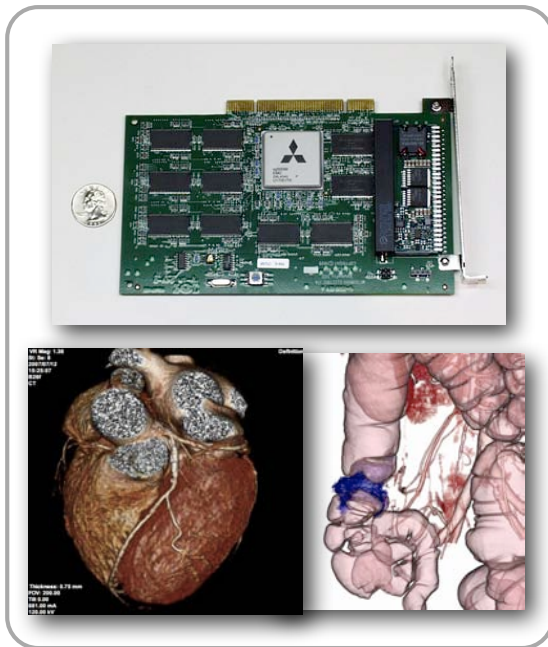


[Bickel et al., SIGGRAPH 2010]

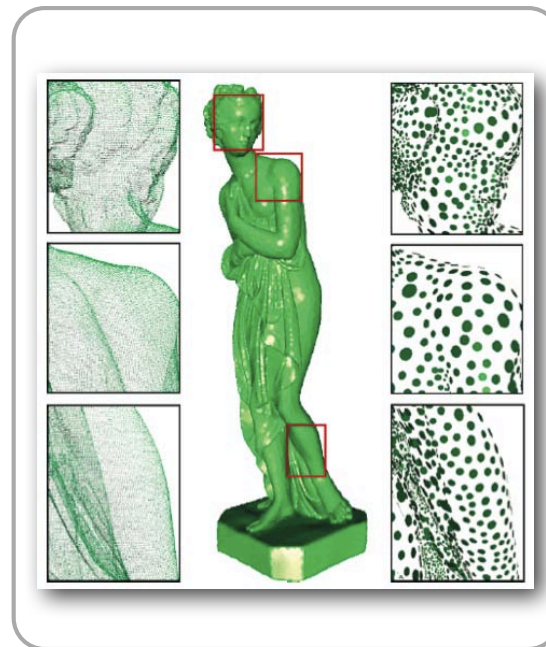
Visual Computing



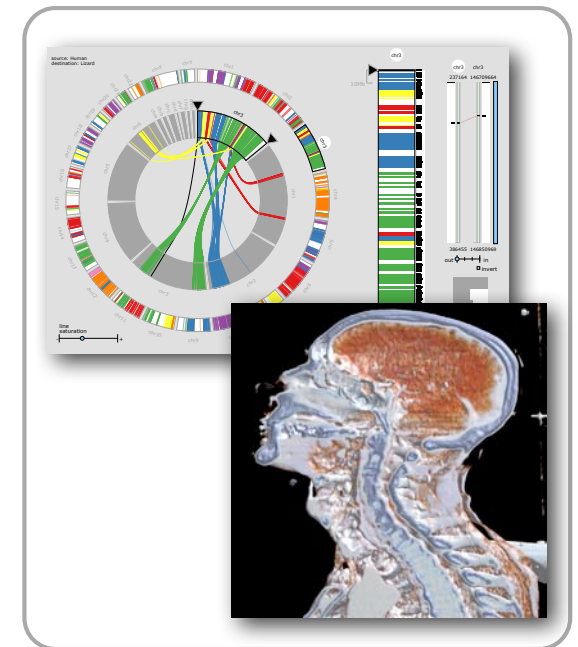
Visualization



VolumePro
Hardware



Point-Based
Graphics



Bio-Medical
Visualization

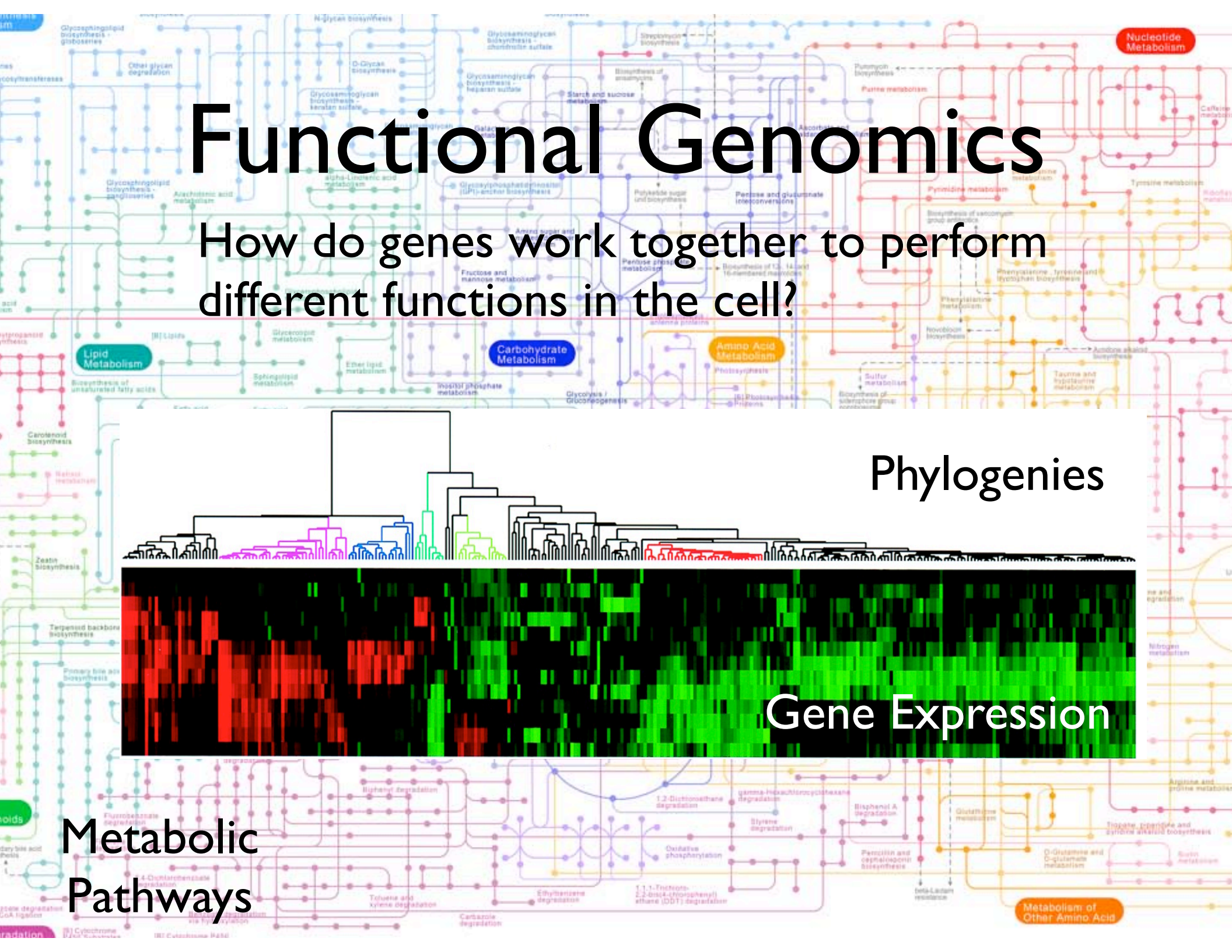
Functional Genomics

How do genes work together to perform different functions in the cell?

Phylogenies

Gene Expression

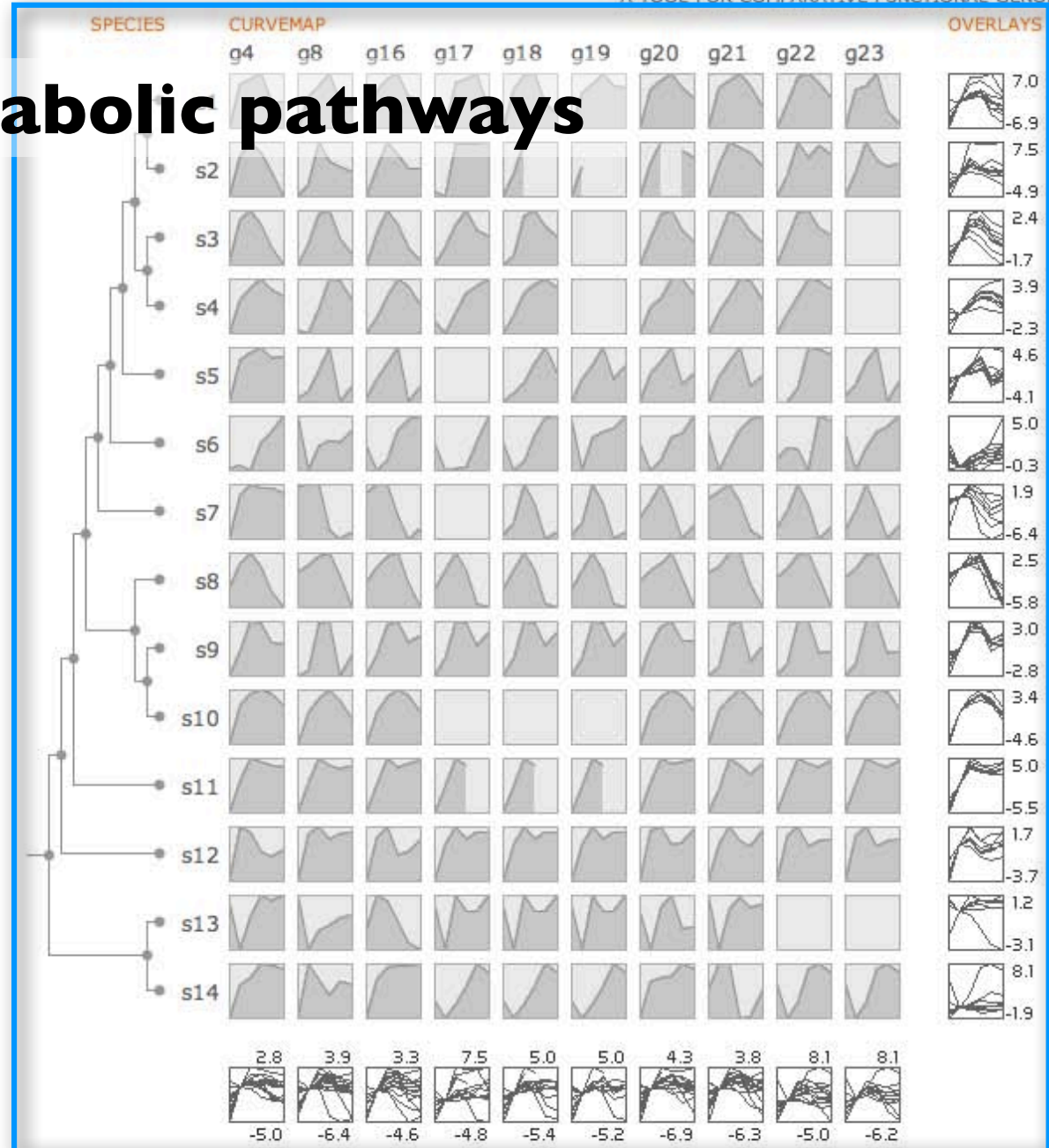
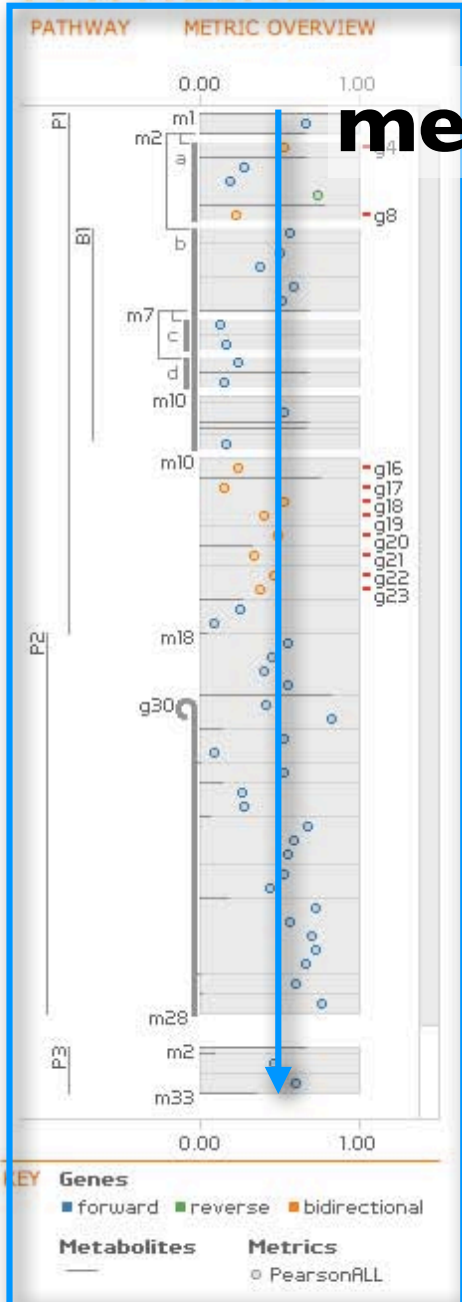
Metabolic Pathways



PATHLINE

A TOOL FOR COMPARATIVE FUNCTIONAL GENOMICS

metabolic pathways



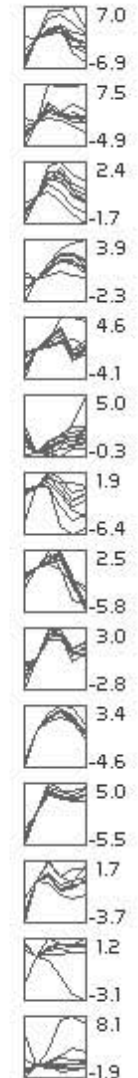
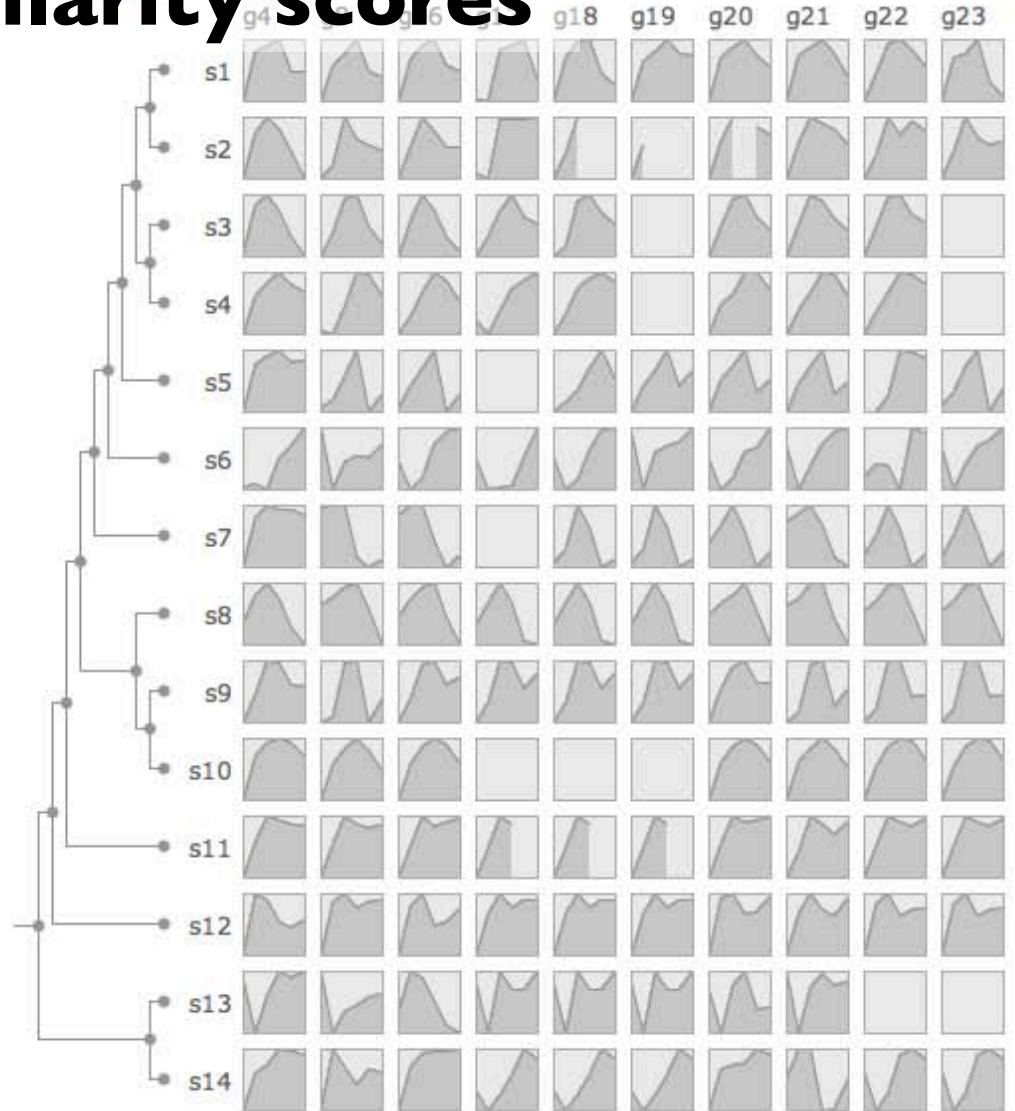
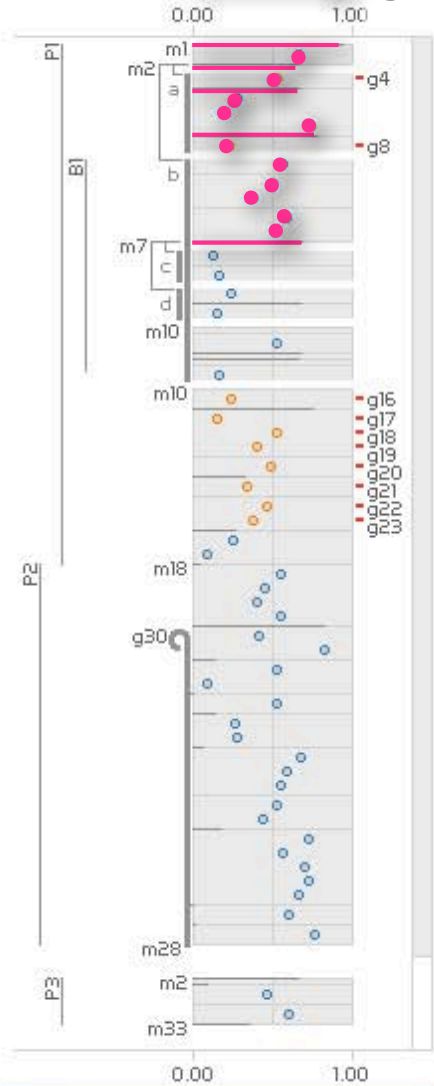
PATHLINE

A TOOL FOR COMPARATIVE FUNCTIONAL GENOMICS

PATHWAY METRIC OVERVIEW

similarity scores

OVERLAYS

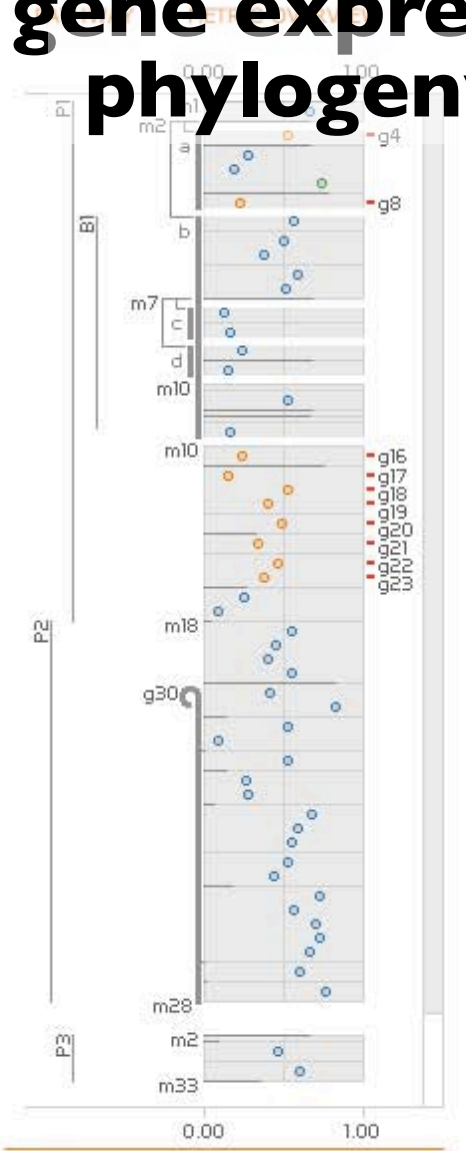


KEY Genes
■ forward ■ reverse ■ bidirectional
Metabolites Metrics
○ PearsonALL

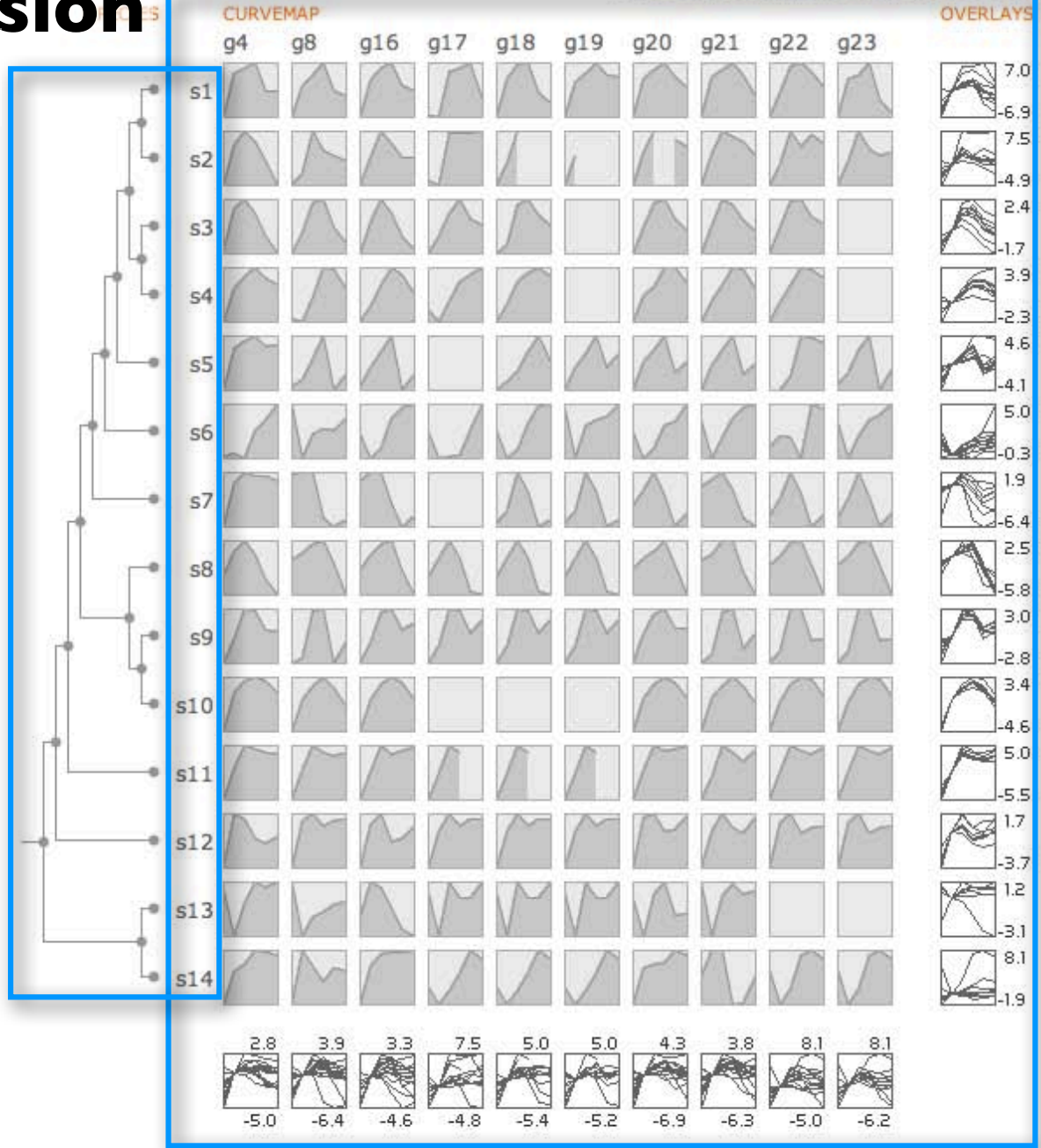
PATHLINE

gene expression phylogeny

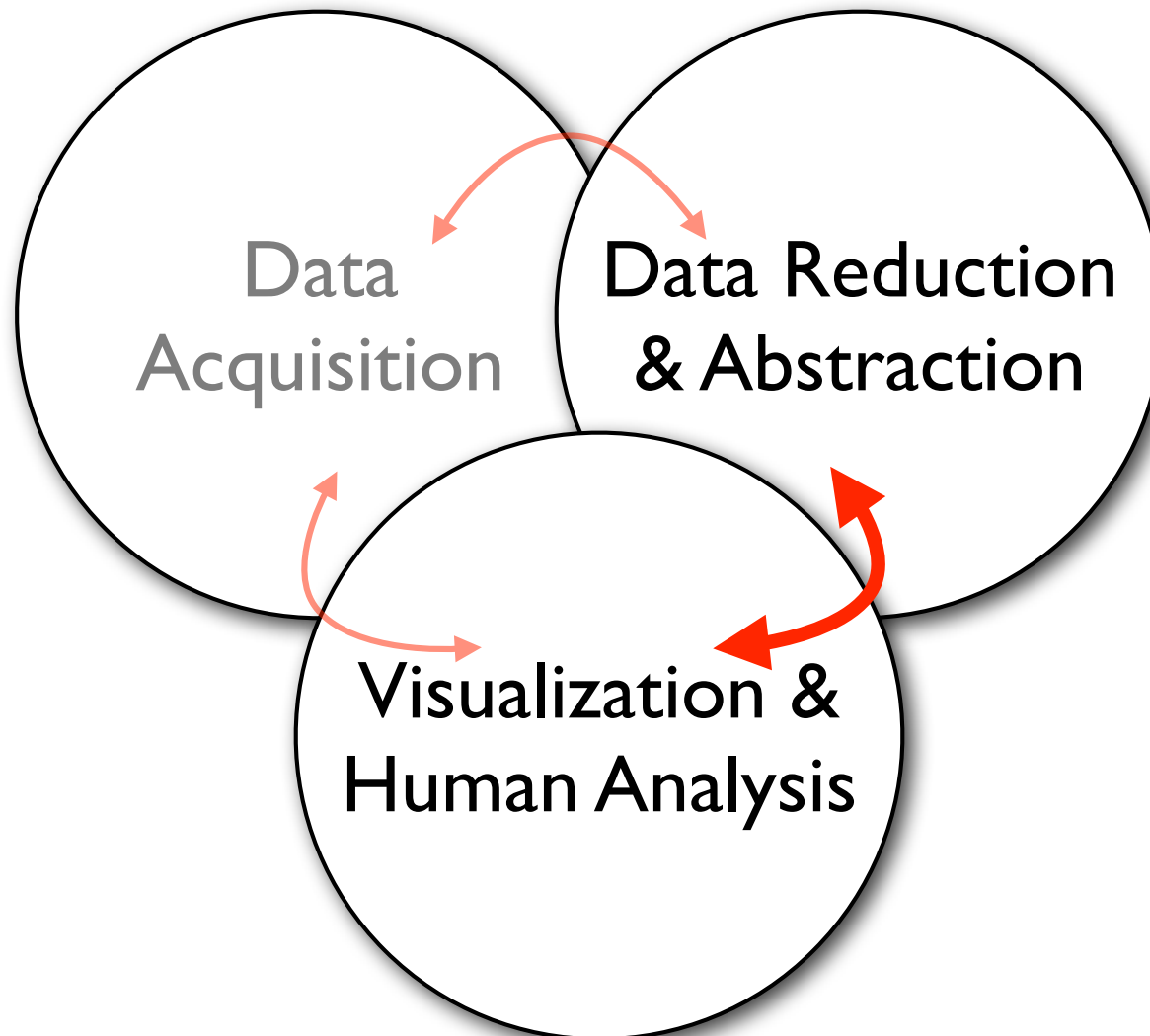
A TOOL FOR COMPARATIVE FUNCTIONAL GENOMICS



KEY Genes
■ forward ■ reverse ■ bidirectional
Metabolites Metrics
○ PearsonALL



Visual Computing



Metabolomics and Integrative Systems Biology Analysis of the
Evolution of the Diauxic Shift in Ascomycota fungi,
M. Styczynski et al., in preparation.

Visual Computing Group

Dr. Won-Ki Jeong
Dr. Verena Kaynig
Dr. Miriah Meyer
Moritz Baecher
Michelle Borkin
Kevin Dale
Amanda Peters
Mike Roberts
Kalyan Sunkavalli
Amelio Vazquez





The Connectome

Discovering the Wiring Diagram of the Brain

Collaborators

Harvard Center for Brain Science

- Prof. Jeff Lichtman & Prof. Clay Reid



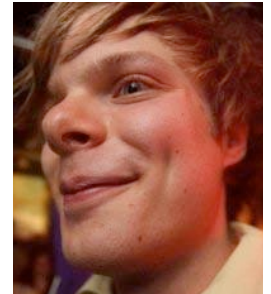
KAUST

- Prof. Markus Hadwiger, Dr. Johanna Beyer



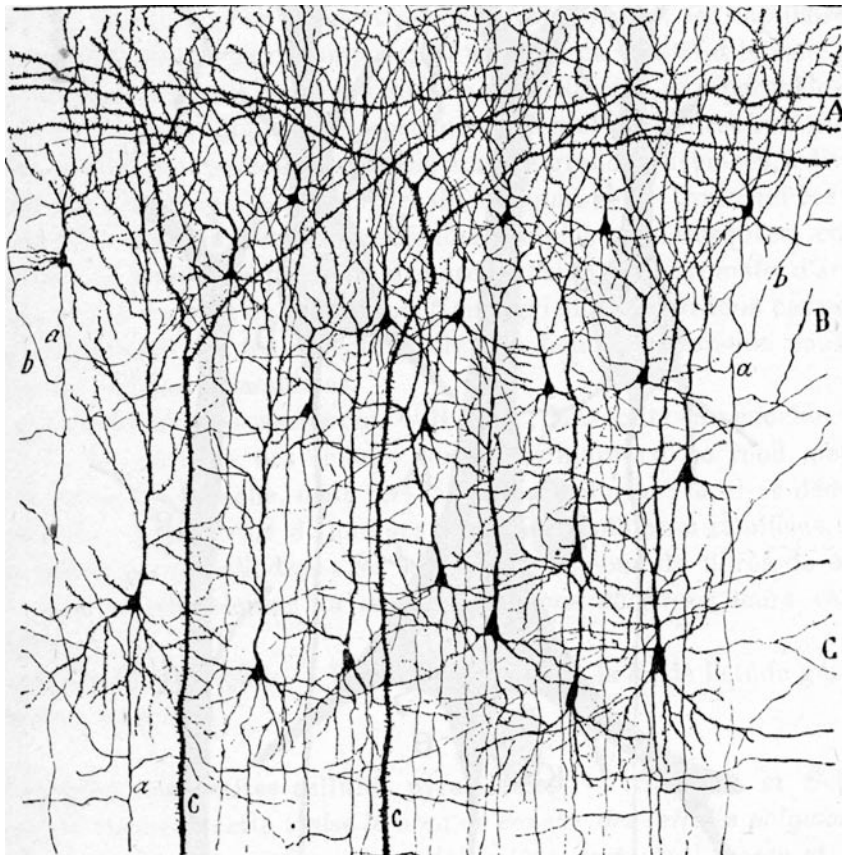
SEAS

- Dr. Won-Ki Jeong
Dr. Verena Kaynig-Fittkau
Amelio Vazquez
Mike Roberts
N

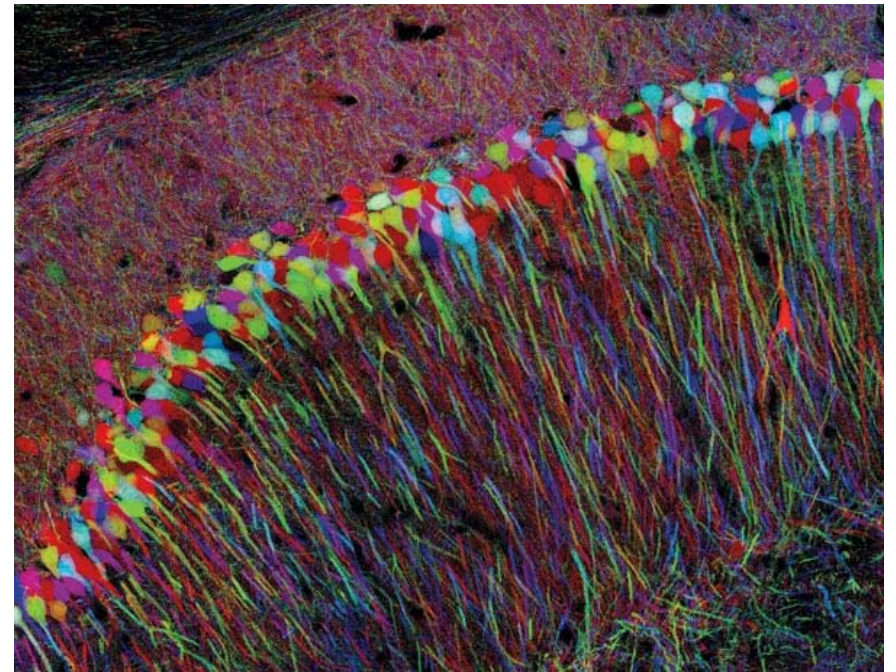


The Scientific Challenge

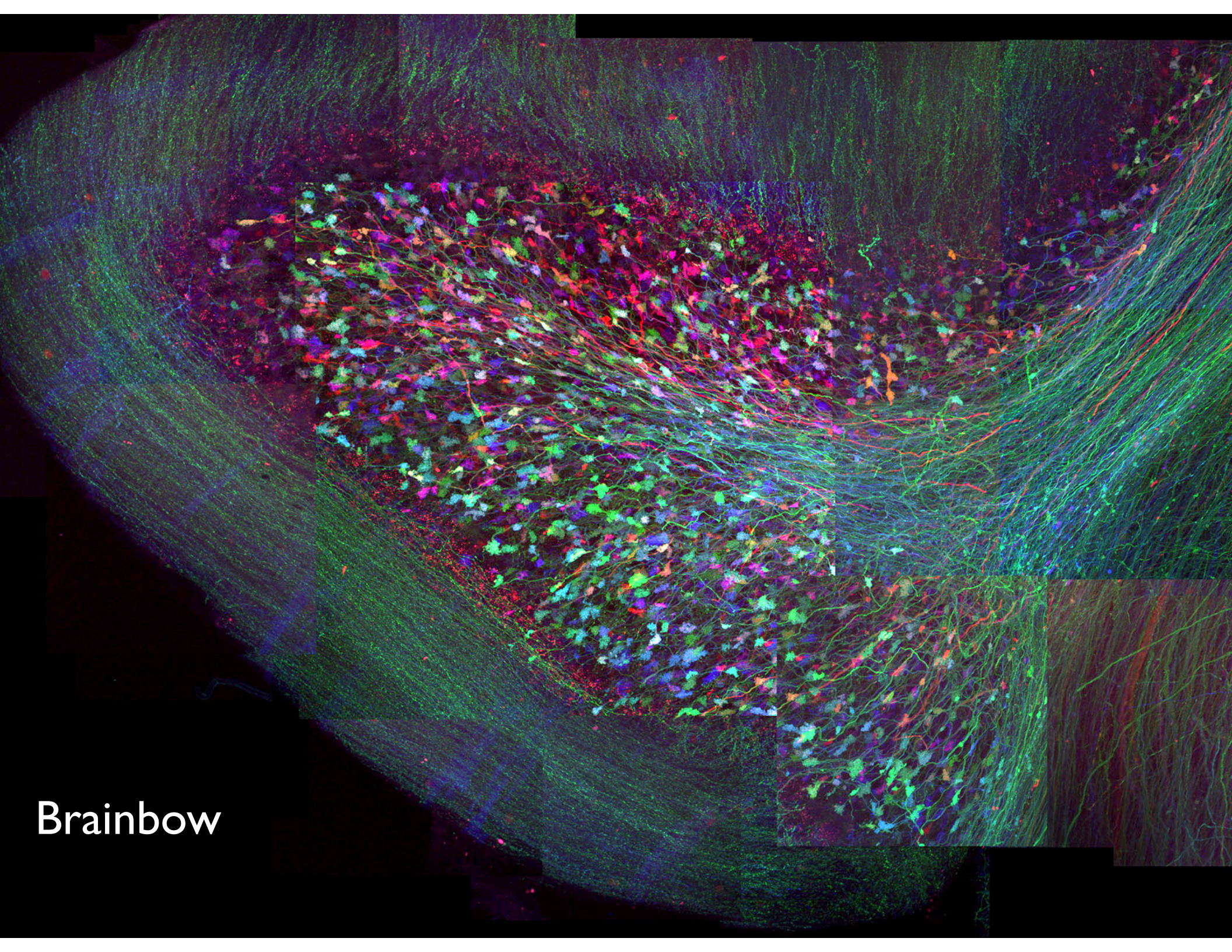
How is the mammalian brain wired?



Ramón y Cajal, 1905

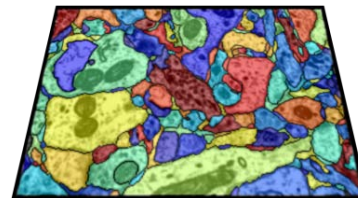
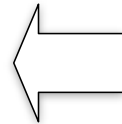
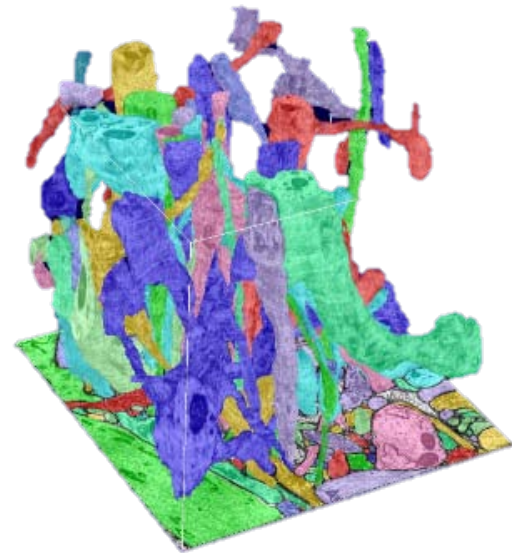
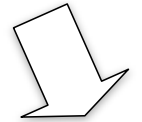
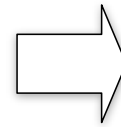
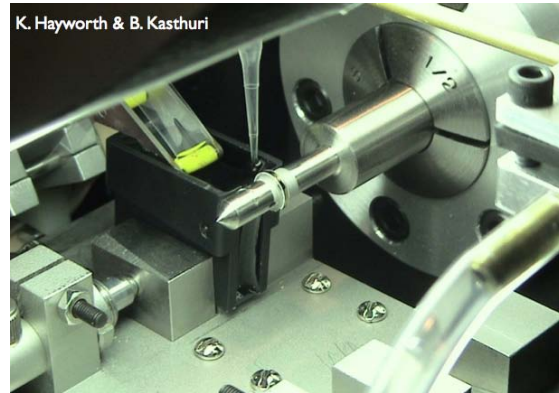
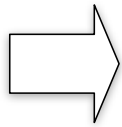
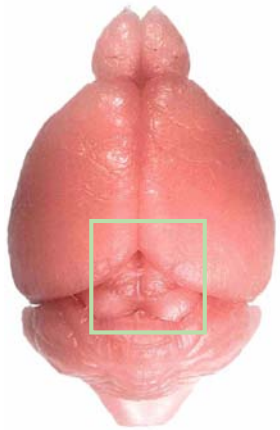


Harvard Center for Brain Science

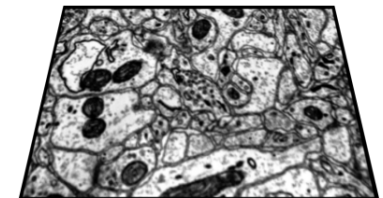
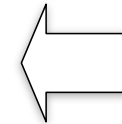
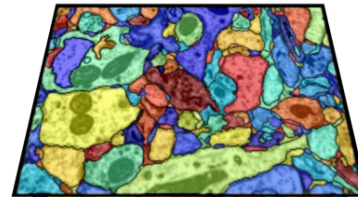


Brainbow

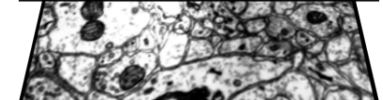
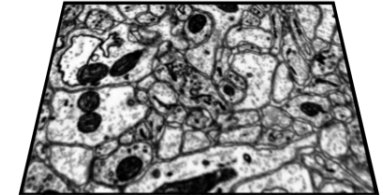
Connectome Workflow



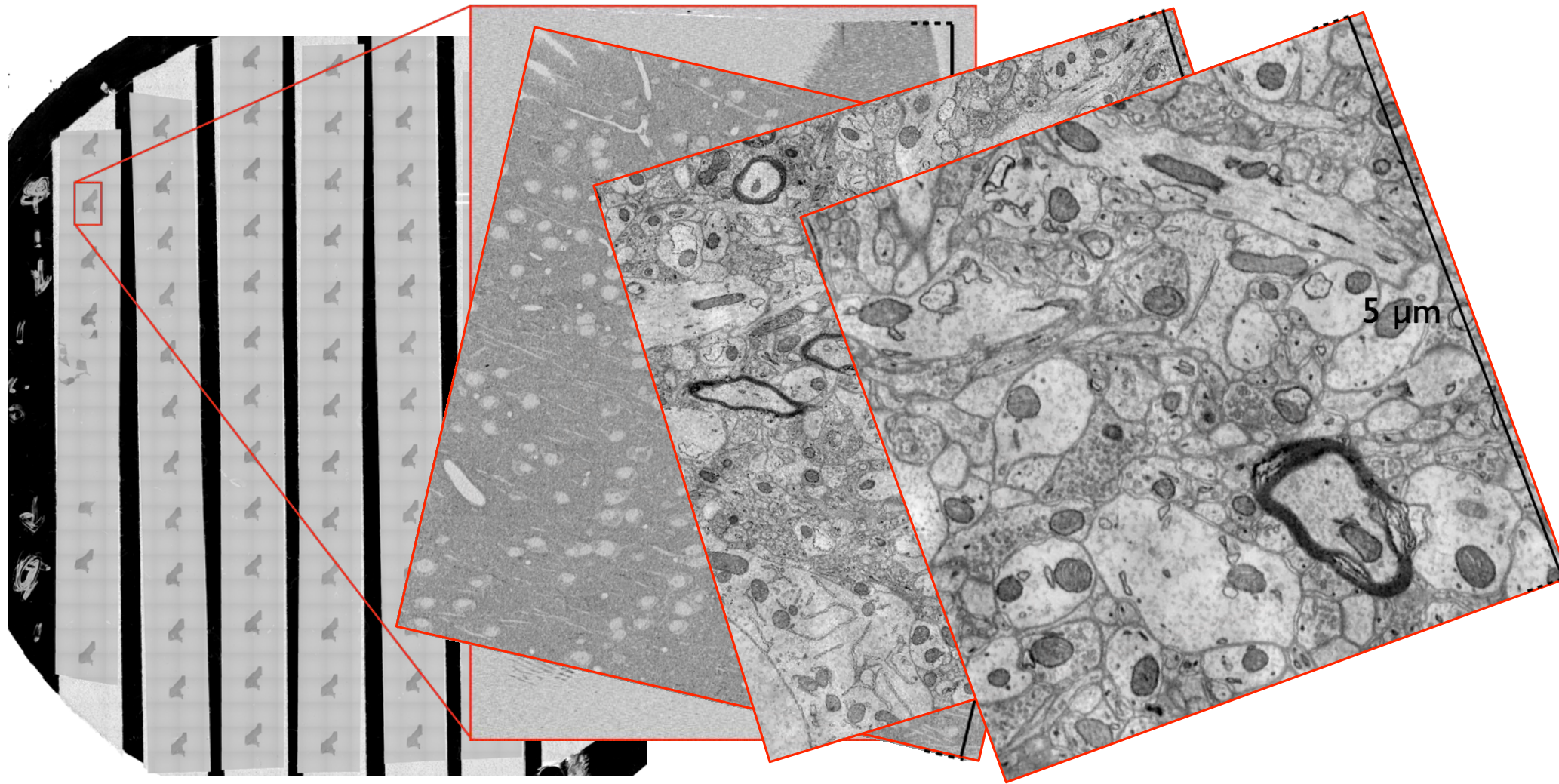
⋮



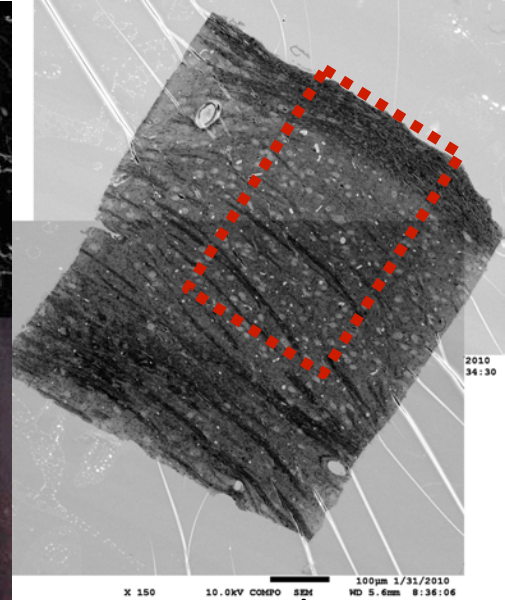
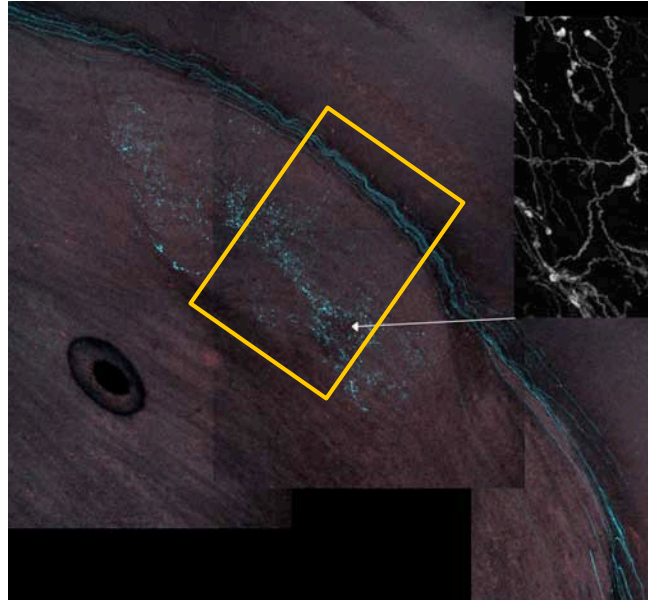
⋮



Electron Microscopy



The Data Challenge



Josh Morgan

- Pixel resolution: 3-5 nm
- Slice thickness: 30-50 nm
- 1 mm³: 200k x 200k pixels x 20k slices
40 Gpixels x 20k = 800 TB

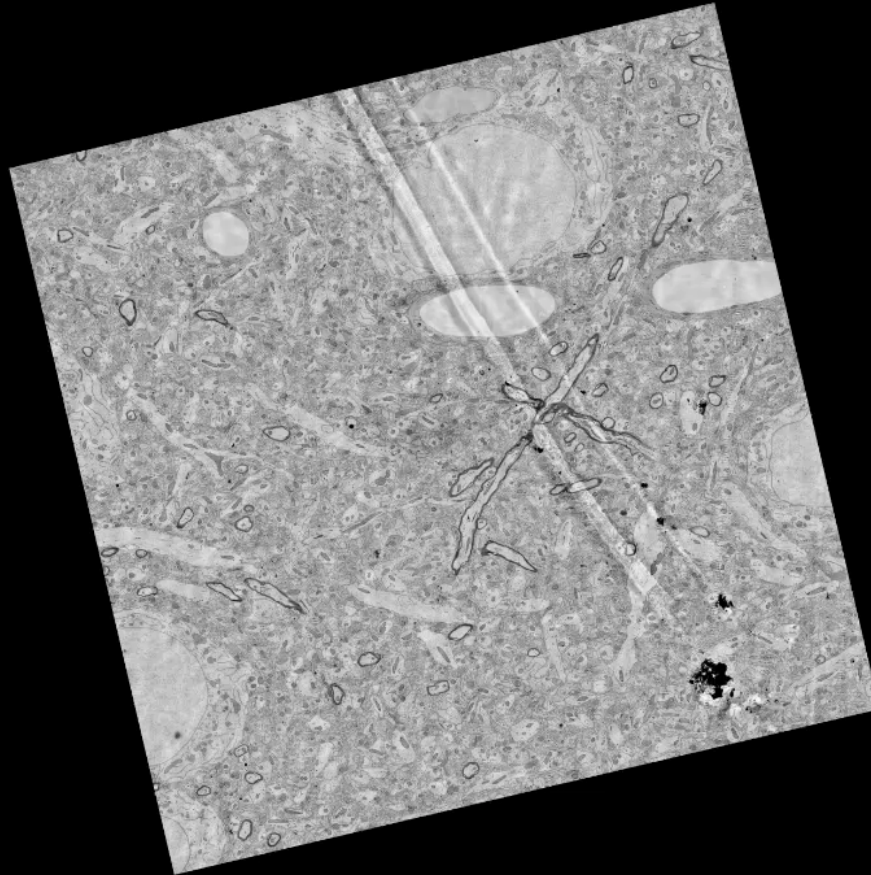
CS Challenges

- Stitching and alignment of overlapping tiles
- 3D registration of sections
- Visualization
- 3D reconstruction
- Synapse detection
- Network analysis

CS Challenges

- Stitching and alignment of overlapping tiles
 - 3D registration of sections
 - **Visualization**
 - **3D reconstruction**
 - Synapse detection
 - Network analysis
- } This Talk

Neurotrace



Interactive navigation of 1TB brain dataset

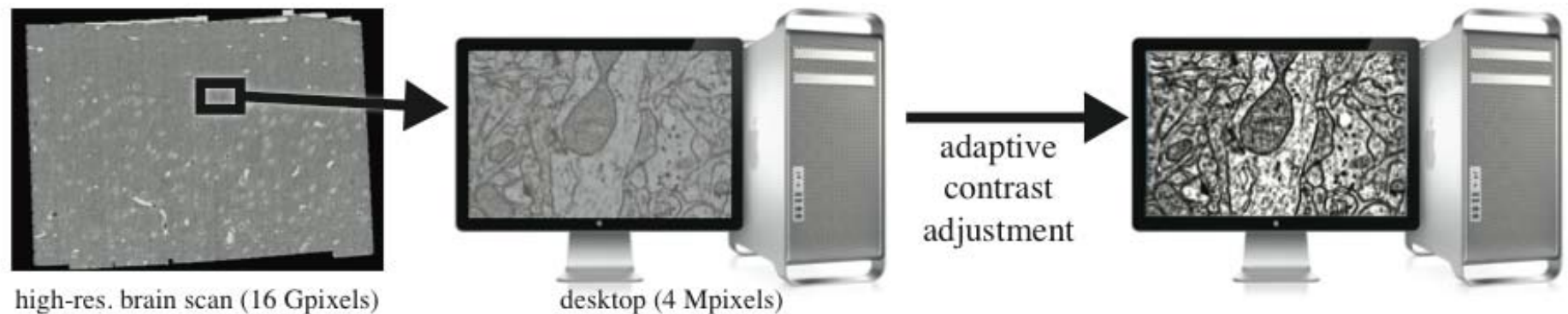
Existing Methods

- Global image pyramid (e.g., Photoshop)
- Preprocessing required
- Not optimal for integration of new image tiles



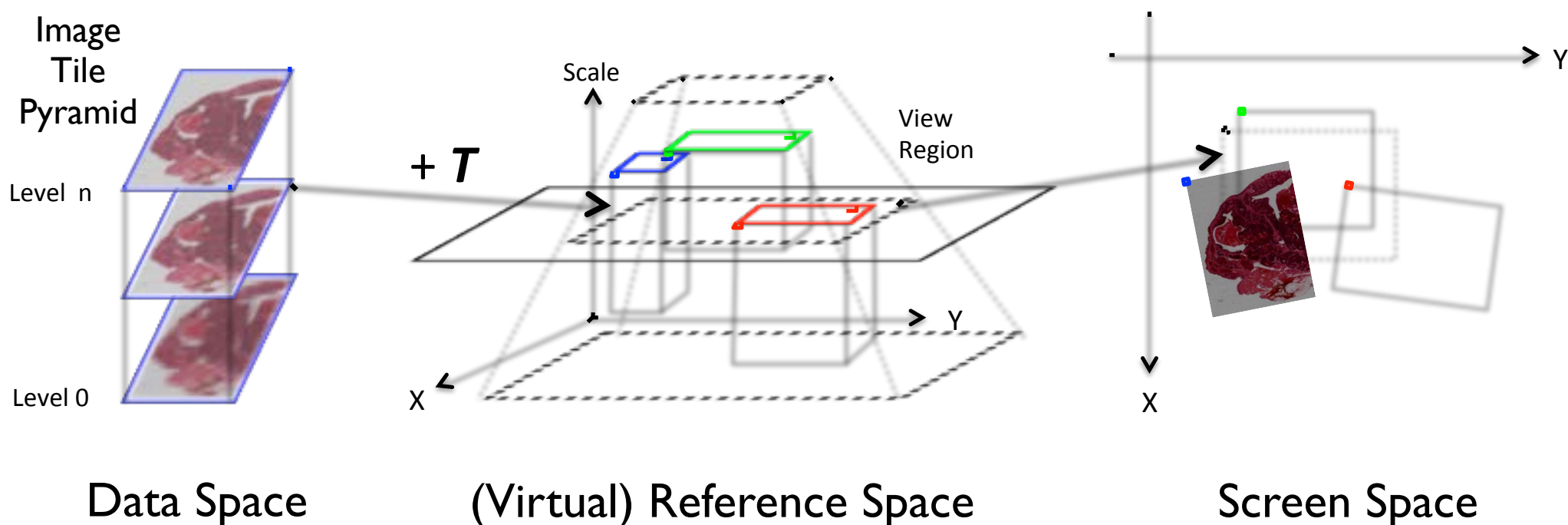
Display-Aware Processing

- Take into account and process only the visible pixels
- Design choice: Displayed image computed on-the-fly each time



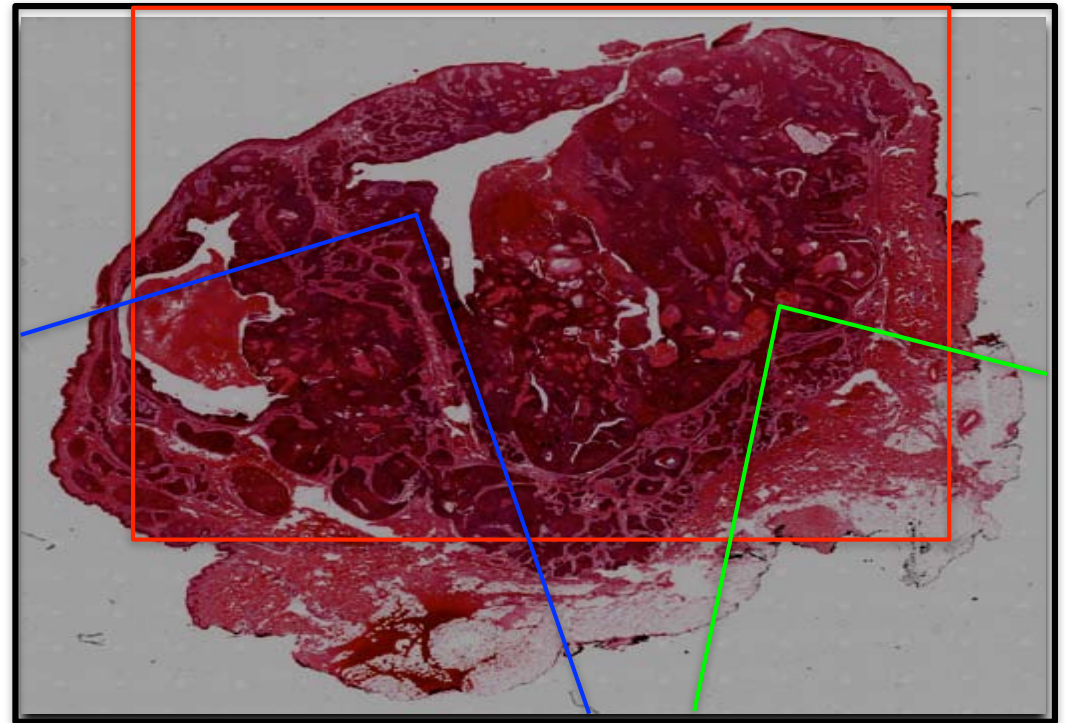
Reference Space

- Virtual coordinate system for images
- On-the-fly resampling of *only* visible pixels

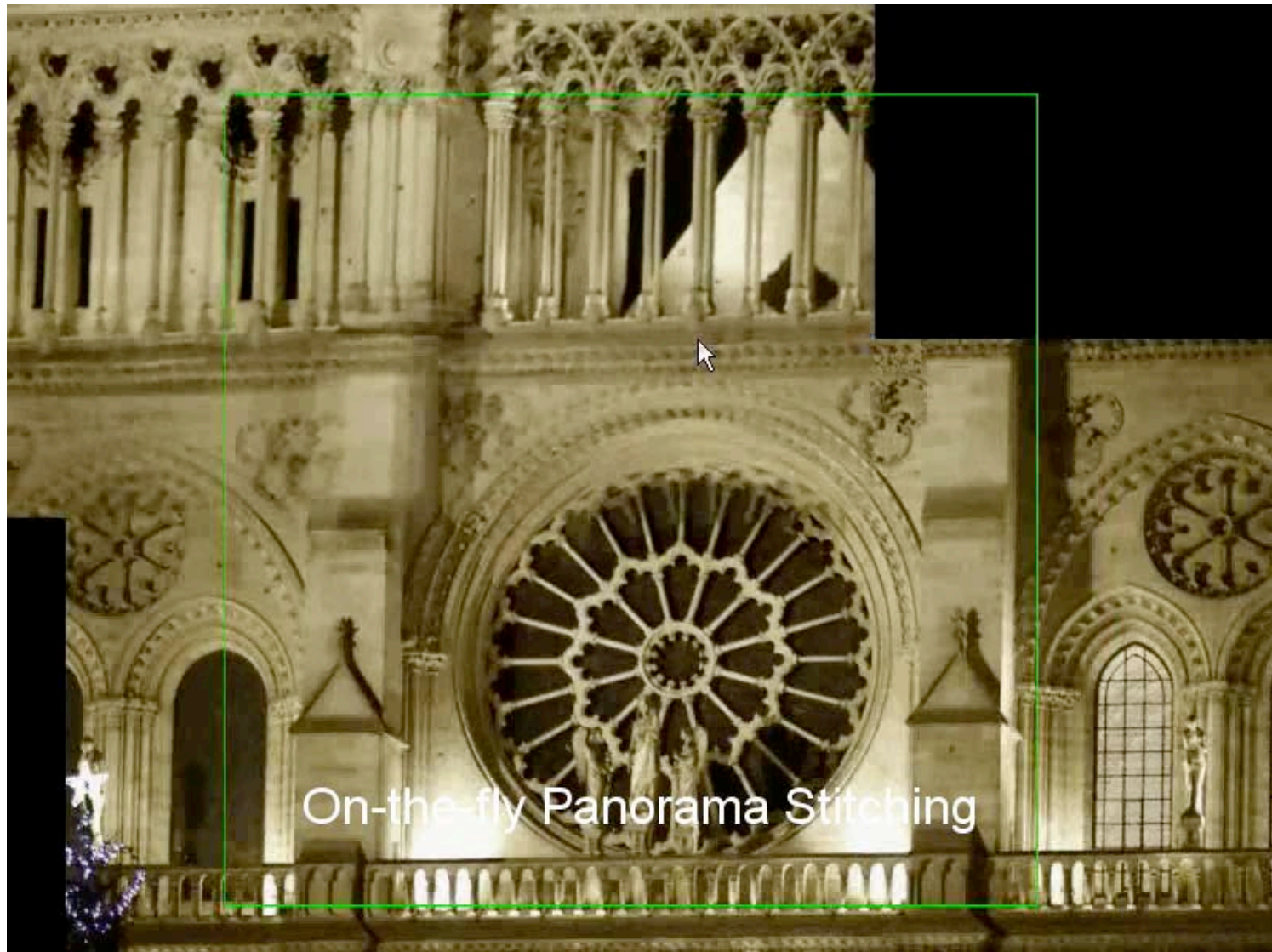


Display Space

- GPU-accelerated resampling
- Display-aware operations
 - Alignment
 - Blending
 - Color editing

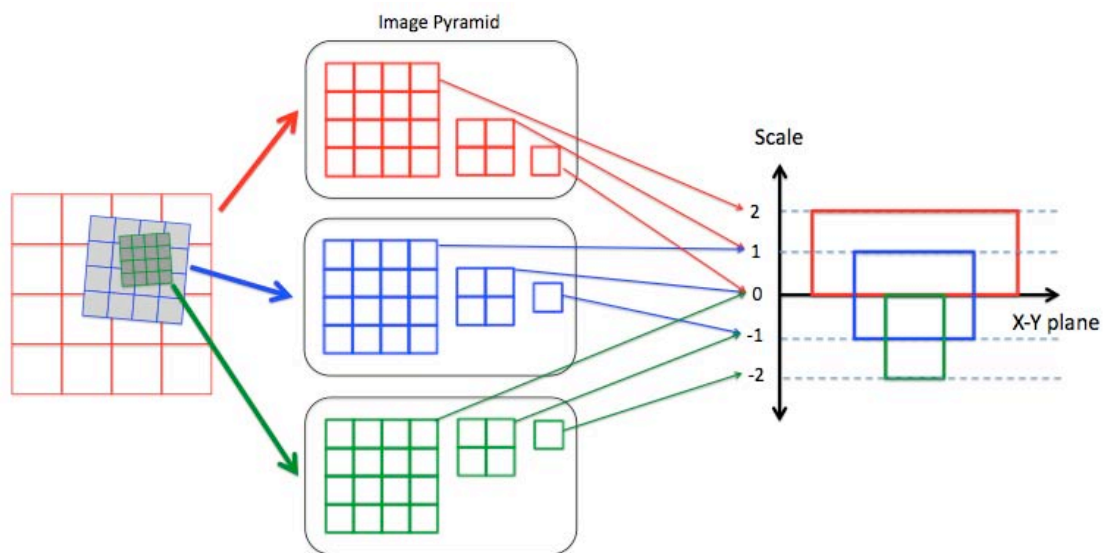
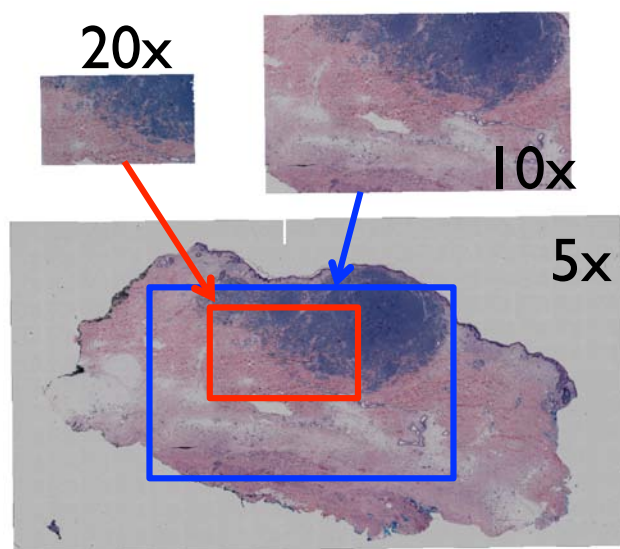


GPU Image Alignment

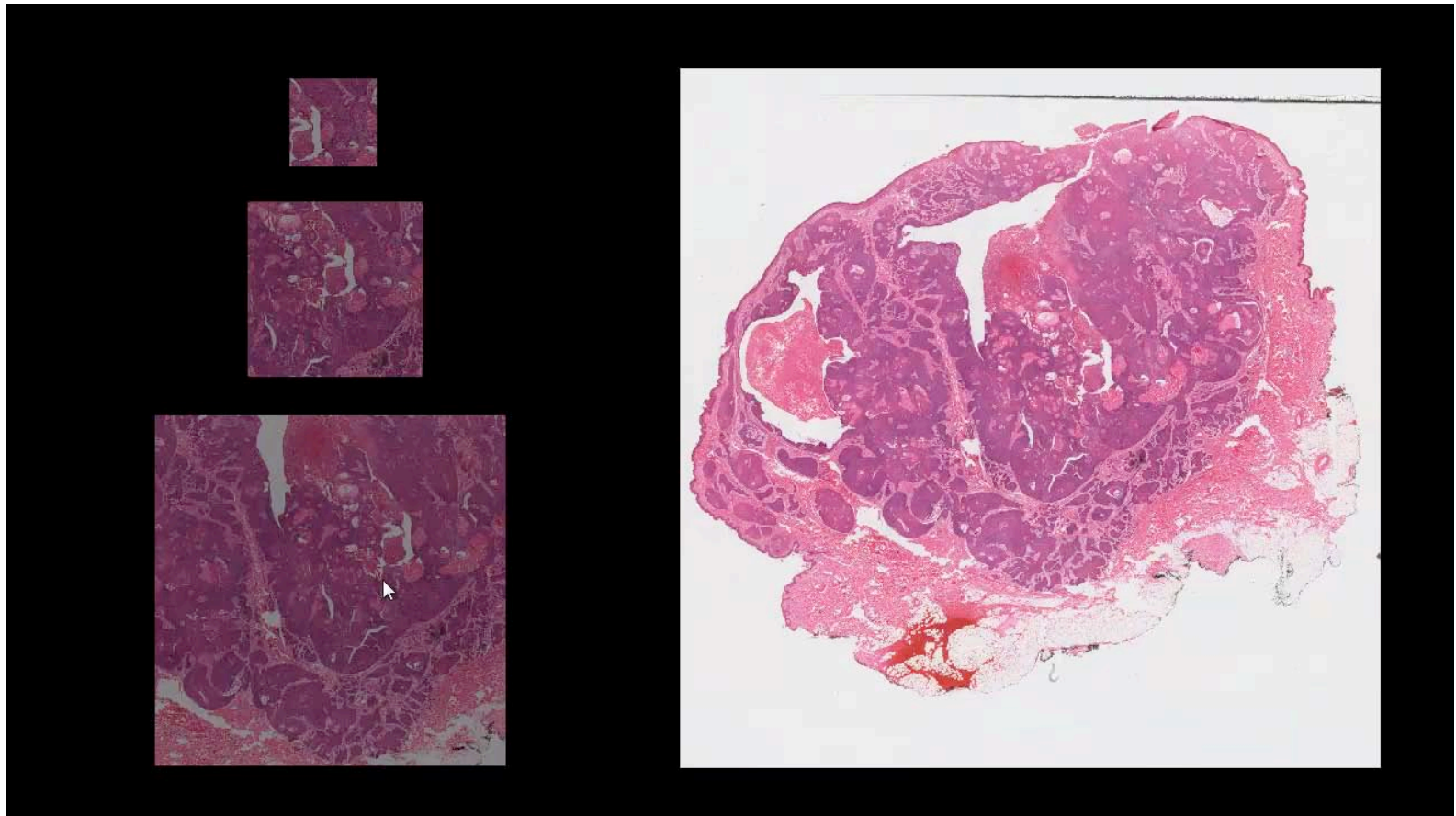


Adaptive Image Hierarchy

- Nested images with varying *resolutions*

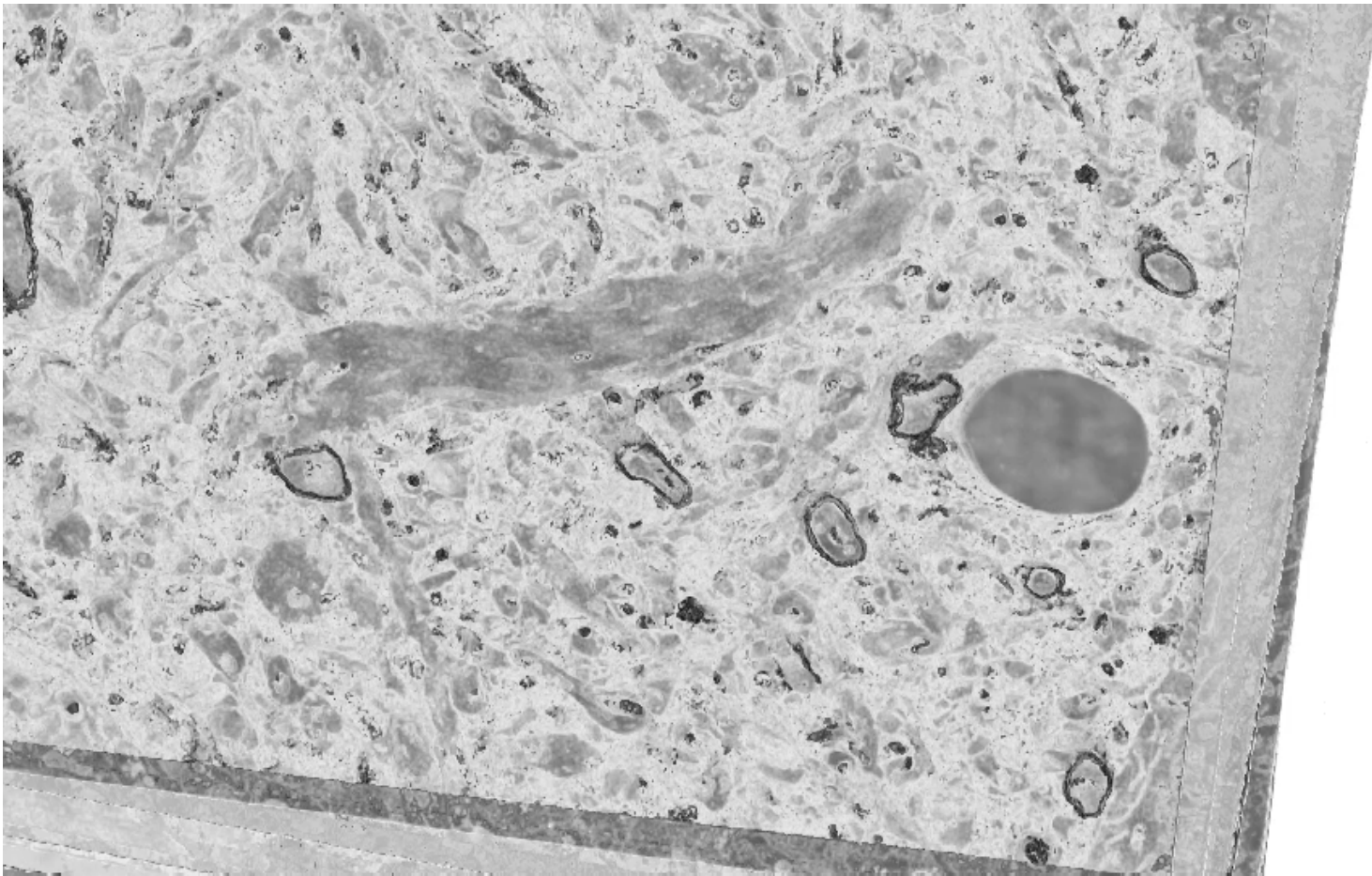


Adaptive Image Hierarchy



Volume Visualization

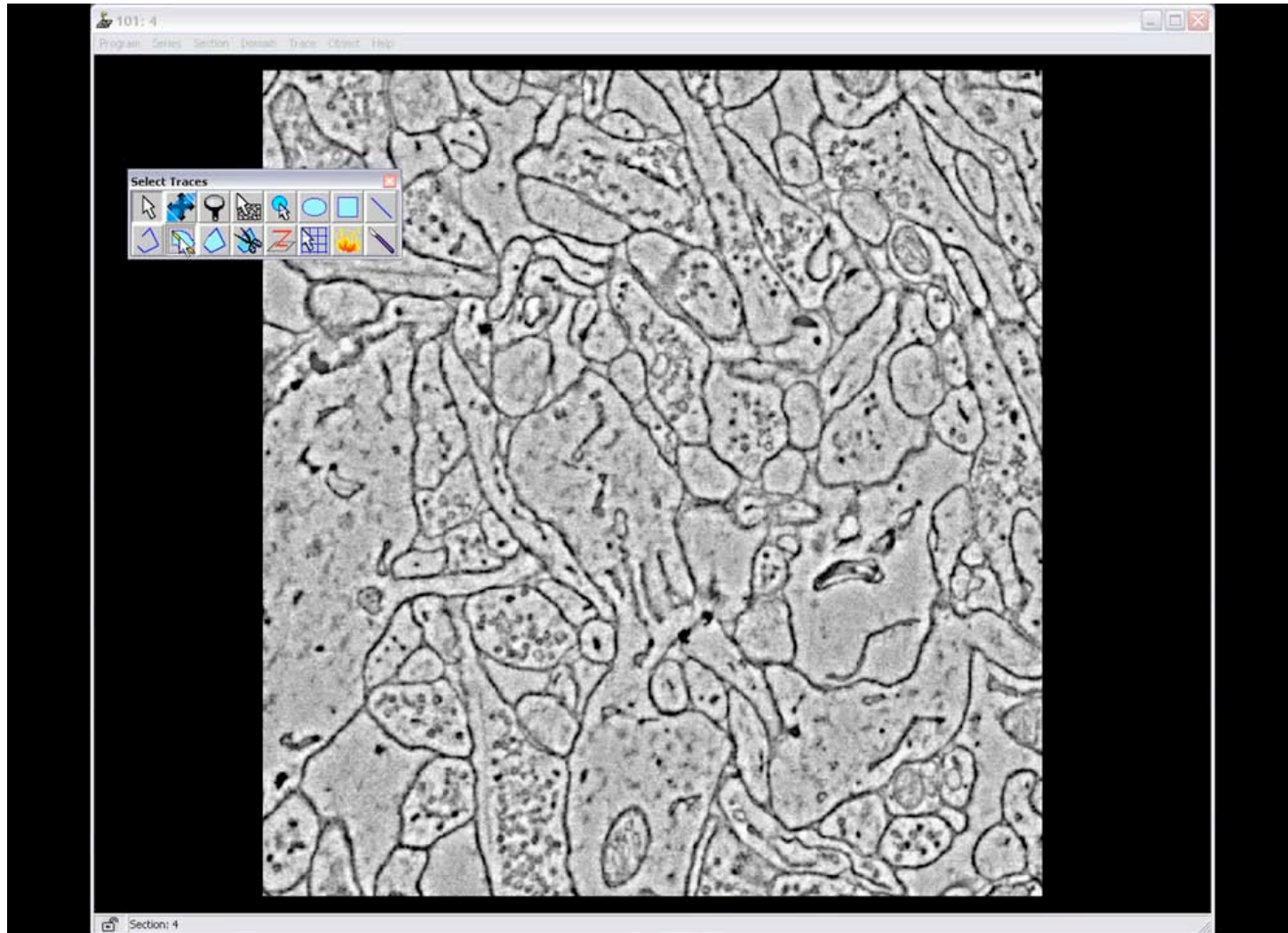
21,494 X 25,790 X 1850 = 955 GB



CS Challenges

- Stitching and alignment of overlapping tiles
- 3D registration of sections
- Visualization
- **3D reconstruction**
- Network analysis

Manual Reconstruction



[Reconstruct]

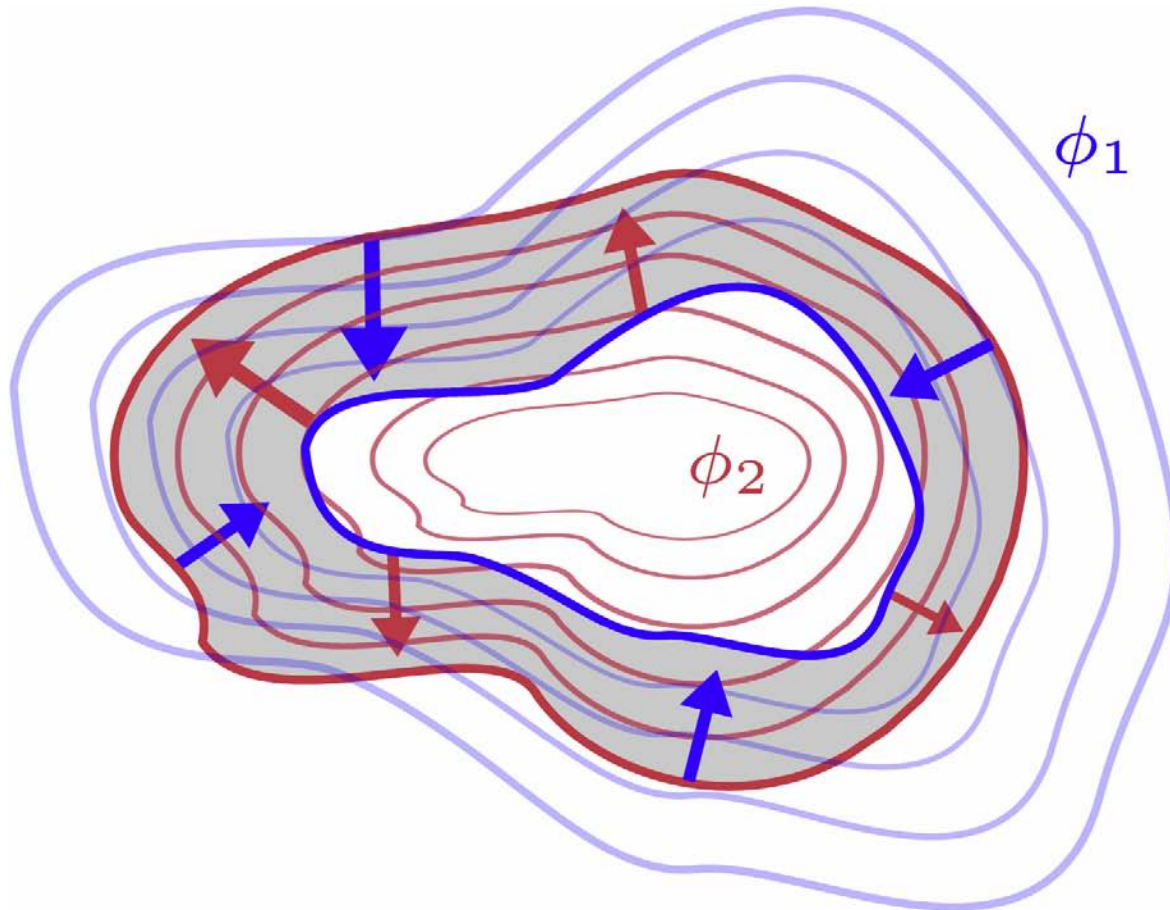
Active Ribbons



Section 1

Active Ribbons

Implicit deformable model based on multiphase level sets and force fields

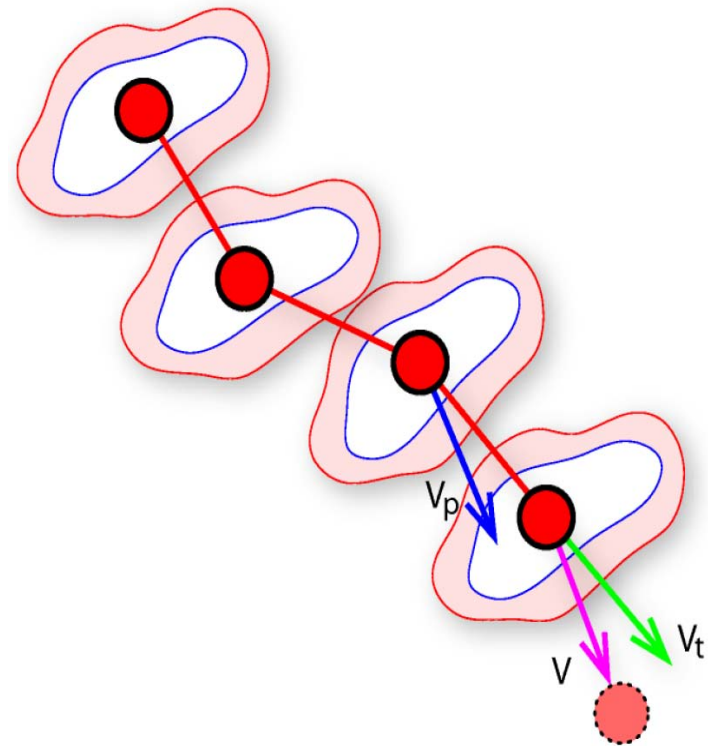


$$v_1 = \frac{\nabla \phi_2}{\sigma_1}$$

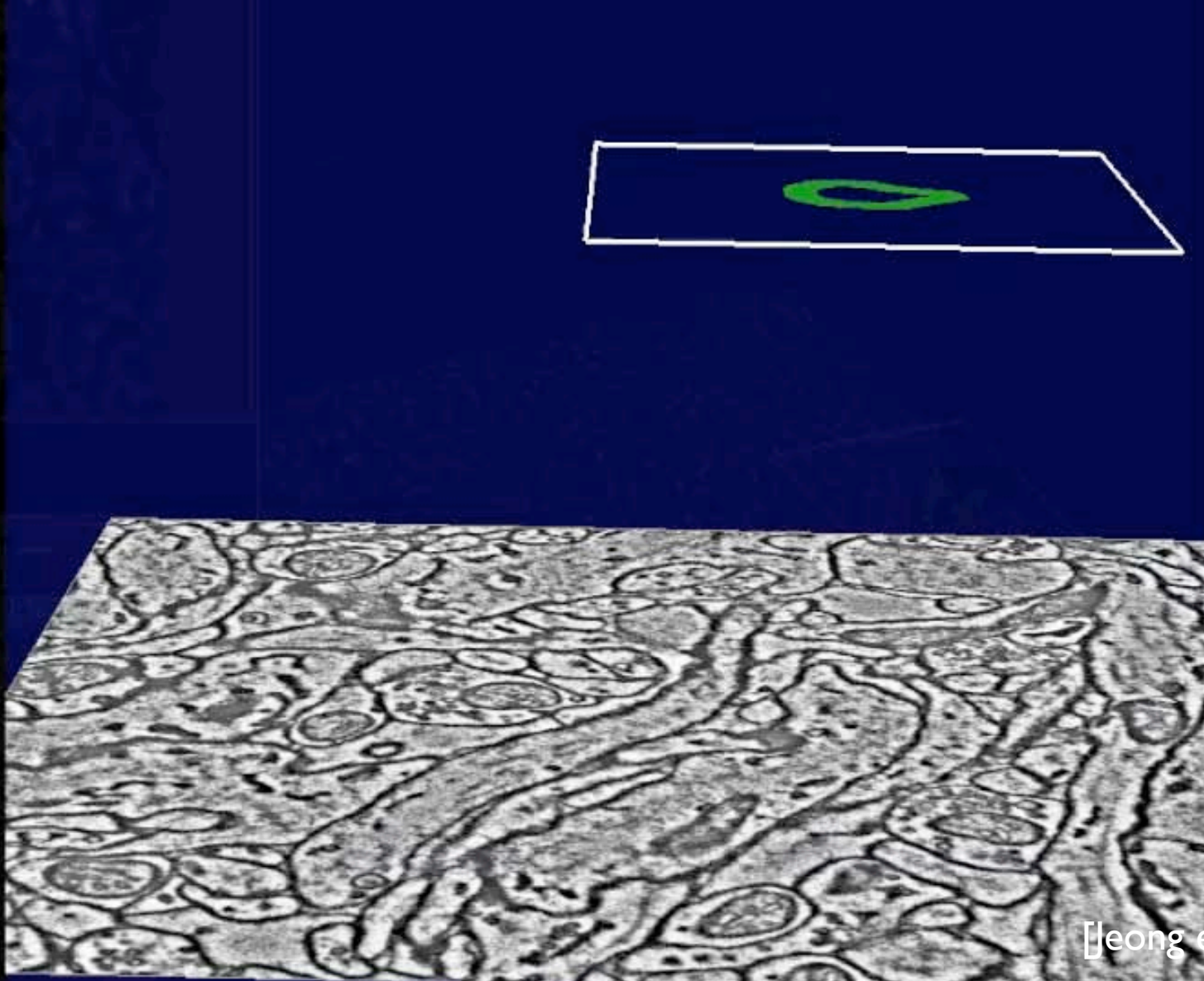
$$v_2 = -\frac{\nabla \phi_1}{\sigma_2}$$

GPU Processing

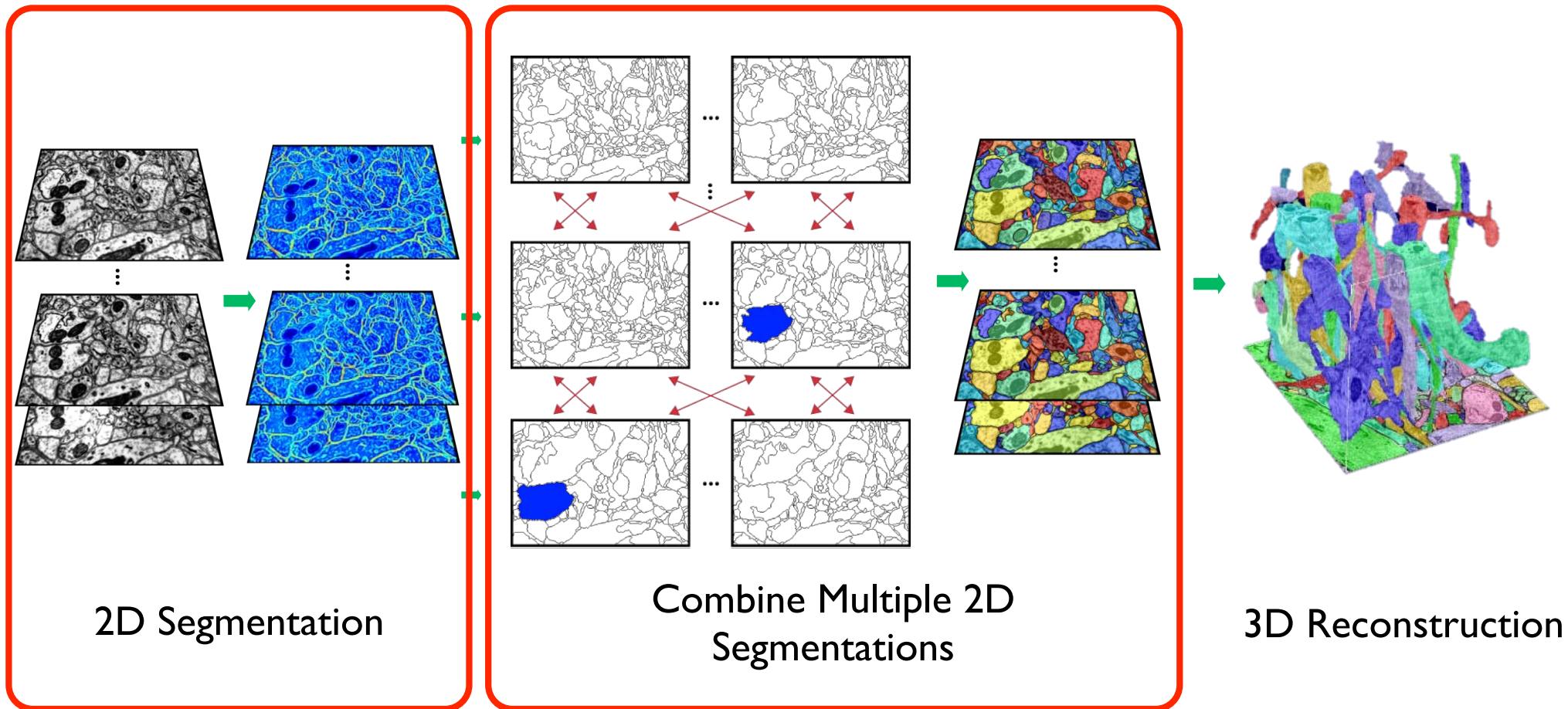
Real-time editing and 3D tracking, 0.5 s/section



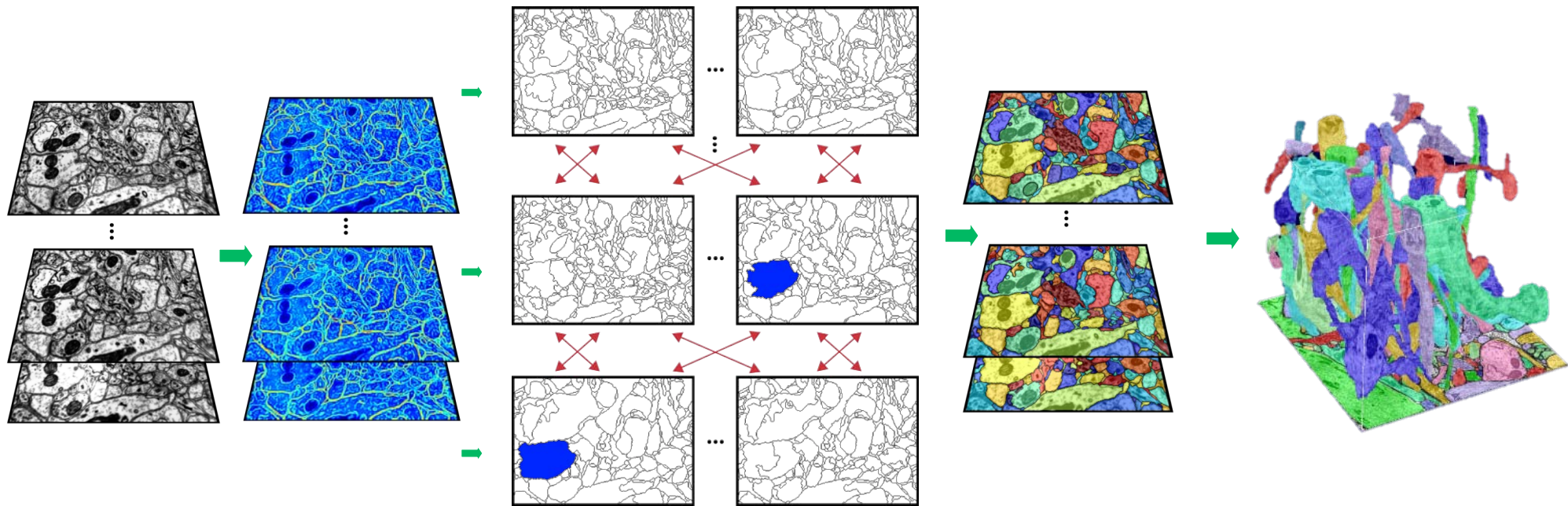
User-Guided Reconstruction



Automatic Reconstruction



Automatic Reconstruction

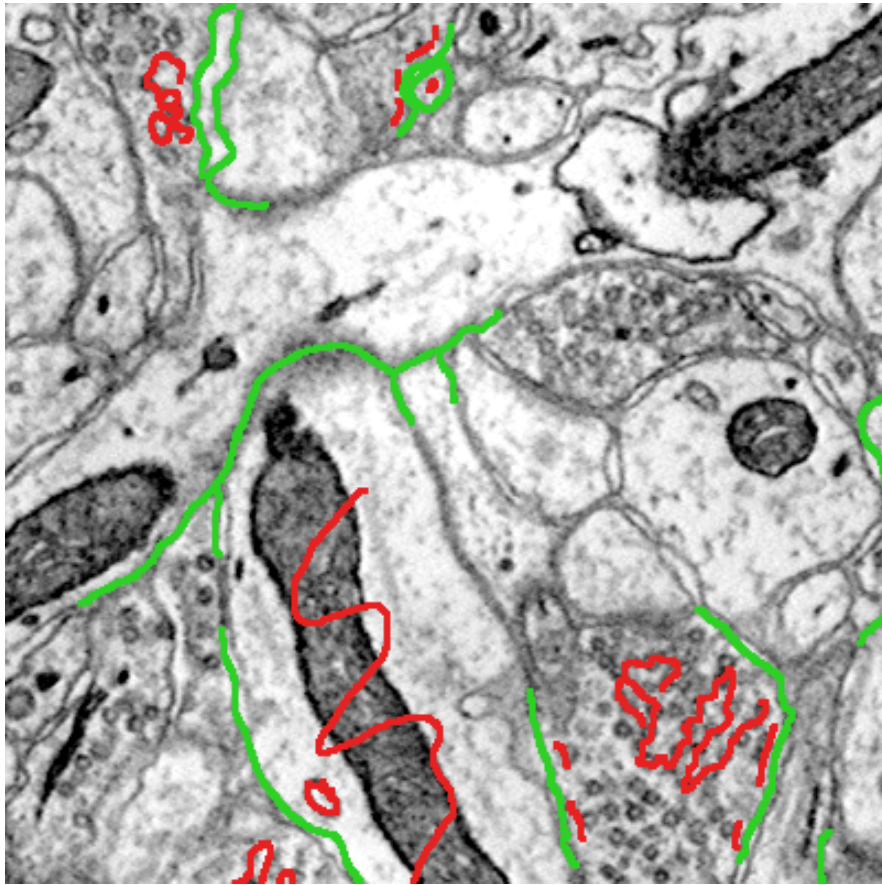


Membrane
probability maps

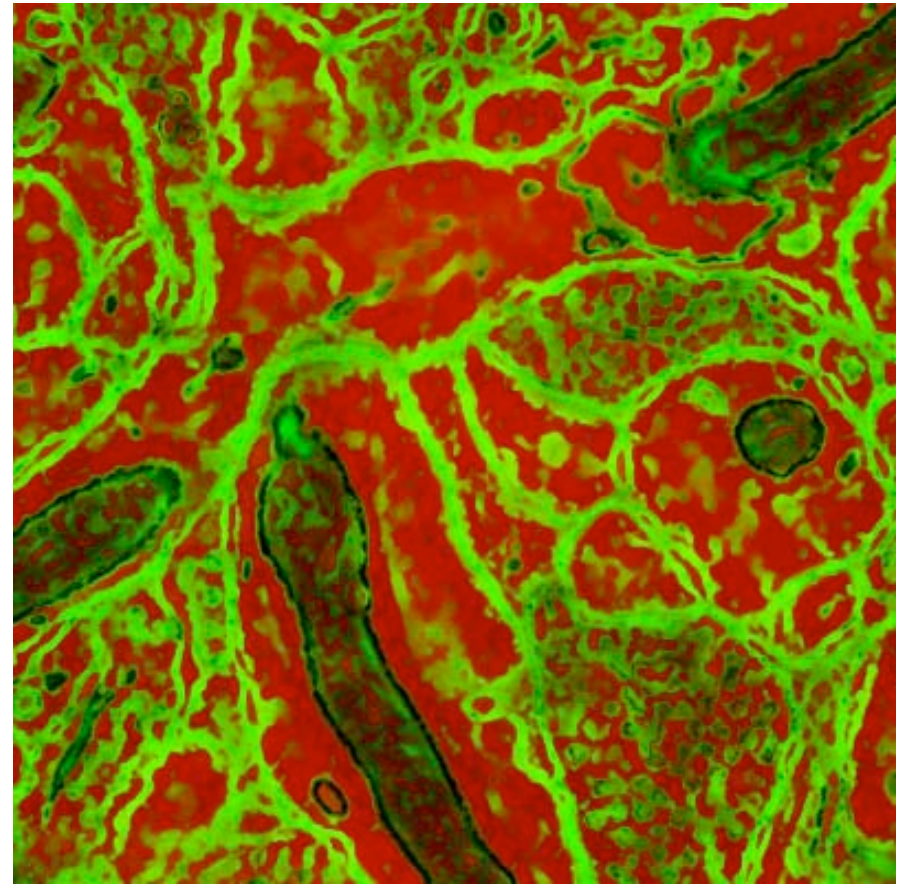
Enumerate and
evaluate combinations

Globally consistent
3D reconstruction

Sparse Interactive Annotation

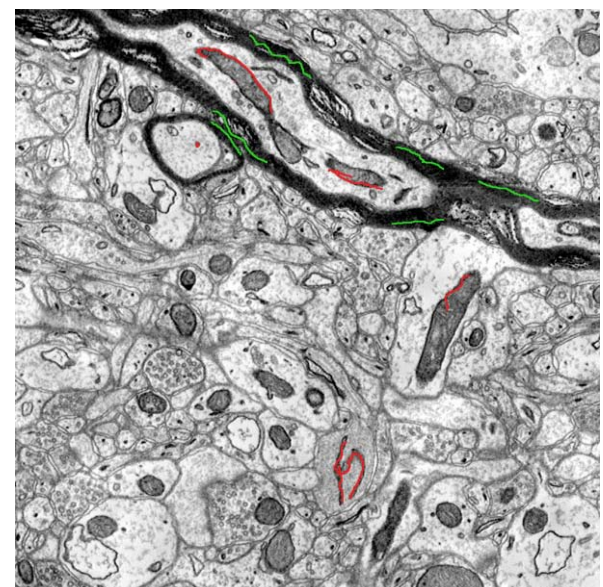
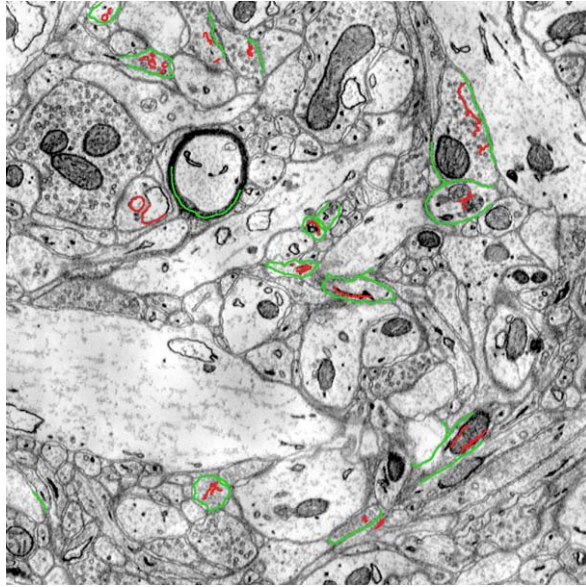
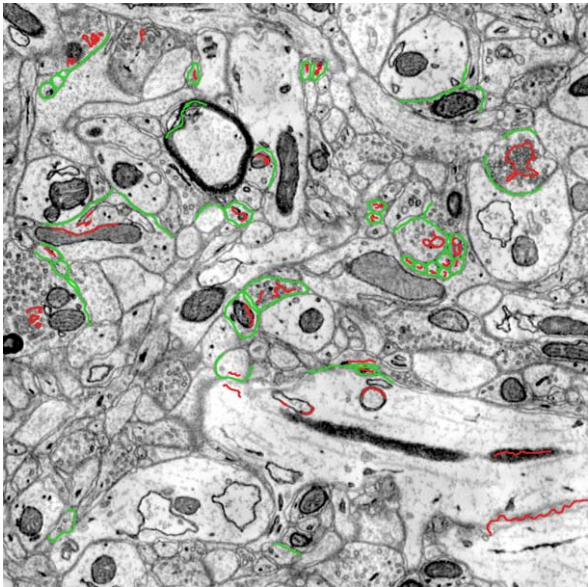
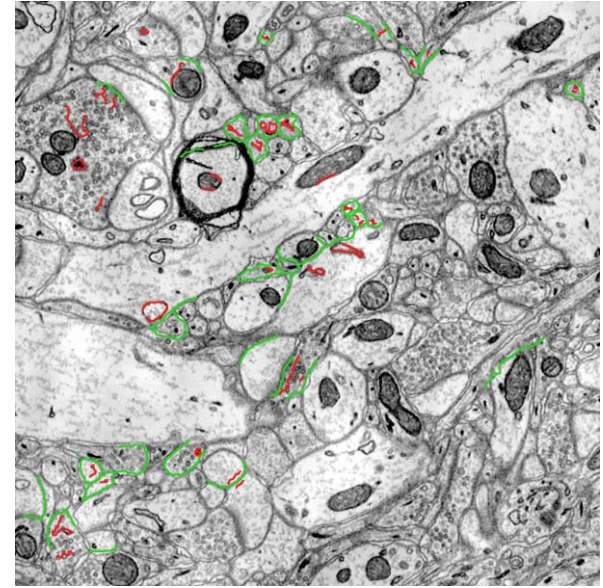
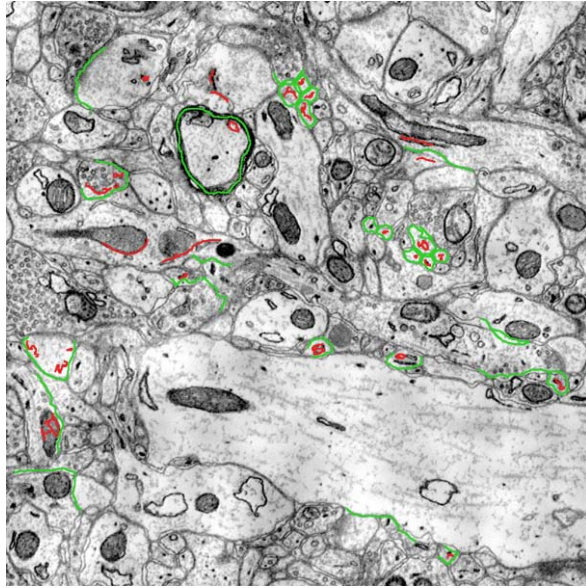
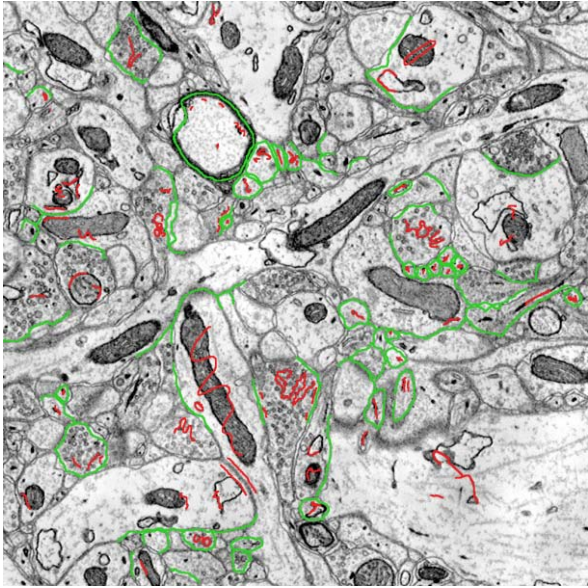


Manual annotation

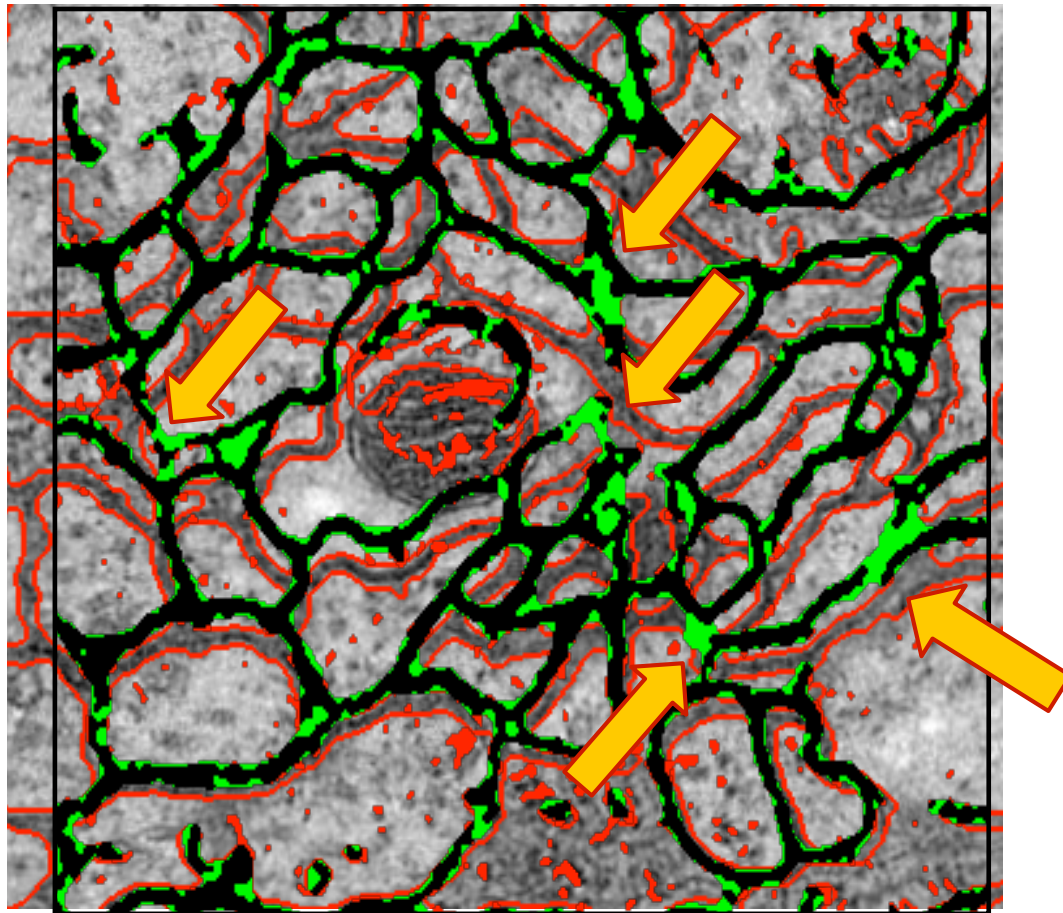


Visual feedback from
classification

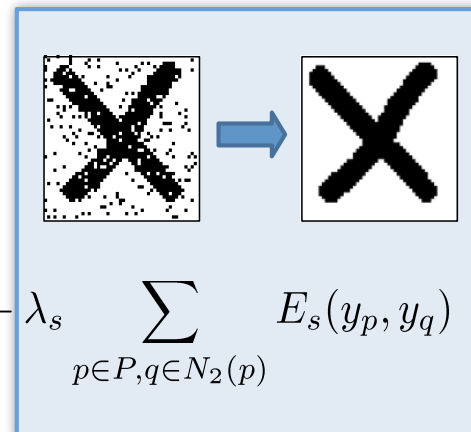
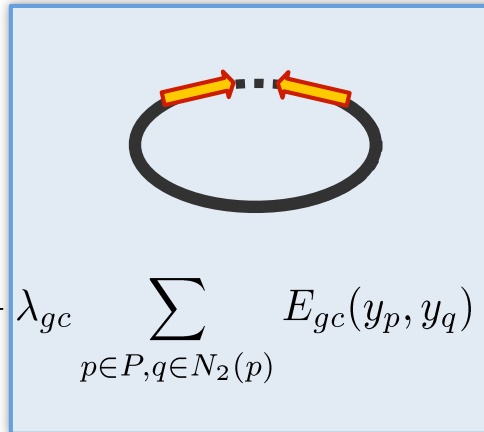
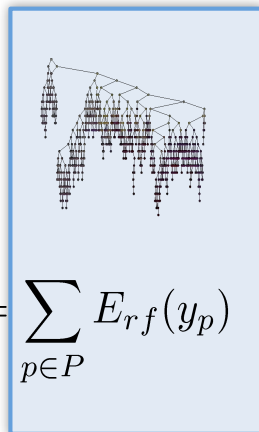
The whole training set! (<1% annotated)



Qualitative Evaluation

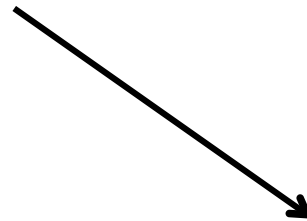
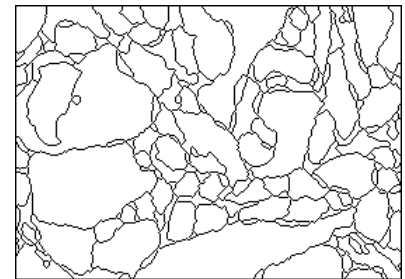
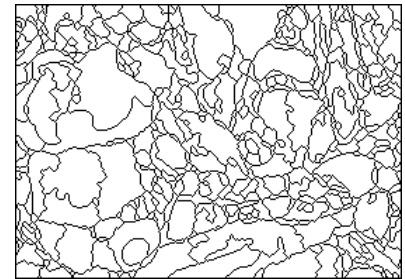
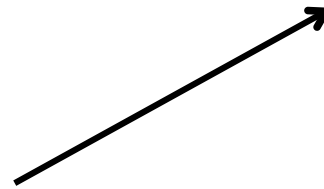
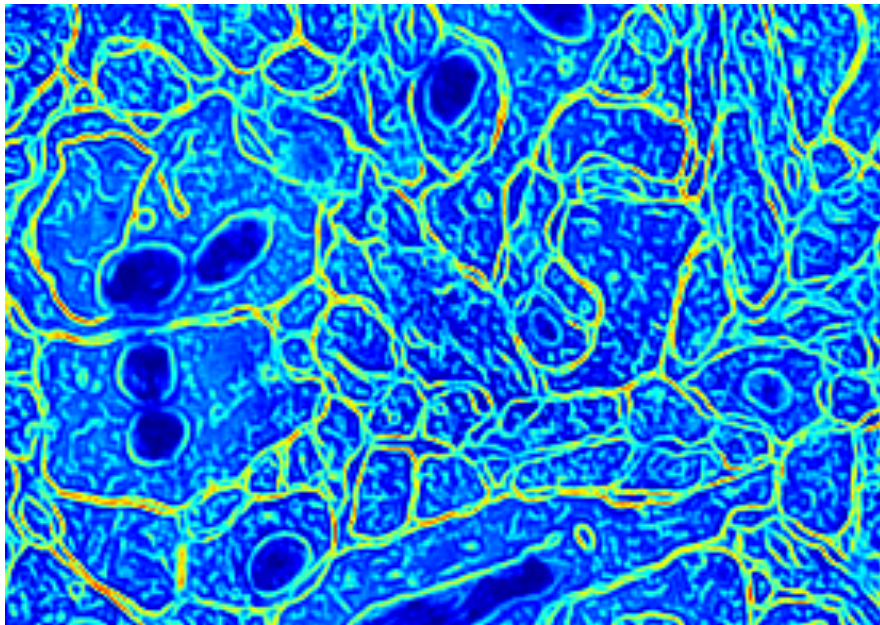


Gap Completion



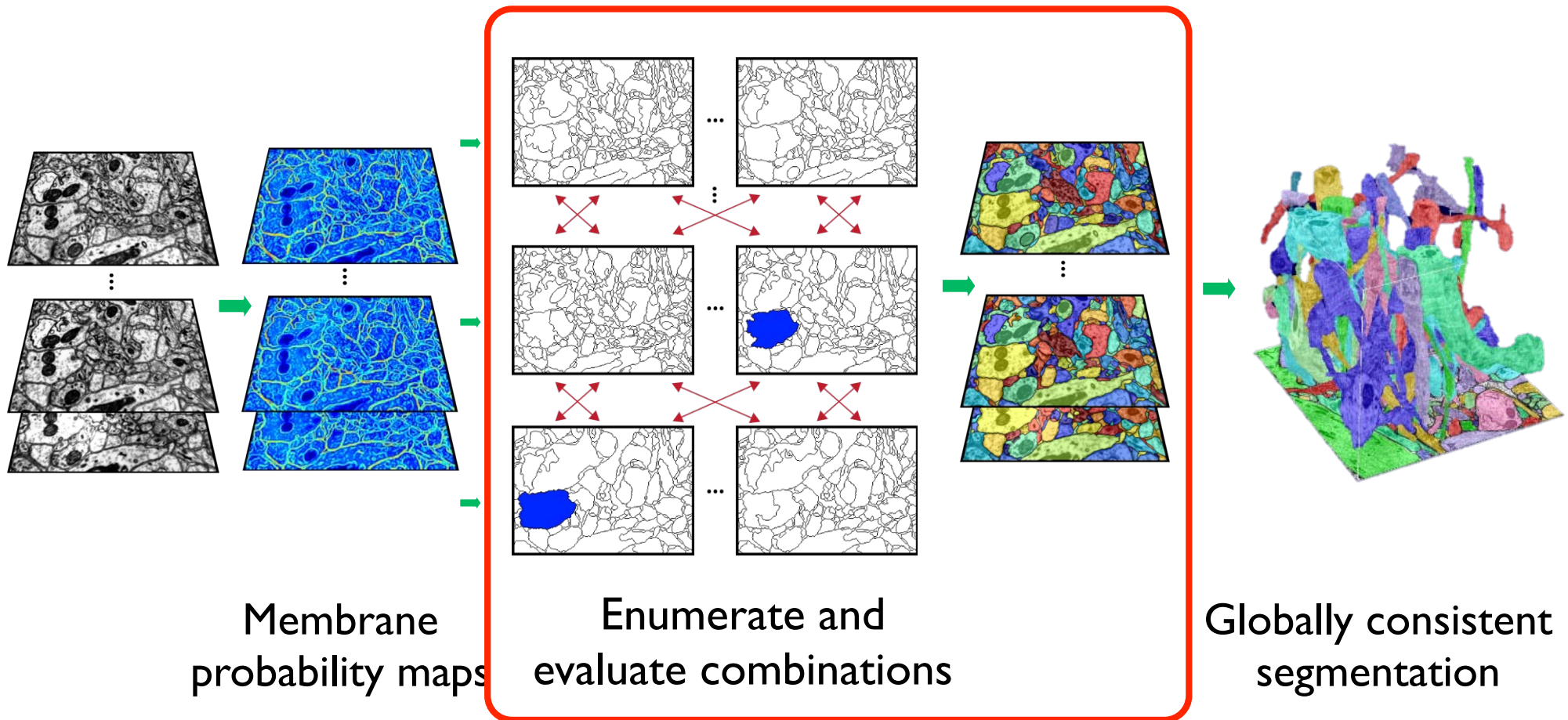
$$E(y) = \sum_{p \in P} E_{rf}(y_p) + \lambda_{gc} \sum_{p \in P, q \in N_2(p)} E_{gc}(y_p, y_q) + \lambda_s \sum_{p \in P, q \in N_2(p)} E_s(y_p, y_q)$$

Multiple Segmentations

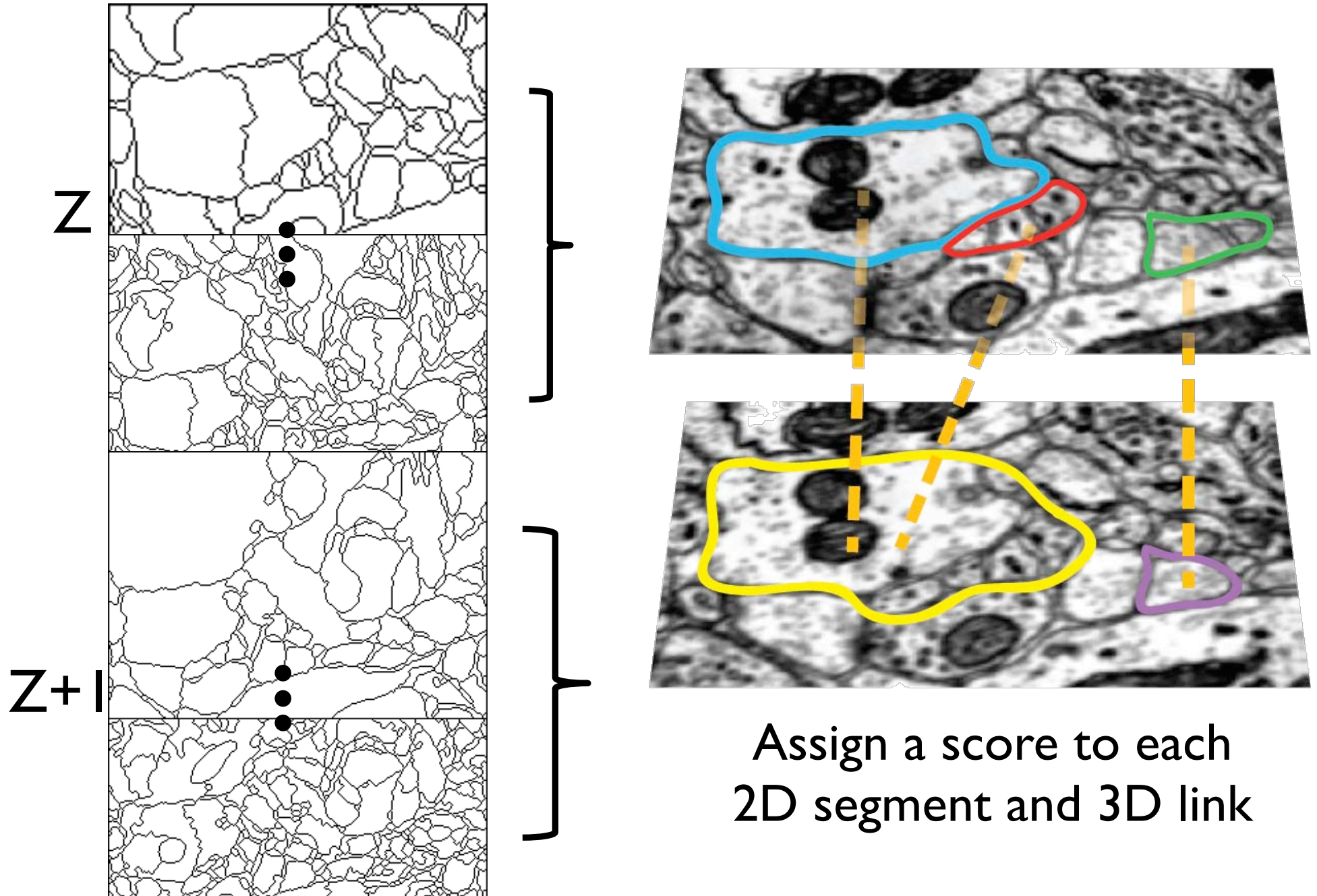


Closed contours

The Pipeline

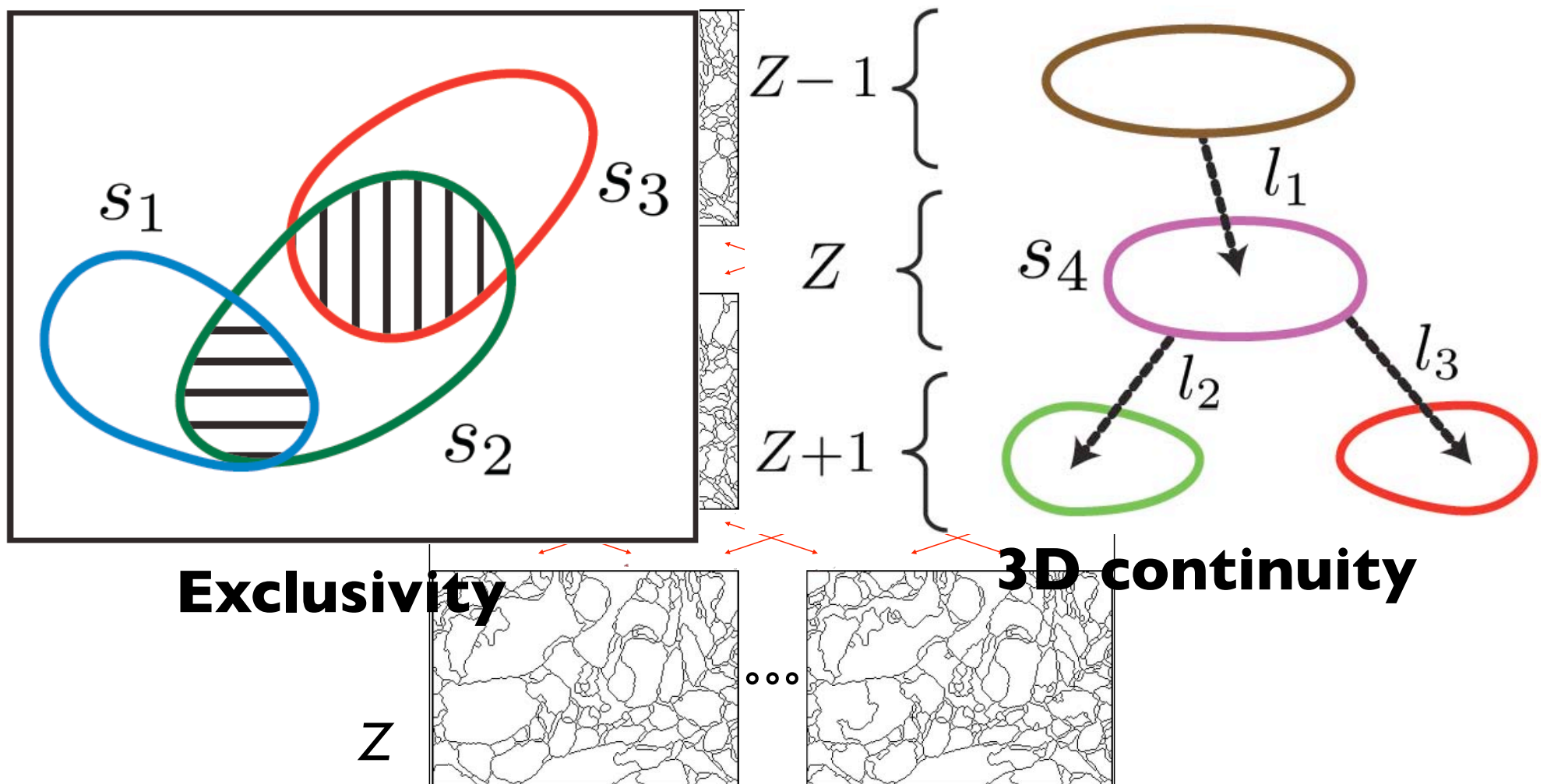


Enumerate 3D links



Modeling Fusion

Find the selection of segments and links that gives us the total highest score



Modeling Fusion

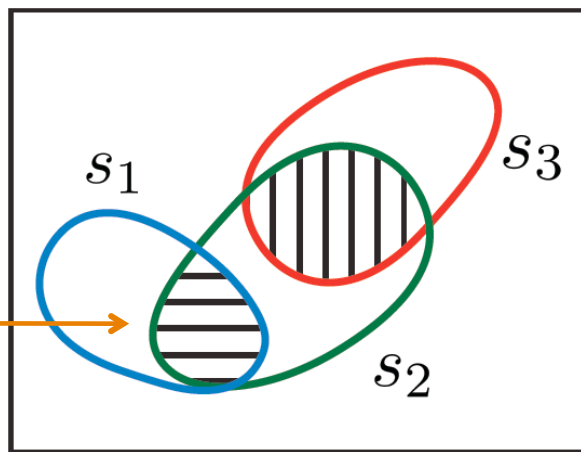
$$\arg \max_{\mathbf{s}, \mathbf{l}} \sum_{i=1}^S \theta_{s_i} s_i + \sum_{j=1}^L \theta_{l_j} l_j$$

Subject to:

Fitness of segment i

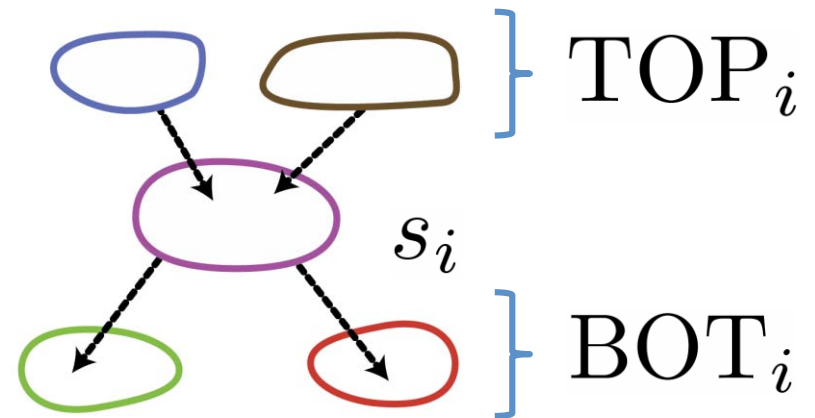
Fitness of link j

Exclusivity



$$\sum_{i \in O_k} s_i \leq 1$$

3D continuity

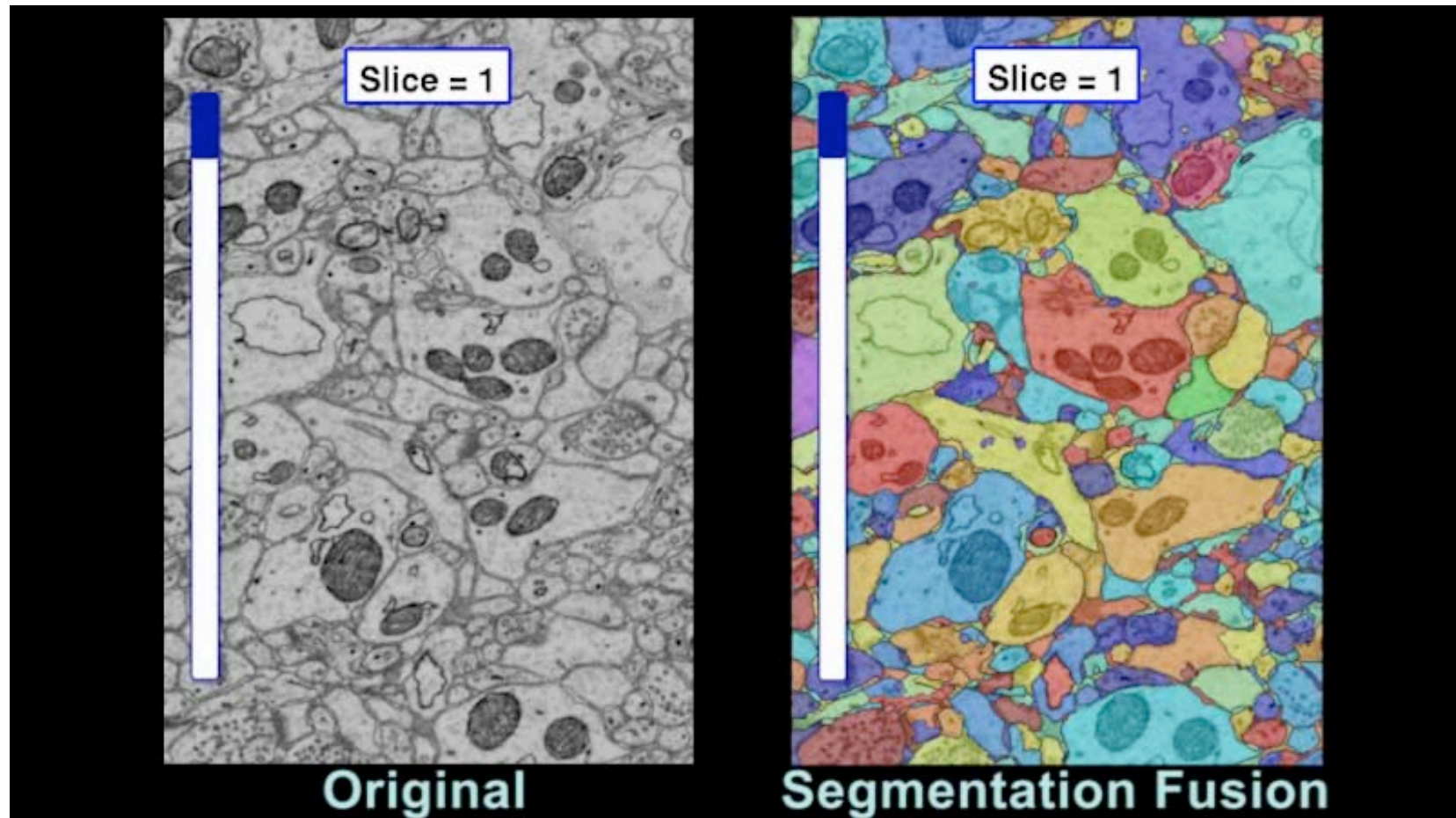


$$\sum_{j \in \text{TOP}_i} l_j \leq s_i, \quad \sum_{j \in \text{BOT}_i} l_j \leq s_i$$

Solving Fusion

- Constrained optimization using ILP solver
- Enumerating links is the slowest part
- 1000 x 1000 x 20 volume takes 30 mins
- Total time, including boundary probability maps, about 1 hour
- Match sub-blocks solving the assignment problem (max. weighted bipartite matching)

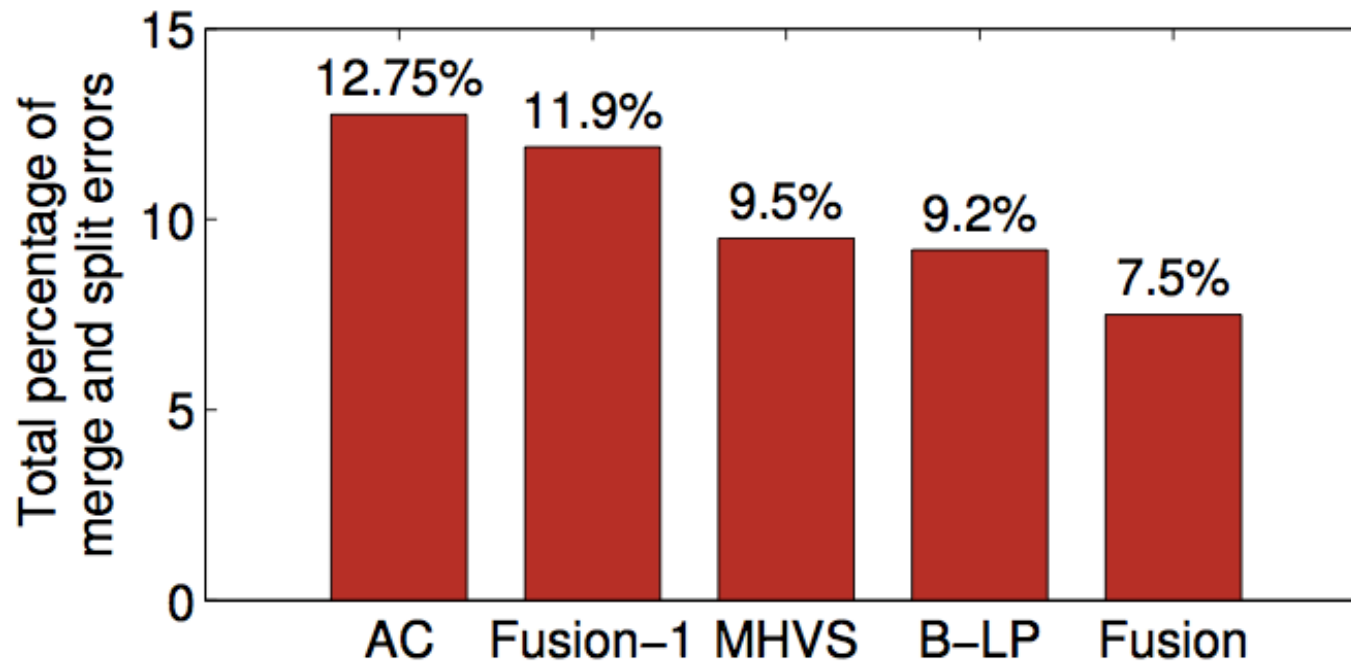
Results

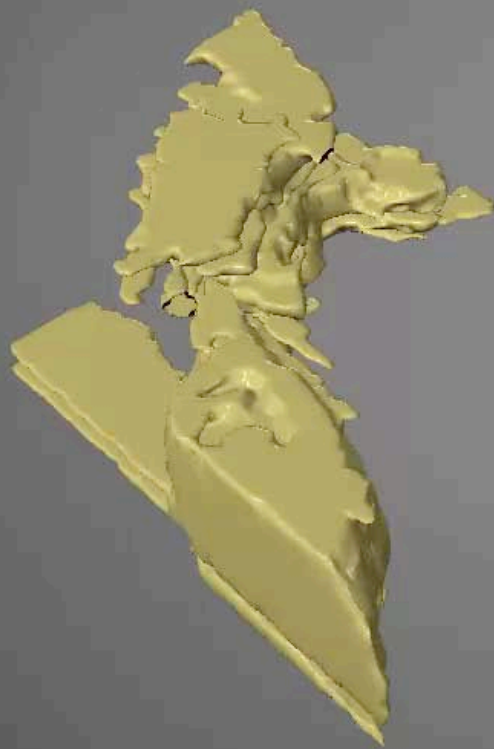


1,000 × 1,000 × 9

[Vazquez et al., ICCV 2011]

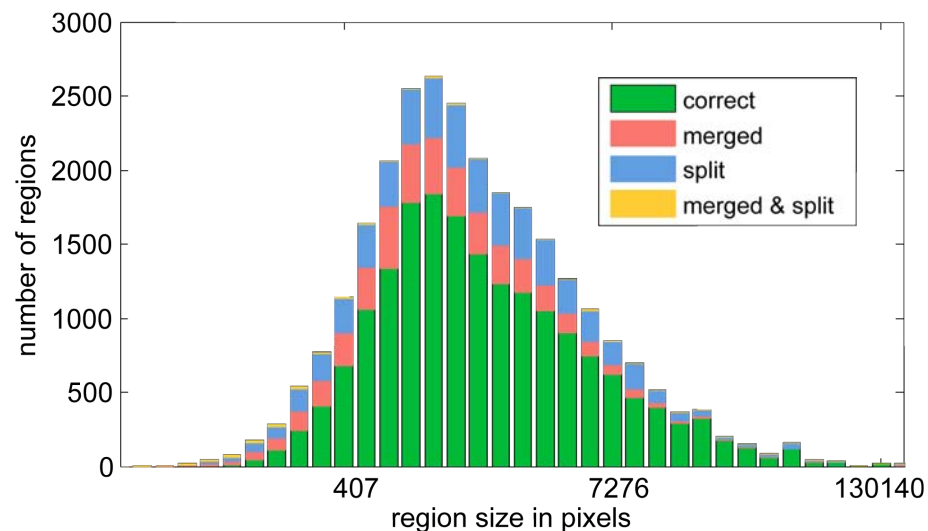
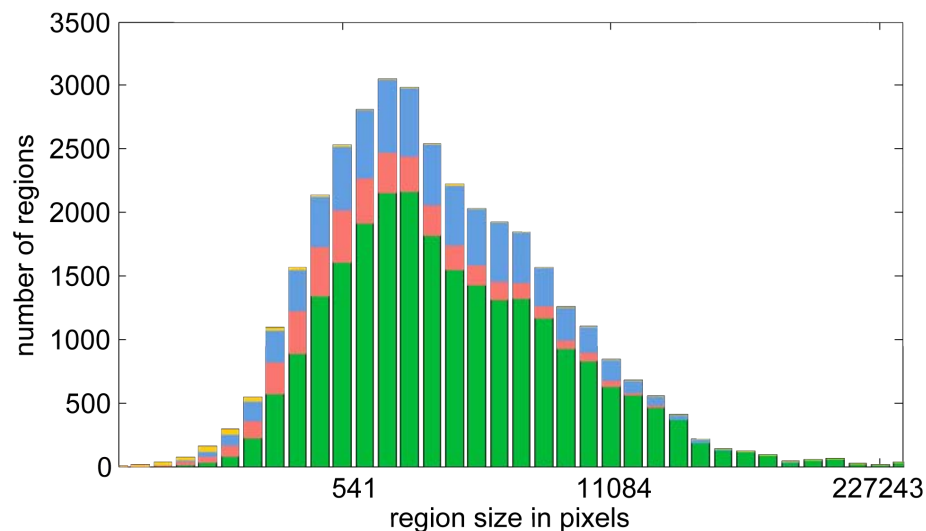
Error Comparison





1024 × 1024 × 100

Quantitative Evaluation



150 sections at 1024x1024 pixels
including 6 training images
correct segmentation covers:
69% ± 4 of all regions
82% ± 5 of all pixels

100 sections at 1024x1024 pixels
completely test performance
correct segmentation covers:
68% ± 4 of all regions
71% ± 7 of all pixels

Mojo

File Edit

Adjust Segmentation
 Merge Segmentation
 Split Segmentation

Show Segmentation
 Constrain Segmentation Merging To Current Slice
 Constrain Segmentation Merging To Connected Component

Proof Reading



Neural Processes

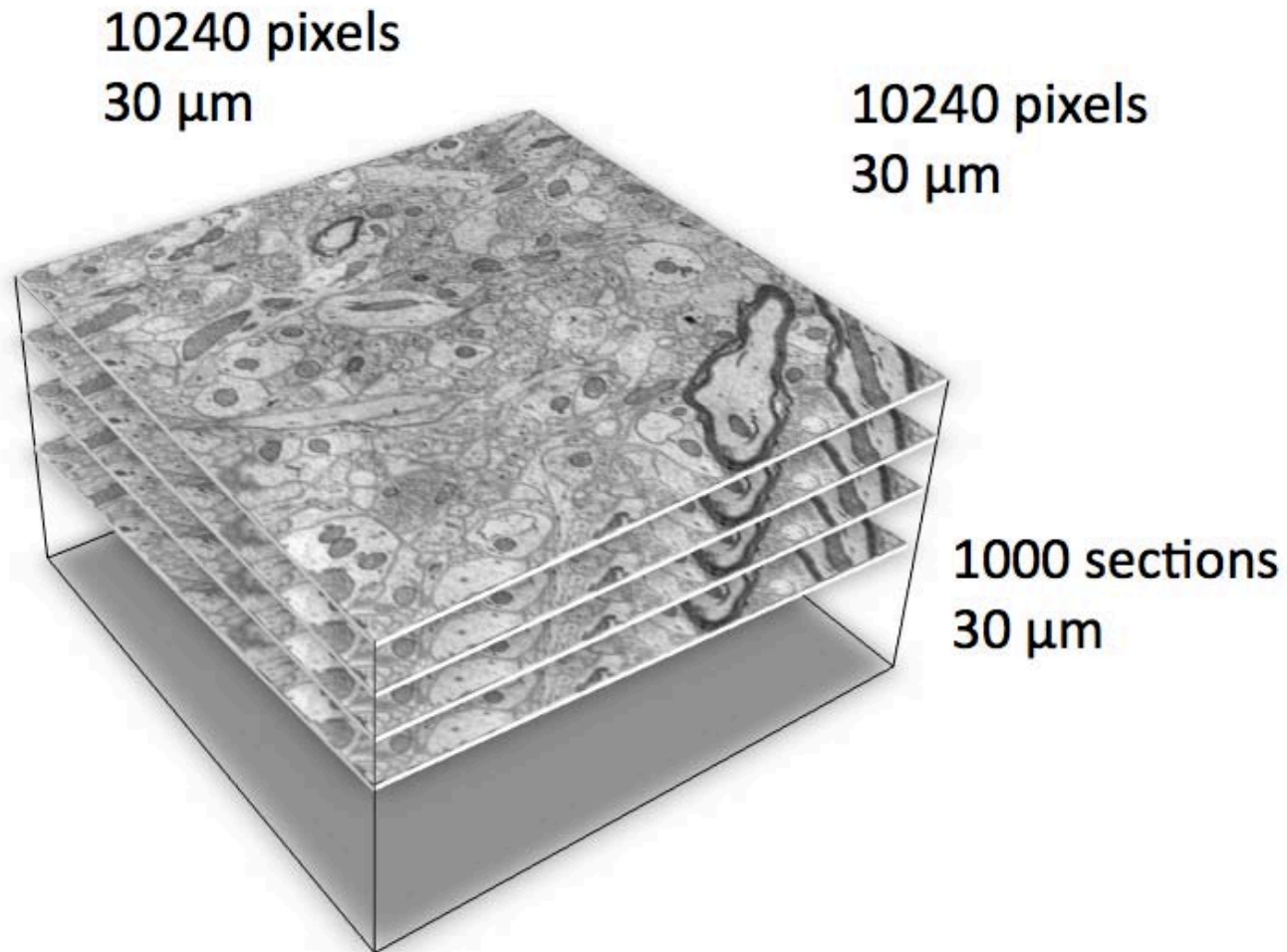
Autogenerated Neural Process (ID 3677)	[64,26,239]
Autogenerated Neural Process (ID 3485)	[96,136,25]
Autogenerated Neural Process (ID 3551)	[156,235,55]
Autogenerated Neural Process (ID 3413)	[72,187,167]
Autogenerated Neural Process (ID 3415)	[175,206,227]
Autogenerated Neural Process (ID 3429)	[212,27,45]
Autogenerated Neural Process (ID 3567)	[128,189,164]
Autogenerated Neural Process (ID 3553)	[98,227,63]
Autogenerated Neural Process (ID 3604)	[190,221,9]
Autogenerated Neural Process (ID 3702)	[208,179,142]
Autogenerated Neural Process (ID 3620)	[65,138,209]
Autogenerated Neural Process (ID 3561)	[245,148,111]
Autogenerated Neural Process (ID 3461)	[24,173,122]
Autogenerated Neural Process (ID 3446)	[188,99,192]
Autogenerated Neural Process (ID 3464)	[197,202,199]
Autogenerated Neural Process (ID 3471)	[108,114,1]
Autogenerated Neural Process (ID 3483)	[97,224,241]
Autogenerated Neural Process (ID 3474)	[21,151,76]
Autogenerated Neural Process (ID 3572)	[254,84,193]
Autogenerated Neural Process (ID 3456)	[57,114,136]
Autogenerated Neural Process (ID 3436)	[78,166,13]
Autogenerated Neural Process (ID 3426)	[23,250,86]
Autogenerated Neural Process (ID 3519)	[43,160,115]
Autogenerated Neural Process (ID 3543)	[102,250,216]
Autogenerated Neural Process (ID 3555)	[226,104,122]
Autogenerated Neural Process (ID 3650)	[129,109,142]
Autogenerated Neural Process (ID 3390)	[60,182,83]
Autogenerated Neural Process (ID 3661)	[58,160,192]
Autogenerated Neural Process (ID 3656)	[148,76,141]
Autogenerated Neural Process (ID 3377)	[2,86,21]
Autogenerated Neural Process (ID 3571)	[102,88,253]
Autogenerated Neural Process (ID 3660)	[235,254,182]
Autogenerated Neural Process (ID 3518)	[12,22,181]
Autogenerated Neural Process (ID 3458)	[123,247,210]
Autogenerated Neural Process (ID 3378)	[95,71,204]
Autogenerated Neural Process (ID 3419)	[225,22,214]
Autogenerated Neural Process (ID 3559)	[108,230,73]
Autogenerated Neural Process (ID 3655)	[117,121,19]
Autogenerated Neural Process (ID 3516)	[123,29,152]
Autogenerated Neural Process (ID 3616)	[220,74,46]
Autogenerated Neural Process (ID 3443)	[31,61,70]
Autogenerated Neural Process (ID 3540)	[149,151,37]
Autogenerated Neural Process (ID 3624)	[211,78,221]
Autogenerated Neural Process (ID 3686)	[173,80,140]
Autogenerated Neural Process (ID 3520)	[66,111,53]
Autogenerated Neural Process (ID 3614)	[166,219,78]
Autogenerated Neural Process (ID 3683)	[97,91,21]

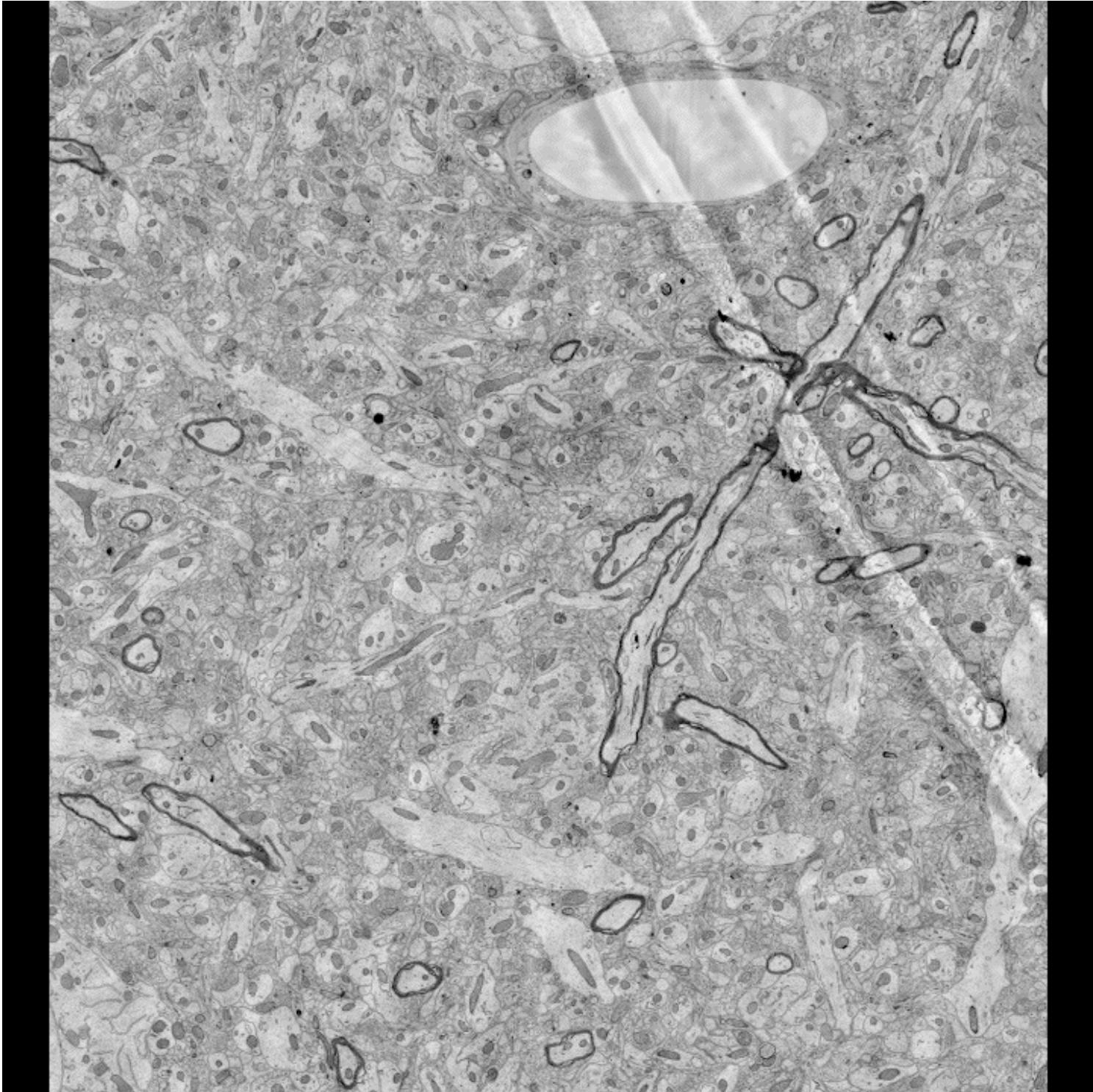
Neural Process Name:

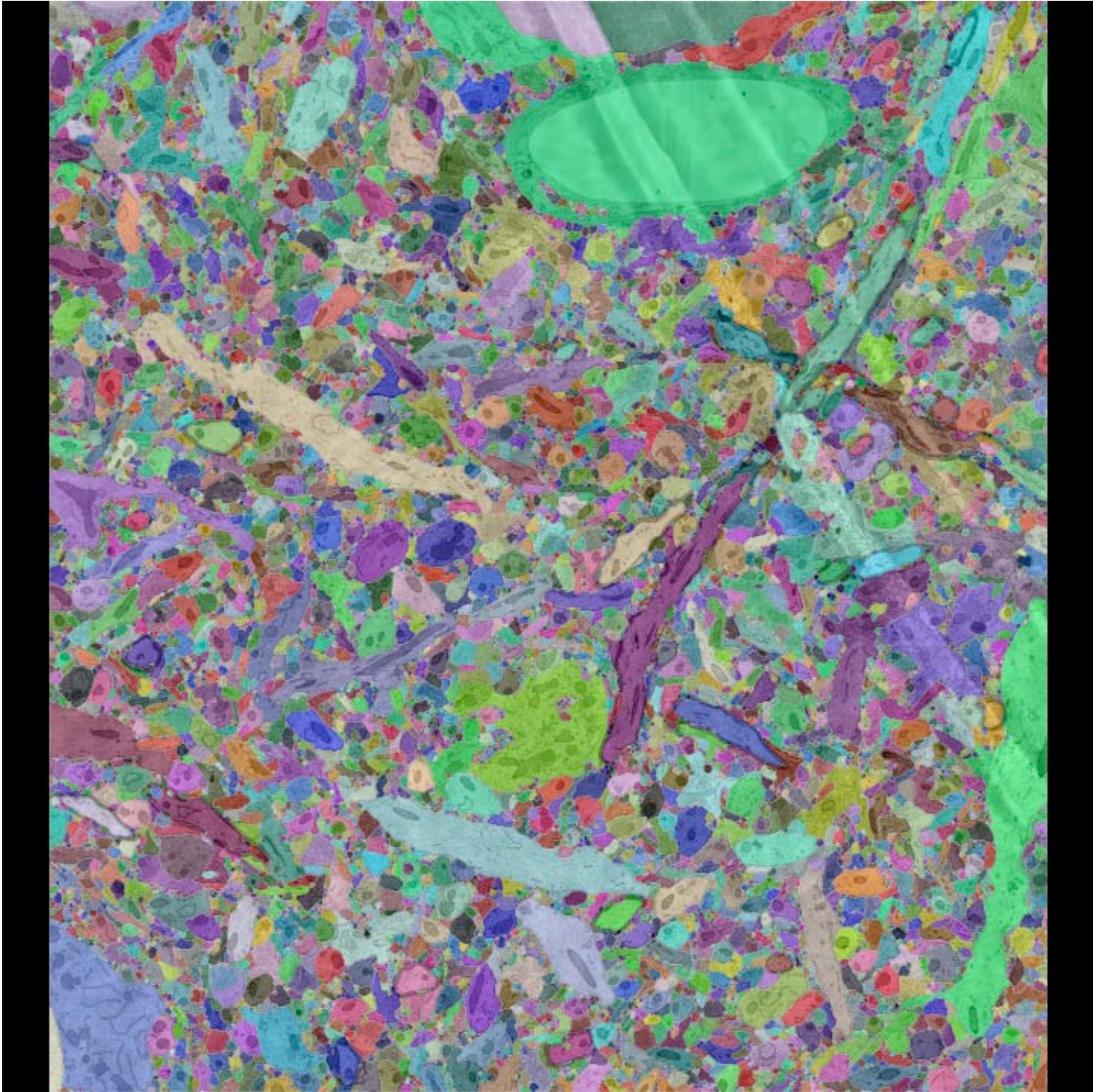
Left mouse button selects process to split.

Jeff's Challenge

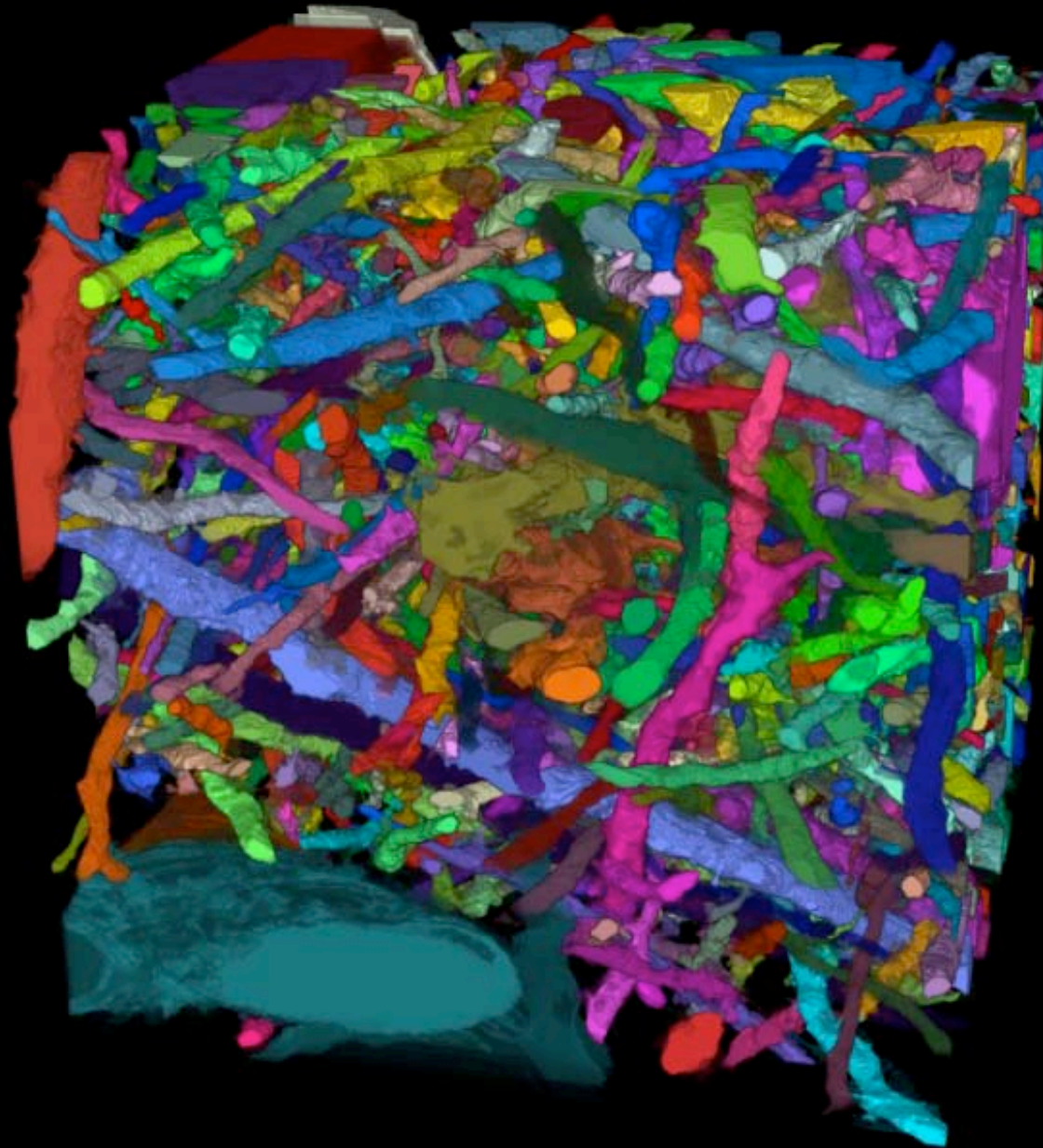
Automatic 3D reconstruction for largest EM stack **ever**

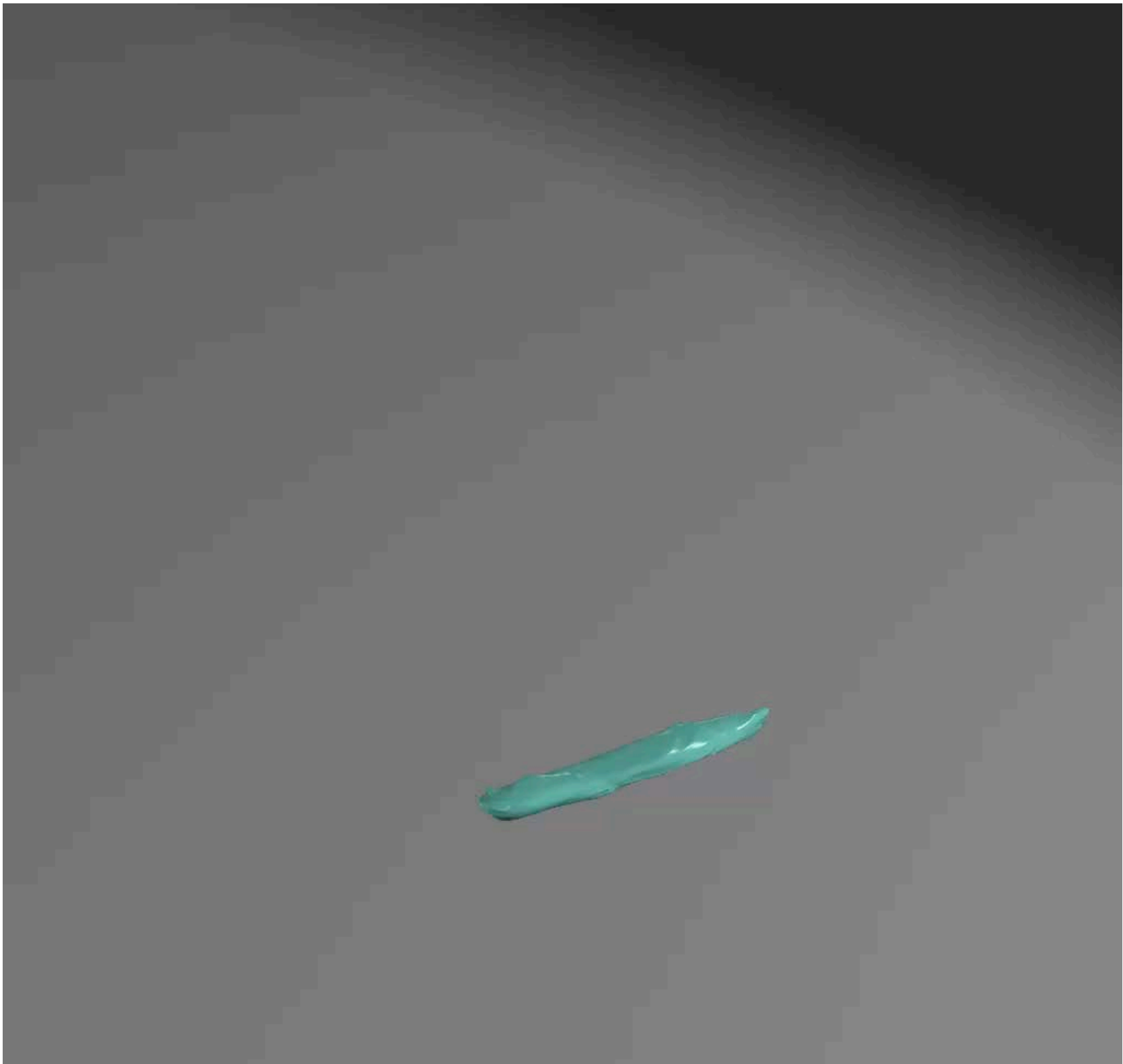


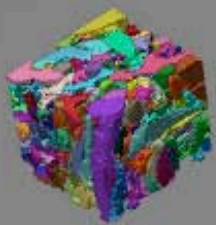




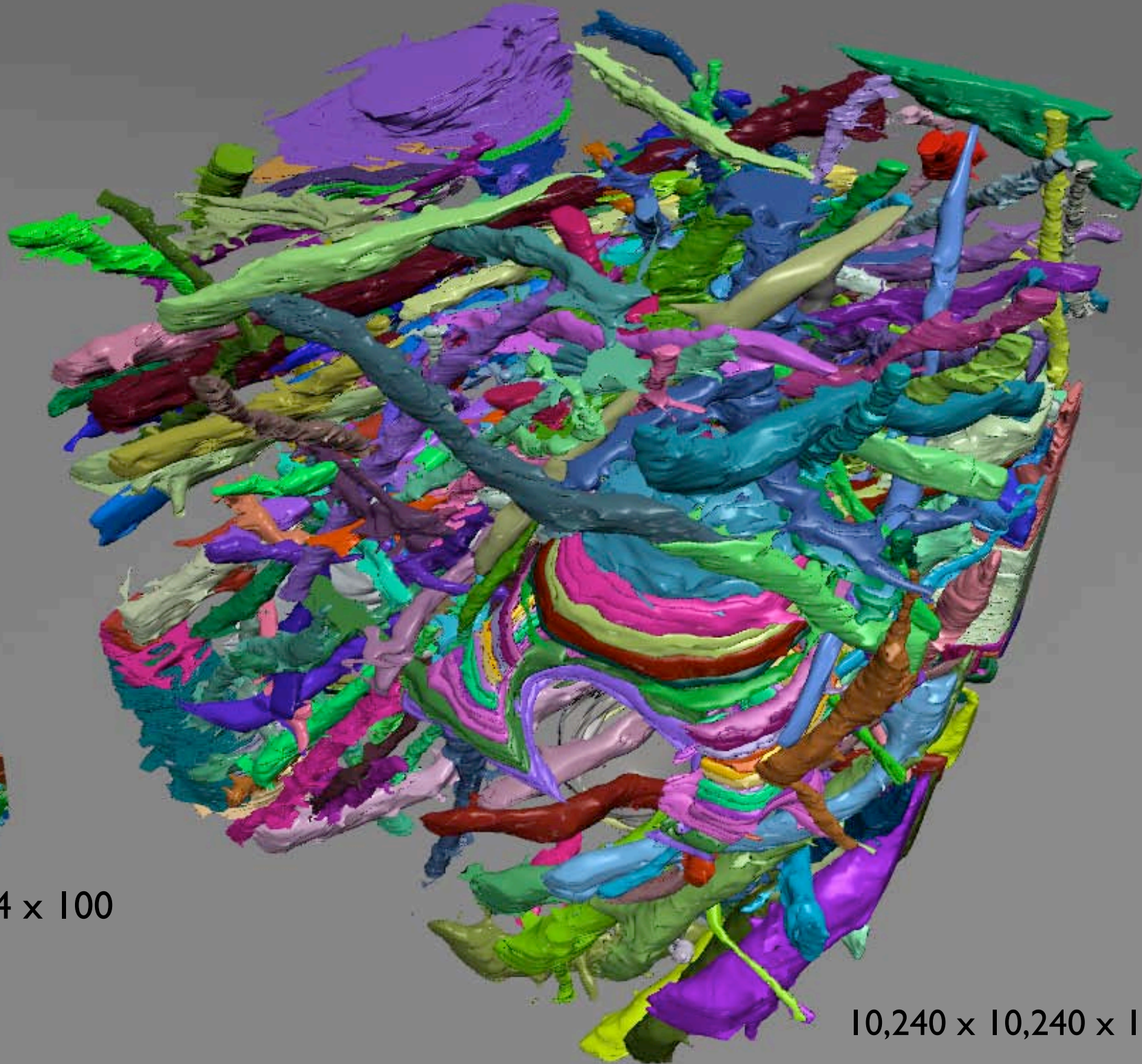
All objects containing at least 100 sections







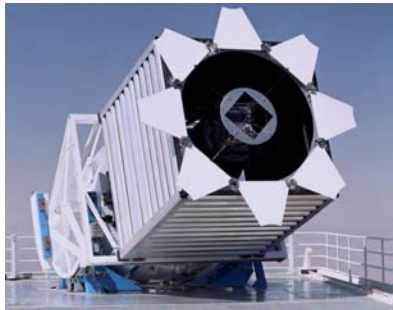
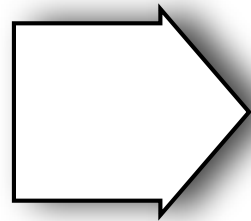
1,024 × 1,024 × 100



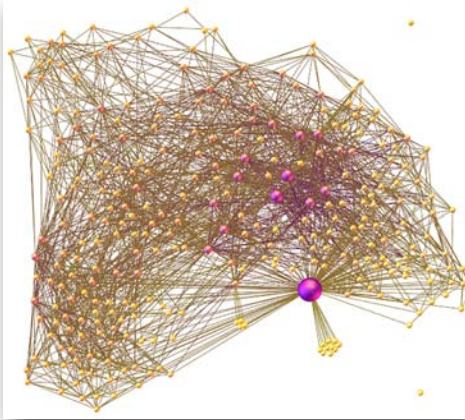
10,240 × 10,240 × 1,000

Future Work

Distributed Image Analysis & Visualization



CBS



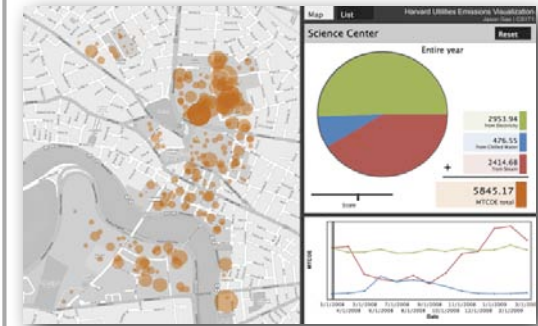
**Network
Visualization**

MGH

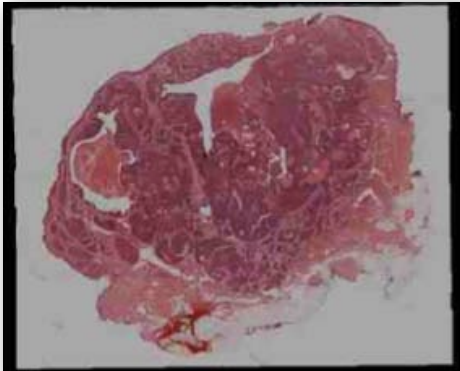


**Radiation
Oncology**

GSD



**Building Energy
Usage**



**Digital
Pathology**

Beth Israel



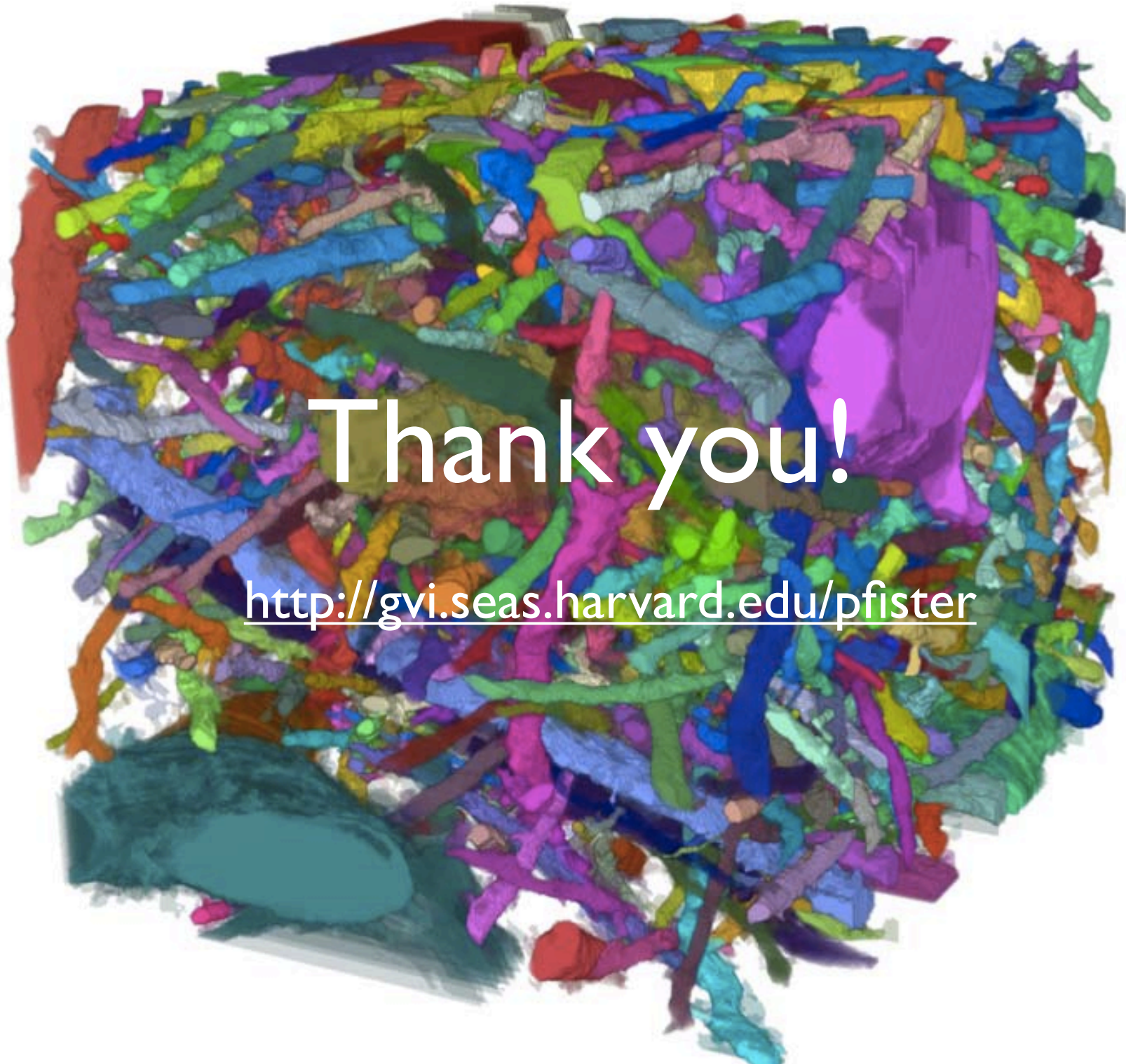
**Object
Replication**

Disney Research



**Video
Browsing**

IQSS



Thank you!

<http://gvi.seas.harvard.edu/pfister>