

ConnectomeExplorer

Query-Guided Visual Analysis of Large Volumetric Neuroscience Data

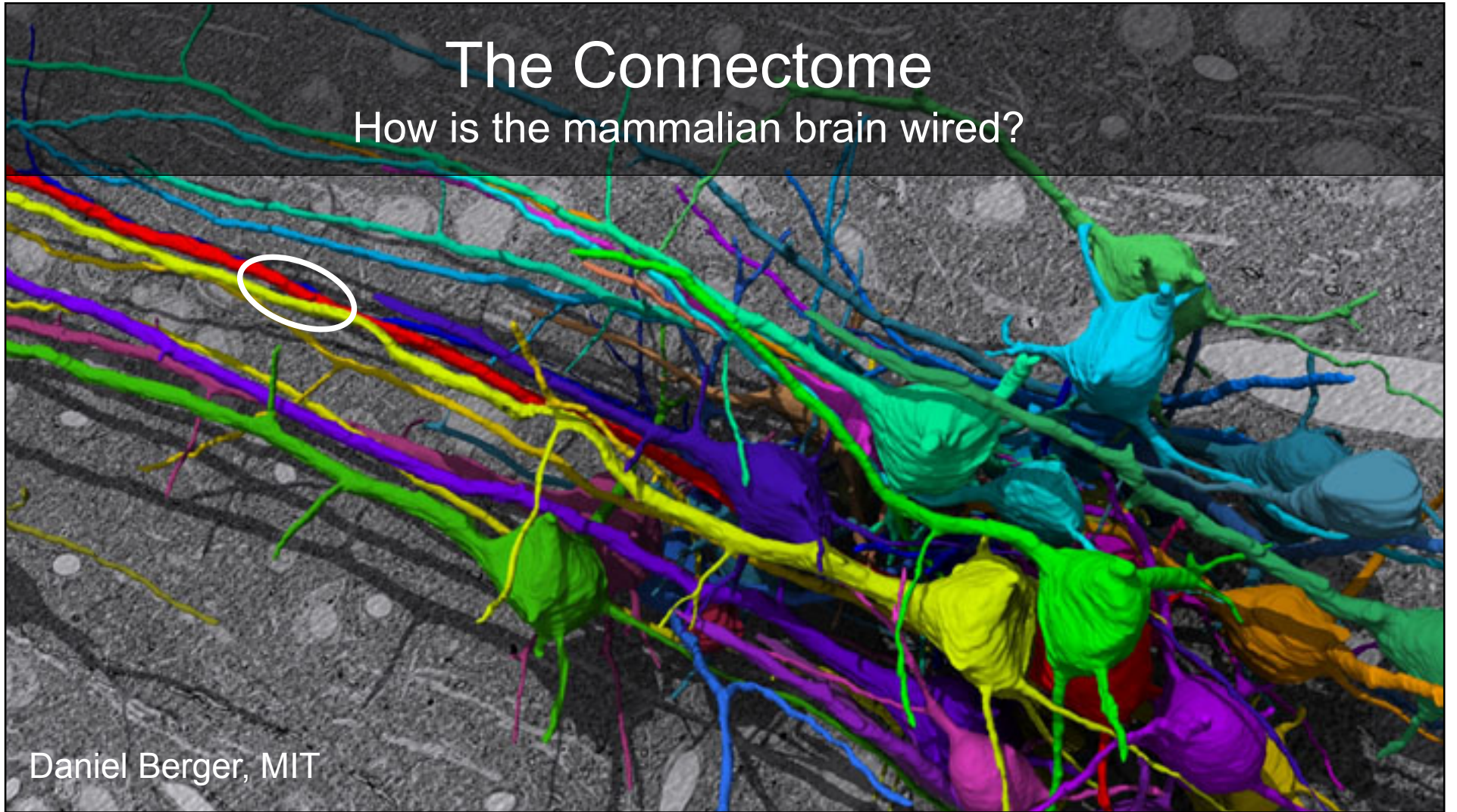


HARVARD
School of Engineering
and Applied Sciences

Johanna Beyer, Ali Al-Awami, Bobby Kasthuri,
Jeff W. Lichtman, Hanspeter Pfister, Markus Hadwiger

The Connectome

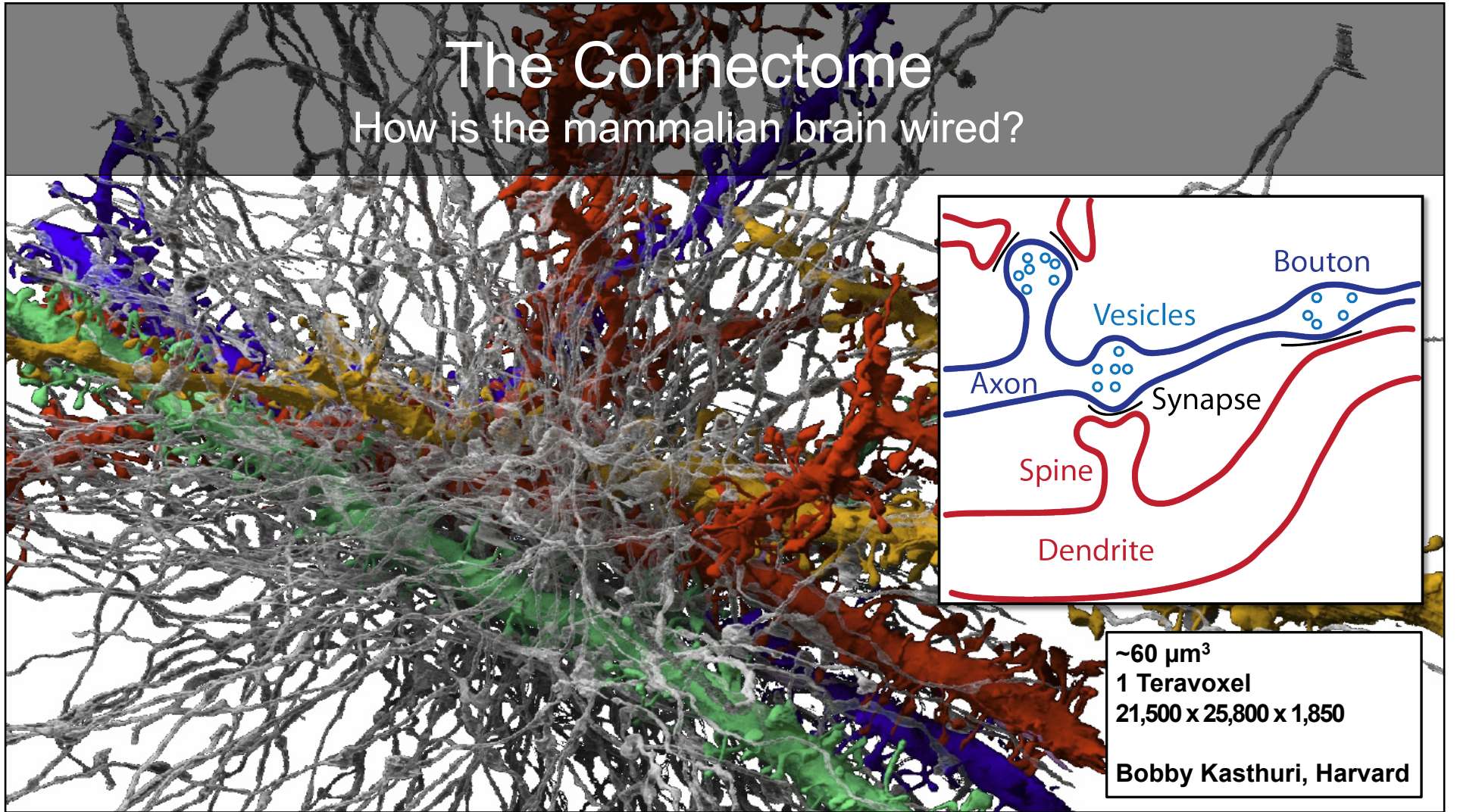
How is the mammalian brain wired?



Daniel Berger, MIT

The Connectome

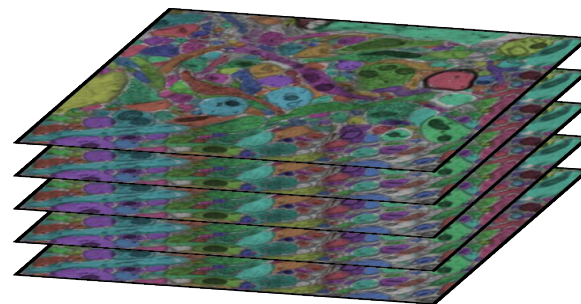
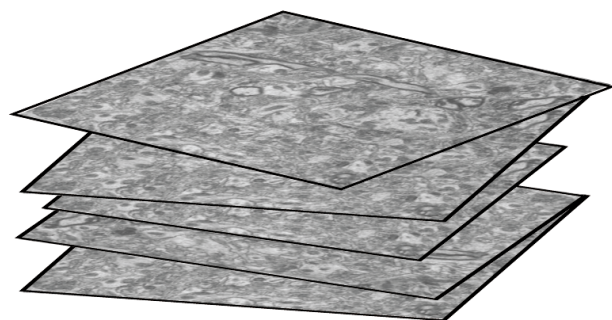
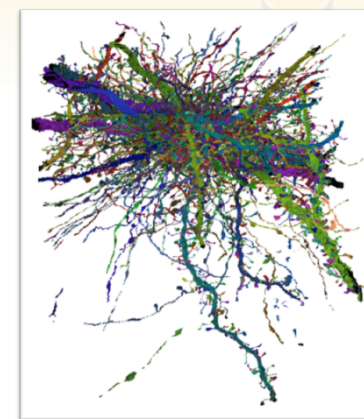
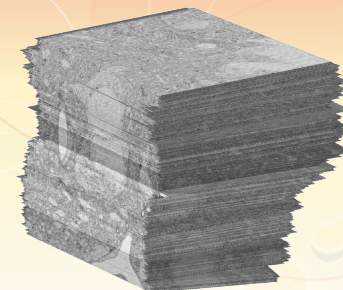
How is the mammalian brain wired?



~60 μm^3
1 Teravoxel
21,500 x 25,800 x 1,850
Bobby Kasthuri, Harvard

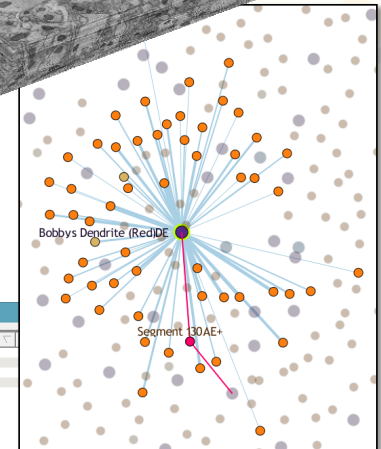
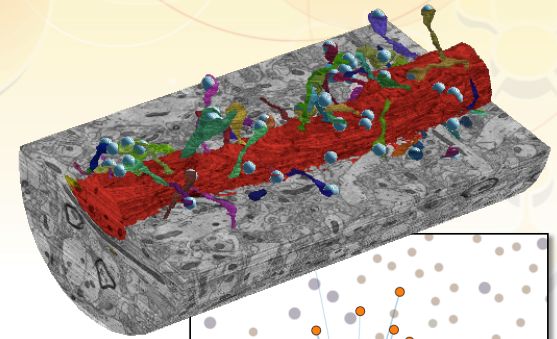
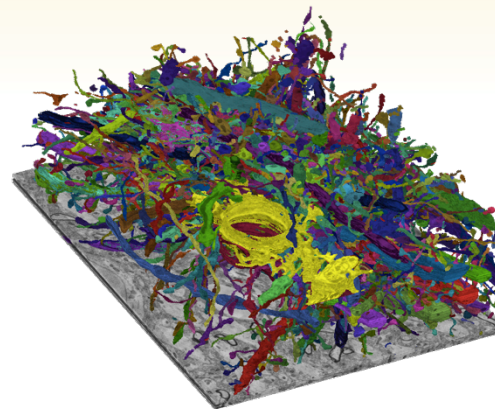
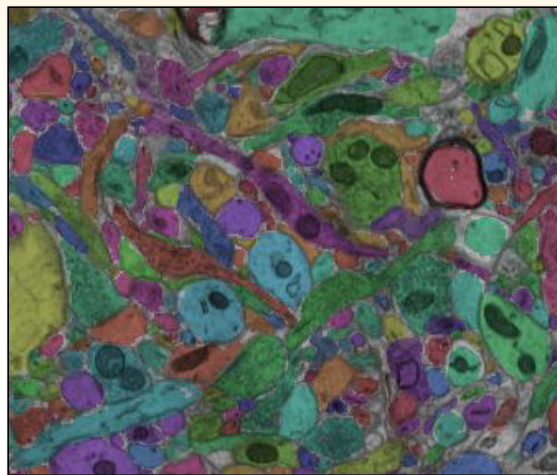
Motivation

- Huge amount of data scanned
- Took years to align and segment
- How to analyze it?

A screenshot of a data table with multiple columns and rows, showing numerical values and some text labels, likely representing the analysis results of the scanned data.

...
...
...
...
...

Towards Visual Connectomics



1

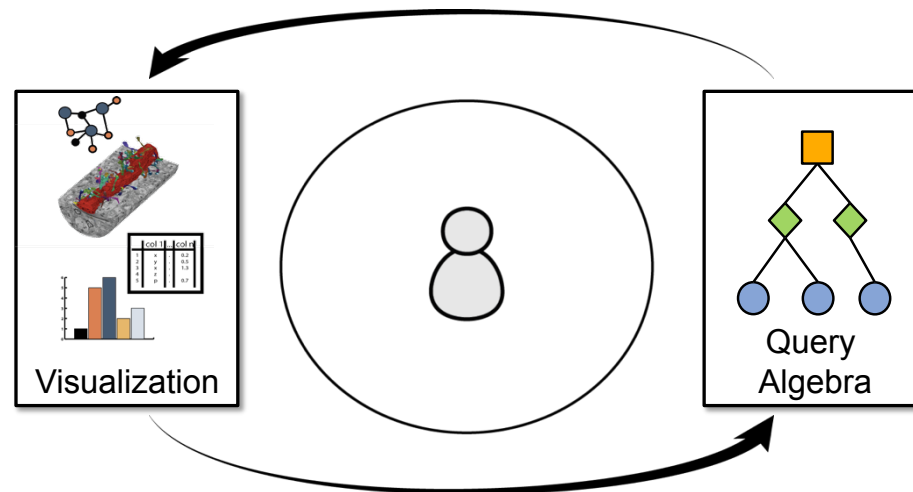
is Connected! (All Objects, b)

- [-] Synapse_Dendrite
- [+] Brodmann_Dendrite
- [+] Axon_Dendrite
- [+] Dendrite

	Axon Name	Axon Voxels	Dendrite Name
29	Segment 134_AE+	1392385	Bobbys Dendrite (Red)_D
30	Segment 133_AE+	1620026	Bobbys Dendrite (Red)_D
31	Segment 132_AE+	1777727	Bobbys Dendrite (Red)_D
32	Segment 130_AE+	4361000	Bobbys Dendrite (Red)_D
33	Segment 12_AE+	2245577	Bobbys Dendrite (Red)_D

Contributions

- Fully dynamic, domain-specific queries
- Simple set-based query algebra
- Iterative, interactive query specification
- Query-driven petascale visualization pipeline (extends [SciVis 2012])



Connectome Options

Options Segmentation Synapses Axons Dendrites ROI 3D Absl

Load Save New Merge Delete

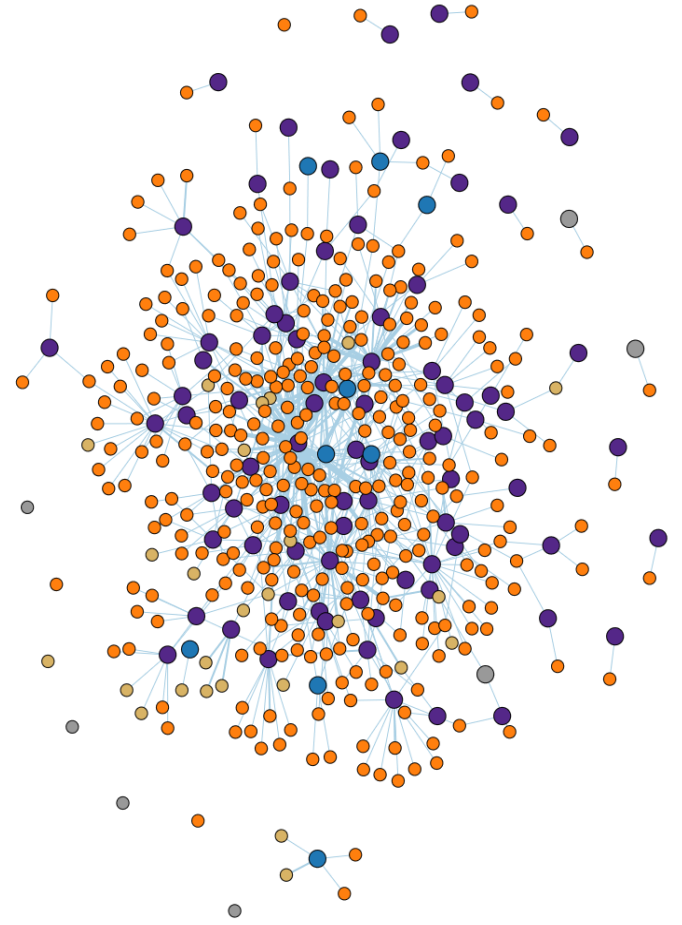
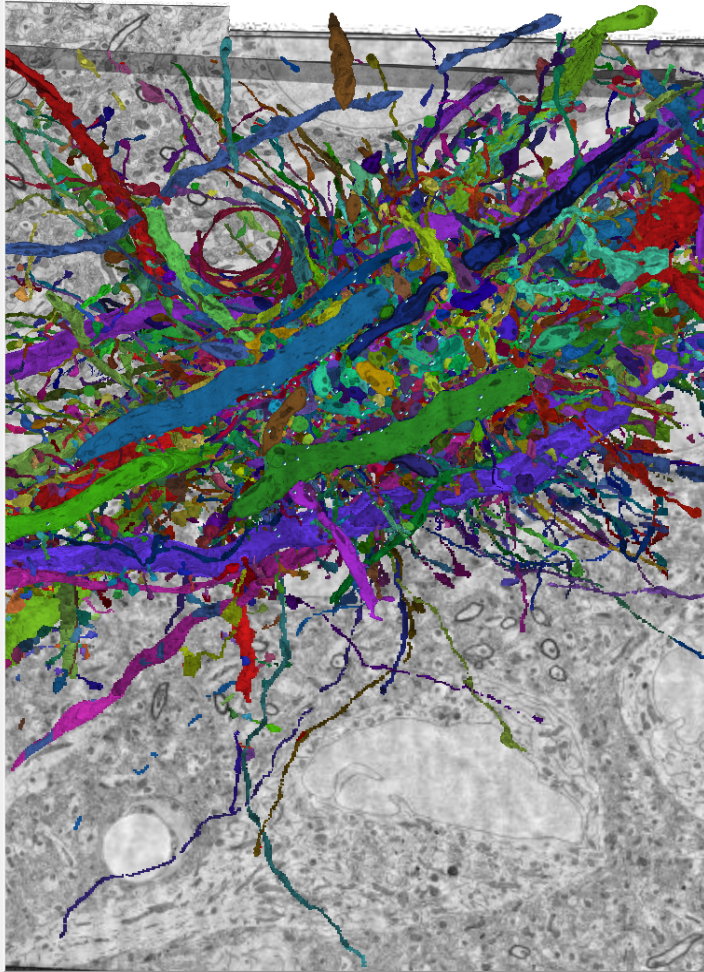
Segment Neuron

- F:\Jose01.vss_NT
- DONE_NT
- Myelinated Axons_NT
- AxonAI+_NT
- AxonA7_NT
- AxonAN_NT
- AxonAE+_NT
- AxonAE+_NT
- DE_NT
 - Segment 4203_DE
 - Segment 4138_DE
 - Emily Bernier 8_DE
 - Solis 38_DE
 - Solis 35_DE
 - Bird SP3 35_DE
 - Dendrites 913_DE
 - Segment 8492_DE
 - Rosenburgd3 2_DE??
 - Dendrites 921_DE
 - Segment 8496_DE
 - MS s4_DE
 - Segment 8335_DE
 - Segment 8431_DE
 - Dendrites 935_DE
 - Segment 8330_DE
 - Dendrites 941_DE
 - Bobbys Dendrite (Red)_DE
 - Dendrites 943_DE
 - Ran 3rd 1_DE
 - Dendrites 900_DE
 - Dendrite 6 Sparse_DE
 - CB Segment 684_DE
 - Segment 8430_DE
 - RRz105_DE
 - Dendrite 7 deep AP02_DE
 - Dendrites 924_DE
 - Upper Dendrite (Green)_DE
 - Dendrites 927_DE
 - Dendrites 911_DE
 - Dendrite 8 AP03_DE
 - Dendrite 5_DE
 - Segment 8476_DE
 - Dendrite 4 Basal_DE
 - Rosenburgd3 3_DE
 - Dendrites 910_DE
 - Lower Dendrite (Blue)_DE
 - Segment 8454_DE
 - Dendrites 987_DE
 - Dendrites 952_DE
 - StuS1106048_DE
 - Dendrites 923_DE
 - Emily Bernier 9_DE??
 - CB segment 176_DE
 - Bird SP 38_DE
 - Dendrites 915_DE
 - Segment 8440_DE

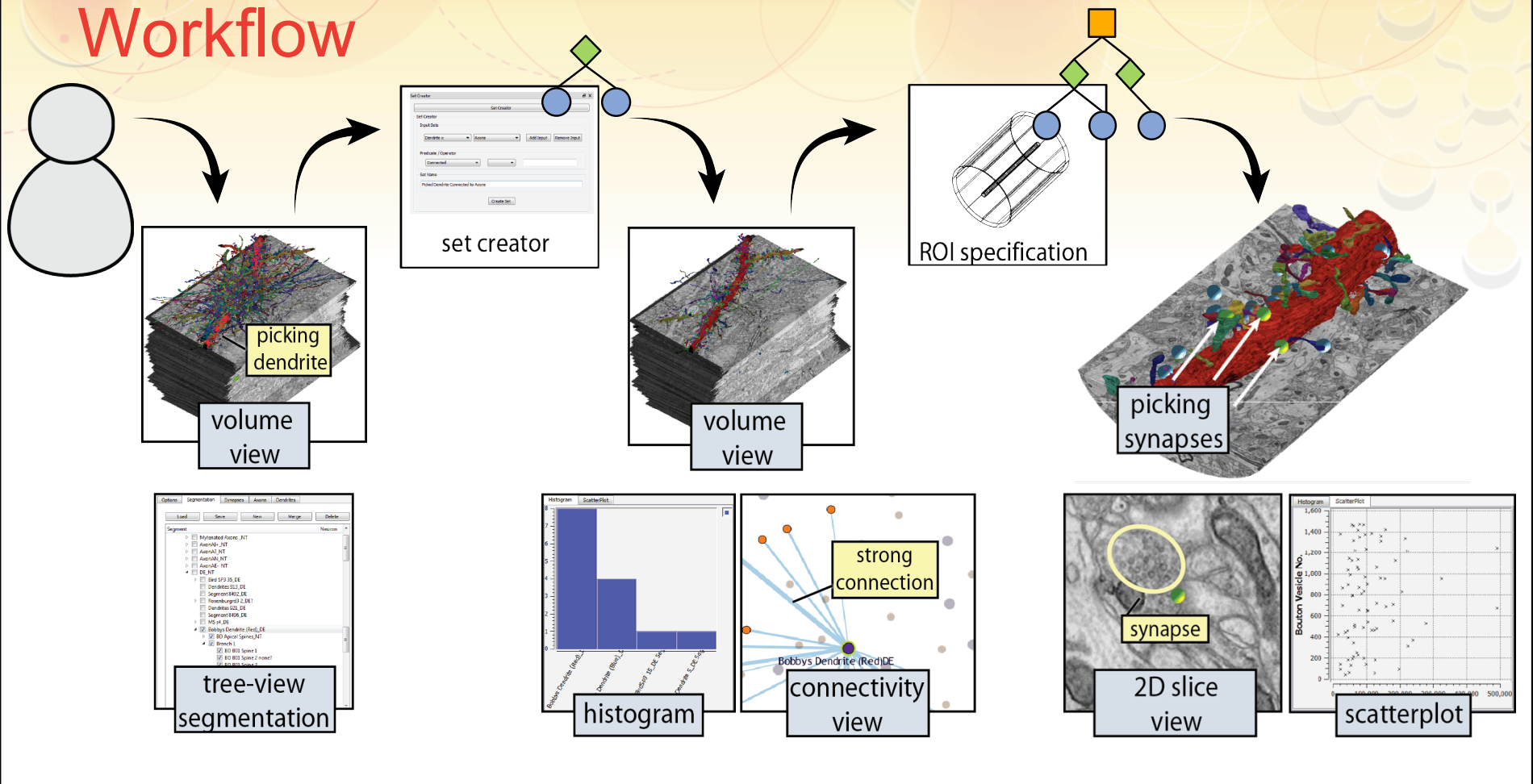
Find

Flat View Mapping Filter Non-Selected Check All Uncheck All

Main Options Connectome Options

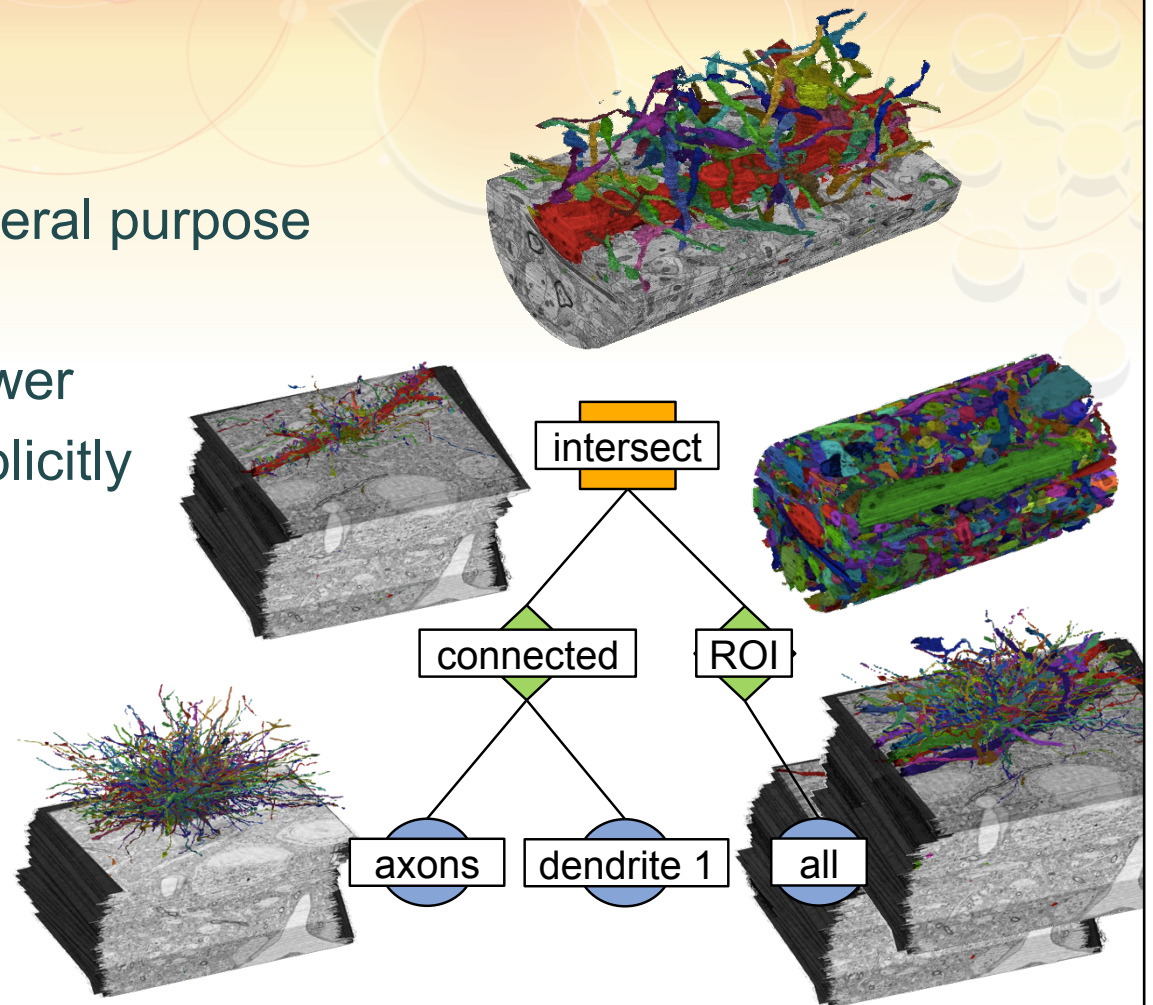


Workflow



Dynamic Queries

- Knowledge-based vs. general purpose
 - Simple to use
 - Enough expressive power
 - Exploit connectivity implicitly
- Components
 - Query algebra
 - Visual Set Creator

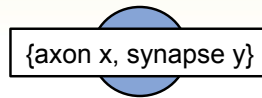


Set Algebra

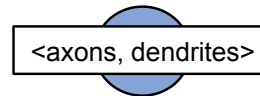
- Result of every query is a set

- Set can contain:

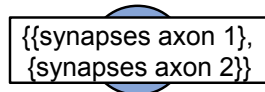
- Objects



- Ordered tuples of objects

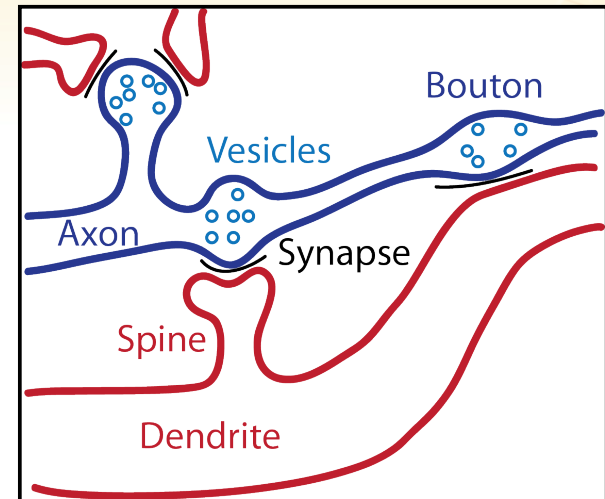


- Sets



- Object attributes

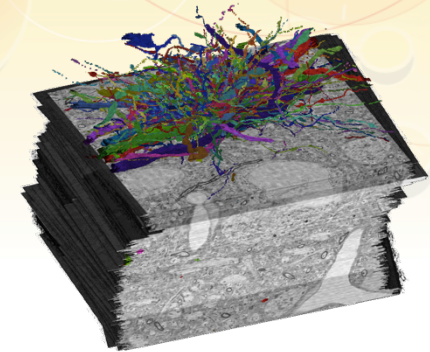
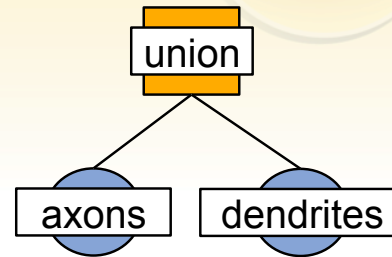
- name, size, excitatory, vesicle number, etc.



Operators

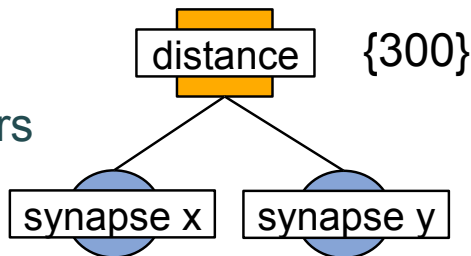
- Set operators

- union, intersect, etc.
- project
- group



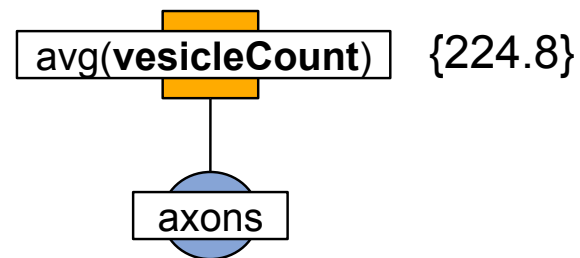
- Spatial operators

- distance



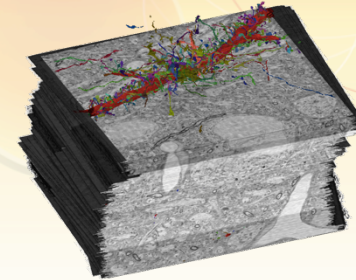
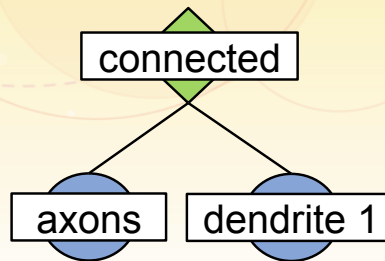
- Statistical operators

- avg, count, dev, etc.

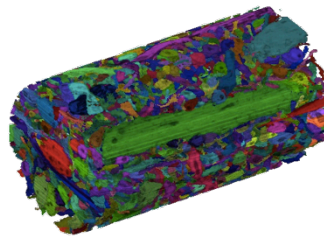
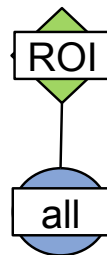


Predicates

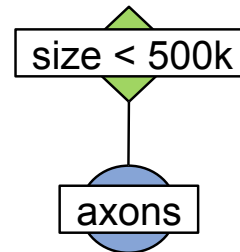
- Topological predicates
 - connected, part of



- Spatial predicates
 - ROI, distance



- Object attribute predicates



Set Creator

Set Creator

Set Creator

Predicate/Operator

Input Sets

Add Input Remove Input

Other Input

Attribute : axon myelinated

Operator : <> Value : Yes

New Set Name

Filter-by-Attribute{ axon myelinated <> Yes }} (All Objects)

Create Set

Sets | Queries

Set	
1	All Objects

Delete Set

Save Sets/Queries Load Sets/Queries

Render ROI if available Display Set Reset Display

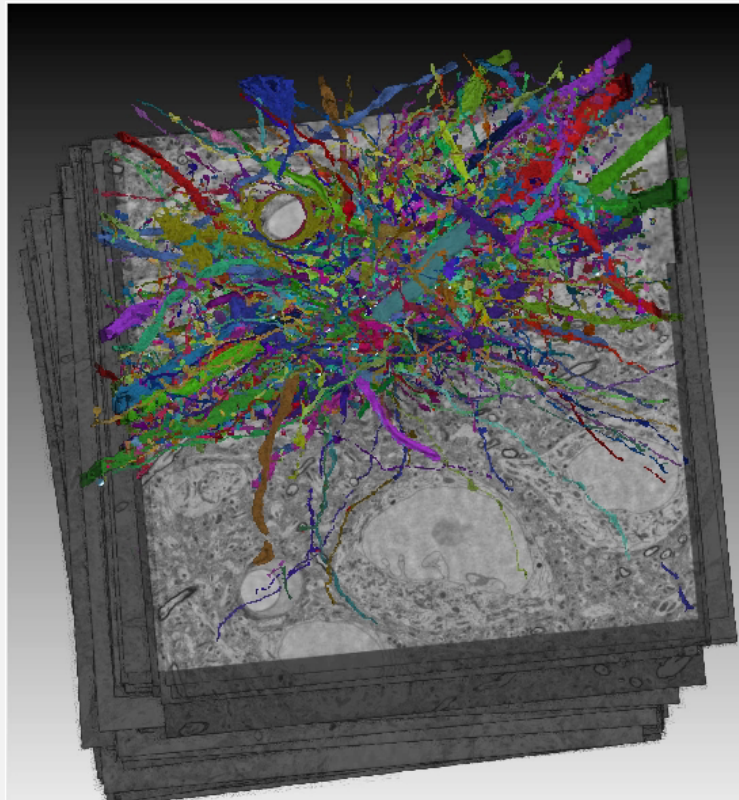
ConnectomeExplorer

File Transfer Functions Segmentation Properties Options Views Help

2D

Set Inspector

1	
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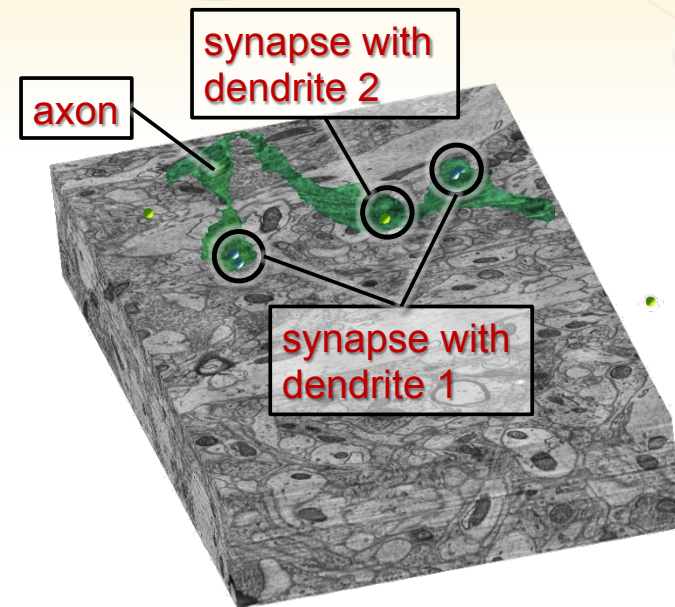


Set Inspe... Main Opti... Connectome Opti...

Use Case – Multiple Hit Axon

Goal:

- Analyze connectivity and morphology
 - Location of synapses
 - Sequential vs. non-sequential arrangement
 - Spatial distribution



User Feedback

- Two main user classes
 - Neuroscientist (hypotheses testing, exploration)
 - Data analyst (proofreading)

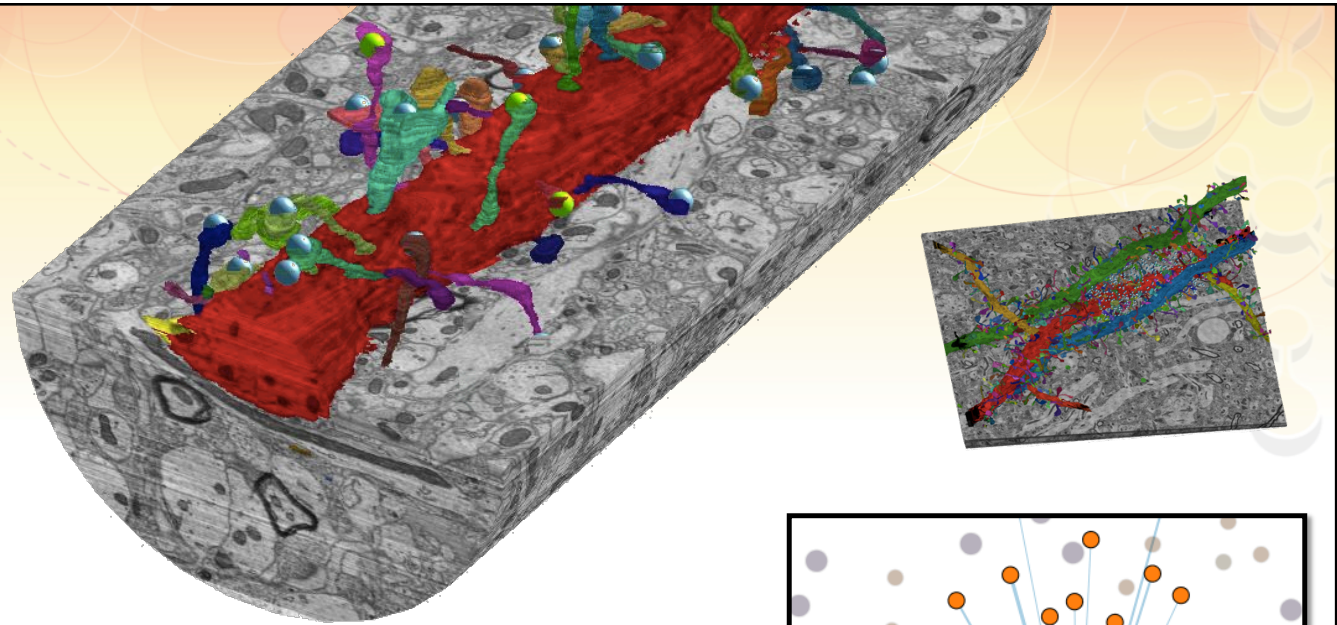
Feedback:

- ✓ Dynamic exploration & queries
- ✓ 3D visualization
- ✓ Connectivity graph view
- ✓ Linked views
- ✓ Quantitative analysis

Conclusions

- ✓ Powerful, yet simple query algebra
 - intuitive interactive exploration & analysis of large neuroscience data
- ✓ Built-in domain knowledge
 - makes queries simpler & more powerful
- ✓ Query-driven approach
 - on-the-fly query evaluation with petascale memory management
- ✓ Iterative query refinement
 - allows to quickly extend, combine, shuffle existing queries

Thanks



Acknowledgements:

Thomas Theußl (KAUST)

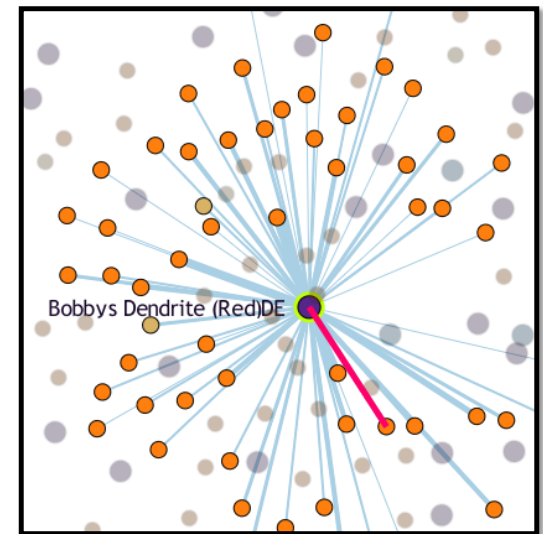
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http://seas.harvard.edu/~jbeyer/connectome_explorer.html



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