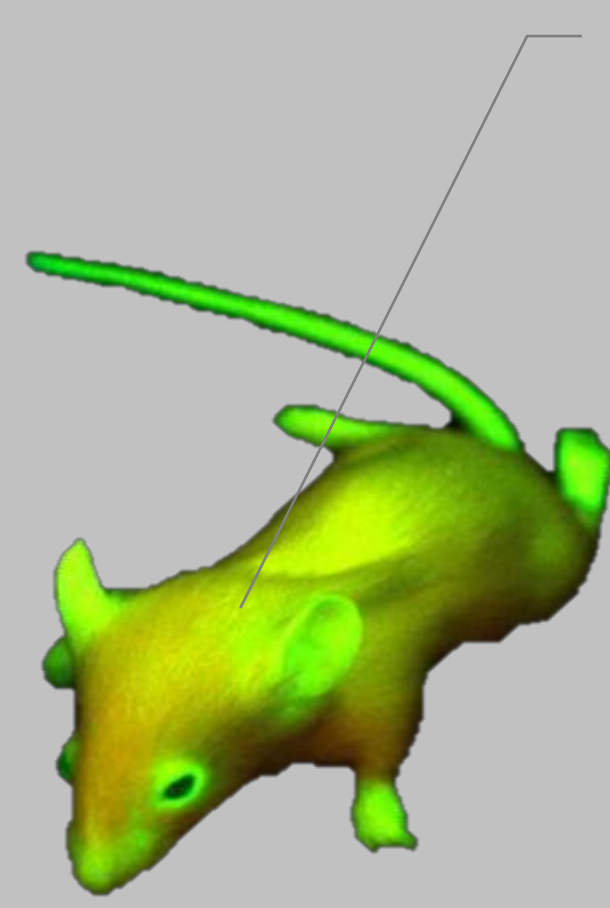



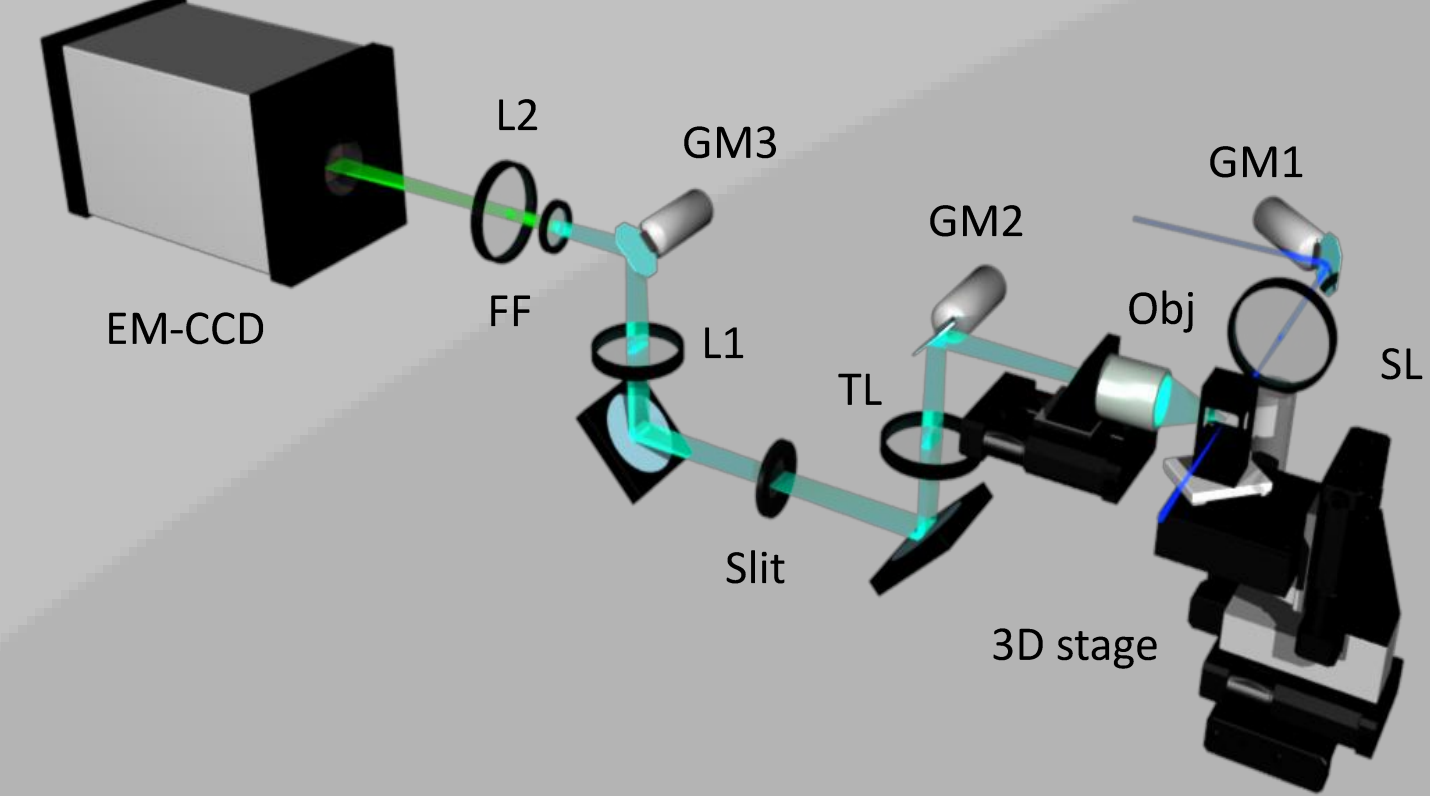
## Set up and testing of a High Performance Computational Infrastructure for processing and visualizing neuro-anatomical information obtained using CLSM



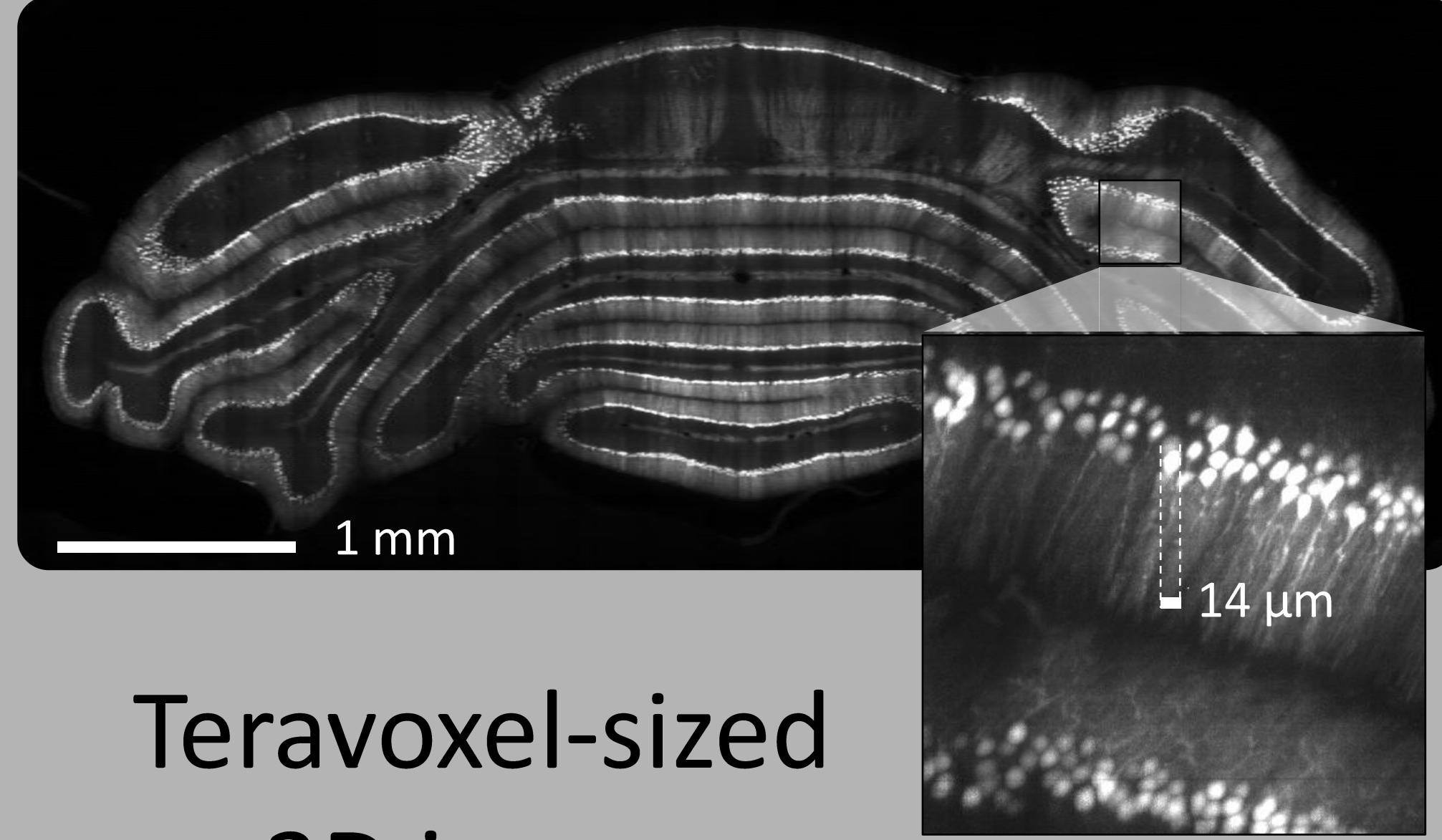
P10 L7-GFP mouse



Cerebellum



Confocal Light Sheet Microscopy (CLSM)



Teravoxel-sized 3D image



Biophotonics Group








### Projectome Environment

High Performance Computing platform running:

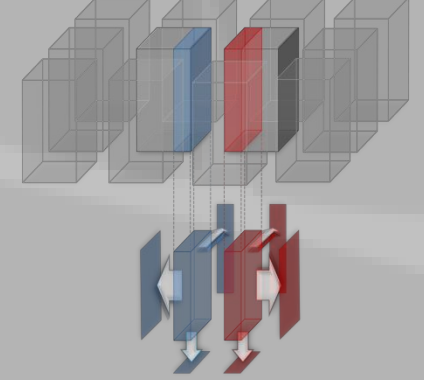
- Projectome Toolkit
- iRODS Data Grid
- Remote Visualisation Service

### Projectome Environment

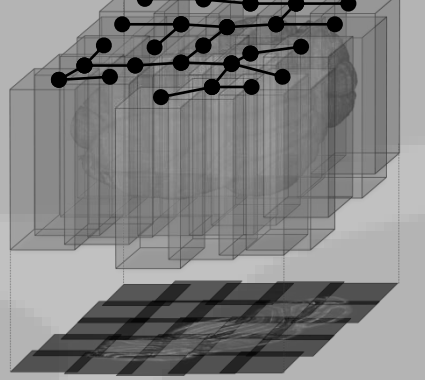


Storage Processing 3D Sticking Sharing

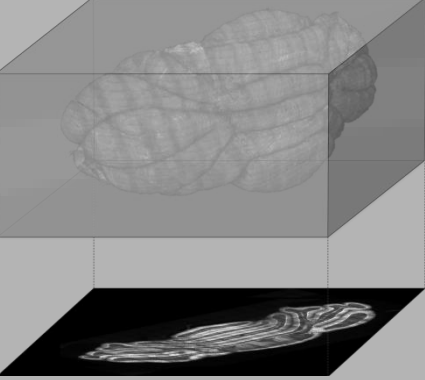
Pairwise displacements computation (Multi-MIP-NCC)



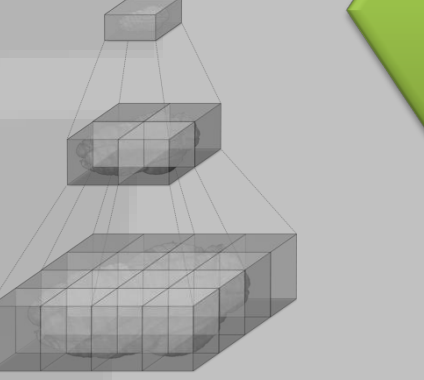
Globally optimal placement of stacks (Minimum Spanning Tree)



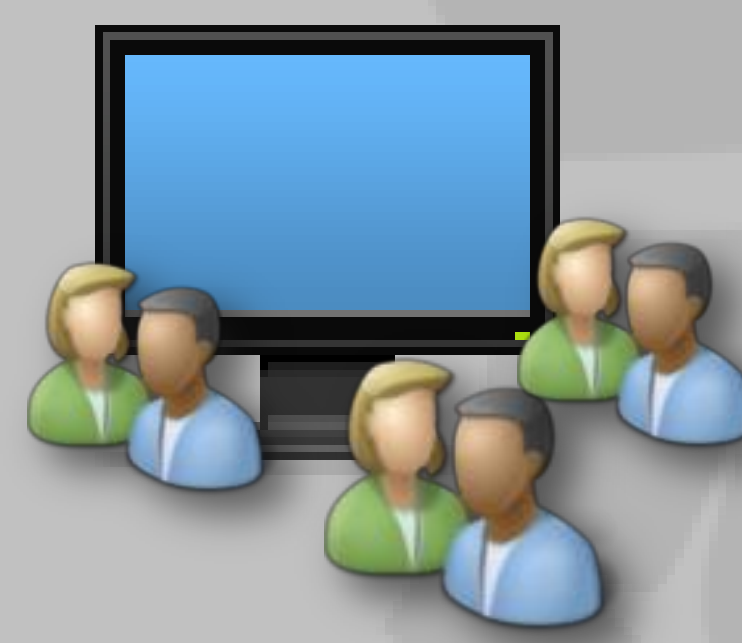
Merging of stacks (sinusoidal blending)



Saving volume at different resolutions

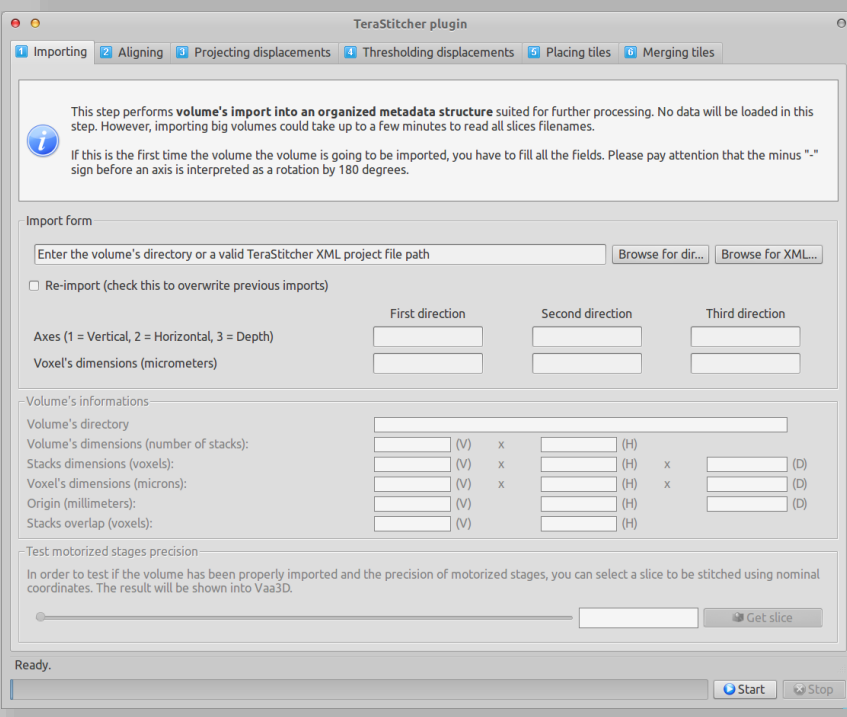


Visualization Annotation (under dev.) Data Mining & Clustering (under dev.) Knowledge extraction (under dev.) Features Extraction



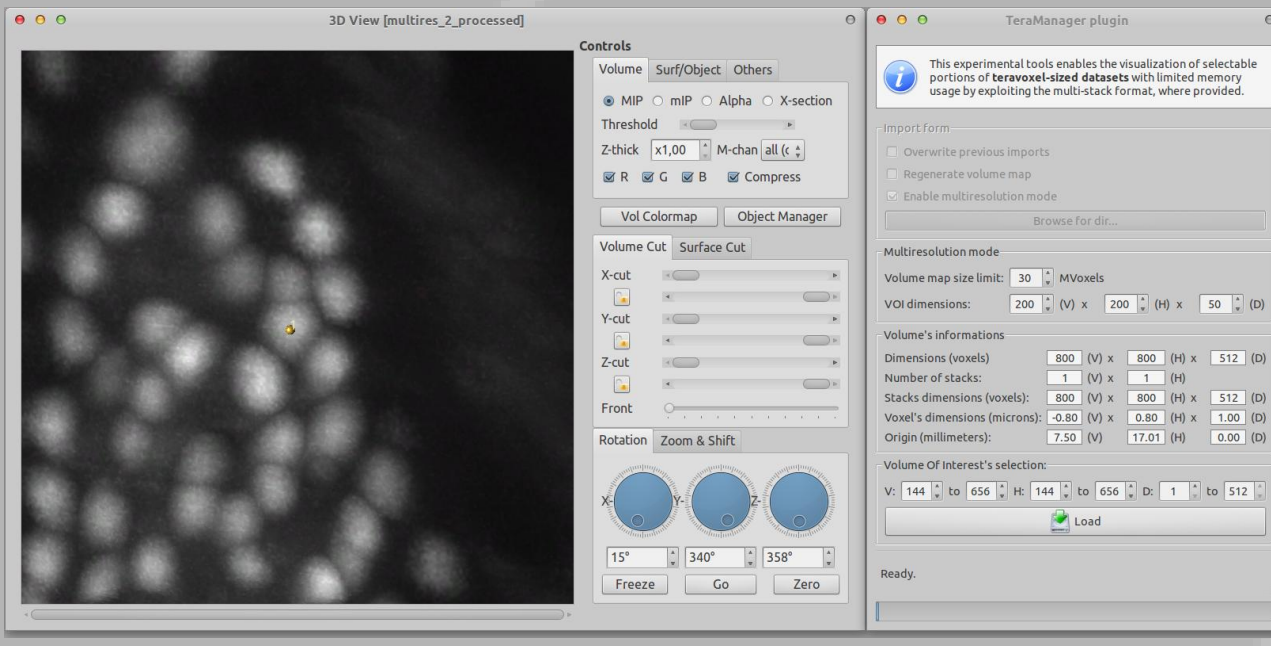
Projectome Scientists Scientists worldwide

### TeraStitcher (Vaa3D plugin)



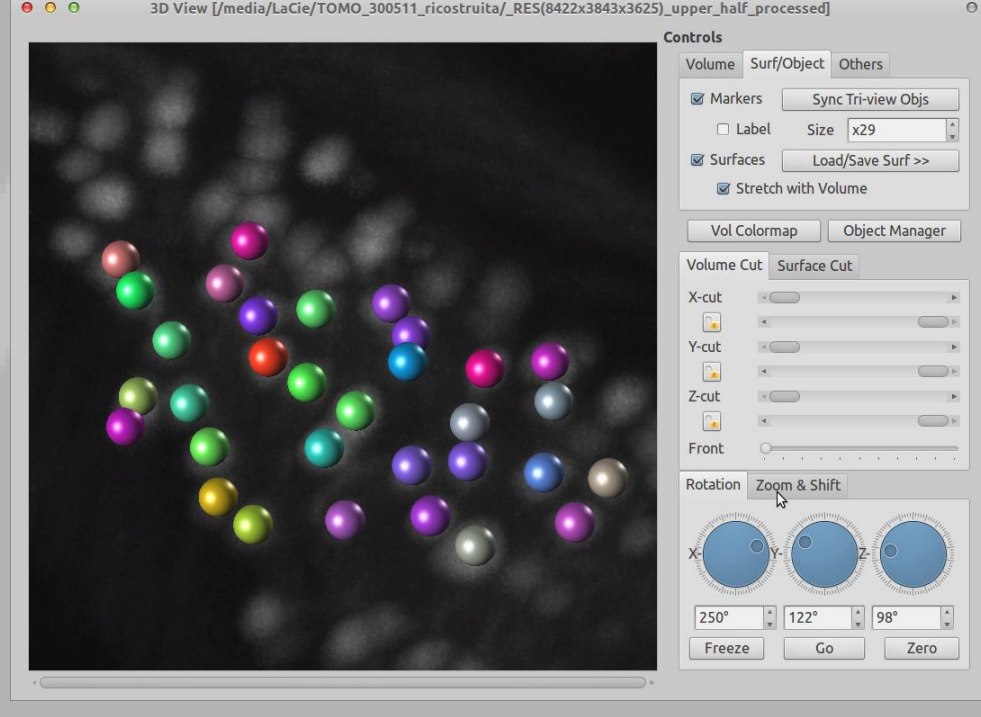
3D Stitching

### TeraManager (Vaa3D plugin)



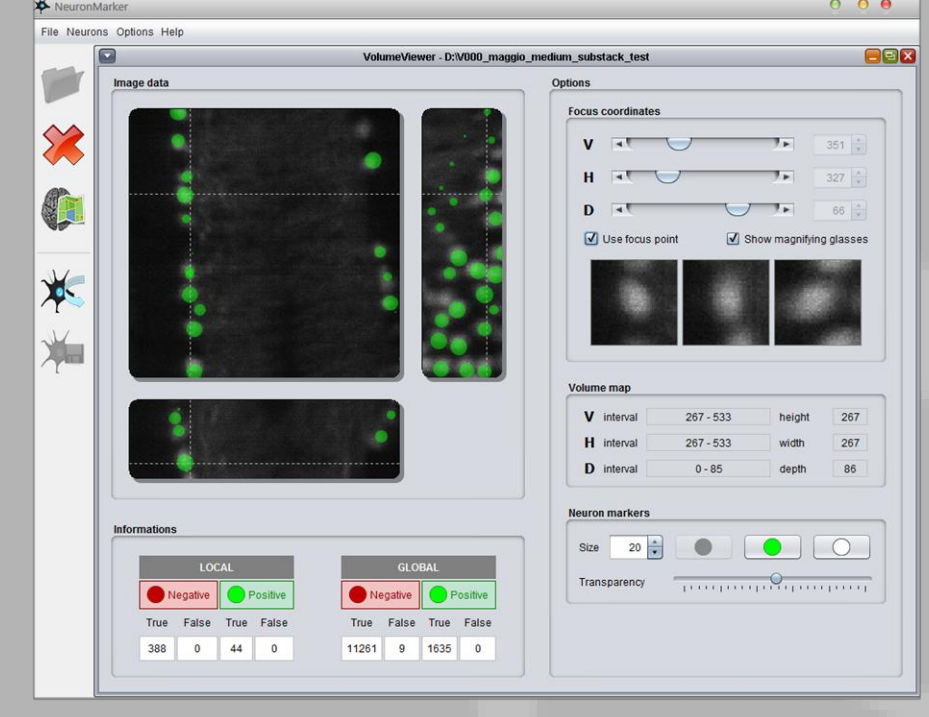
3D Visualization

### Vaa3D



Annotation, neuron tracing, brain registration

### NeuronMarker



Proofreading of neuron detector