

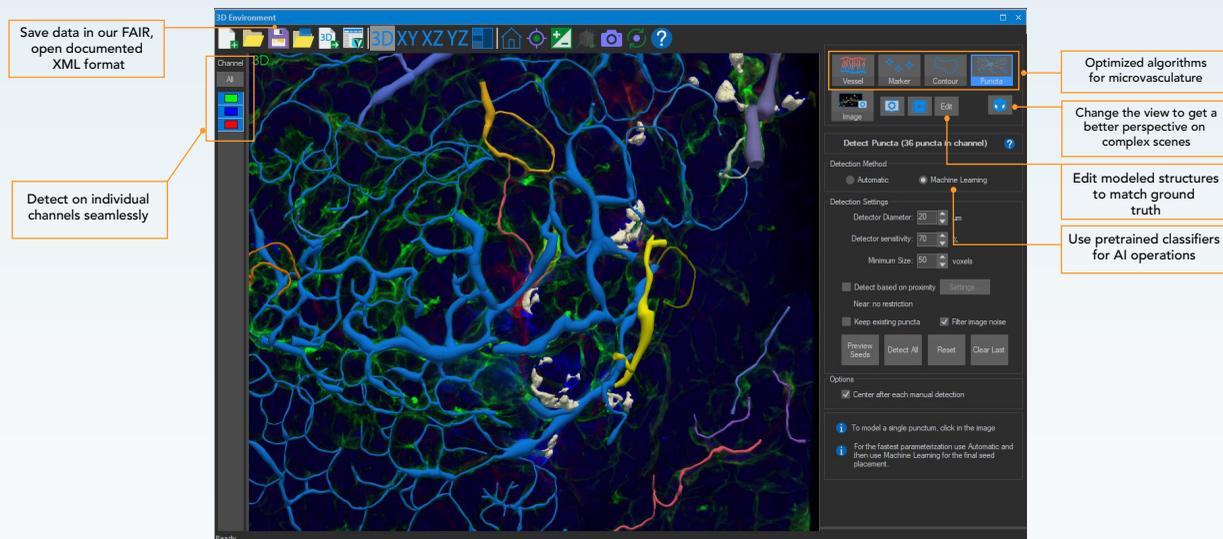
# Vesselucida® 360



## Quantitative and Visual Analysis for Blood Vessels and Microvasculature

### Quantify Vasculature Accurately

Use Vesselucida 360 to quickly model complex vasculature in an intuitive 3D workspace. Generate reliable, quantitative data on the length, connections, and complexity of microvessels to empower your research on angiogenesis in cancer, diabetic retinopathy, stroke and traumatic brain injury, and other conditions that affect microvasculature.



### Automatically Segment and Reconstruct Vasculature in 3D

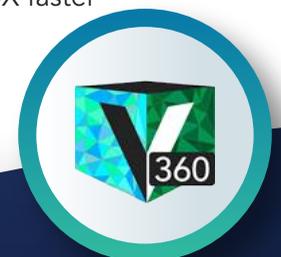
Collect more data in less time using Vesselucida 360.

- Automatically detect microvasculature, even in challenging specimens using sophisticated Vesselucida 360 algorithms
- Examine networks at multiple levels, including nodes, anastomoses, puncta, and multiple orientation and distance measurements
- Use fully automatic, semi-automatic, or manual tracing tools, all with editing capabilities, to generate data based on ground truth
- Process and trace many images at once using the intuitive batch-pipeline tool
- Analyze individual segments' surface and volume, as well as numbers of branch points and reconnections
- Work dynamically with large images in manageable sub-volumes, cut planes, and partial projections for better visualization

### Work with Images of Virtually Any Size in Virtually Any Format

Visualize your specimens using techniques that fit your research paradigm. Images obtained using micro-CT, confocal, two-photon, multichannel fluorescence, and light sheet microscopes are all compatible. Open large, complex images instantaneously using the new image and data handling engines that load large images up to 5000X faster in 3D and up to 250X faster in 2D.

Learn more: [mbfbioscience.com/products/vesselucida-360](https://mbfbioscience.com/products/vesselucida-360)



# Vesselucida® Explorer

## Turn Your Data into Results



### Quantitative Analysis of Morphological Data from Vesselucida 360

Vesselucida Explorer is the analytical software companion included with Vesselucida 360. Use it to perform extensive quantitative morphometric analyses designed specifically for microvasculature models and serial-section reconstructions.

### Far More than Just Numbers

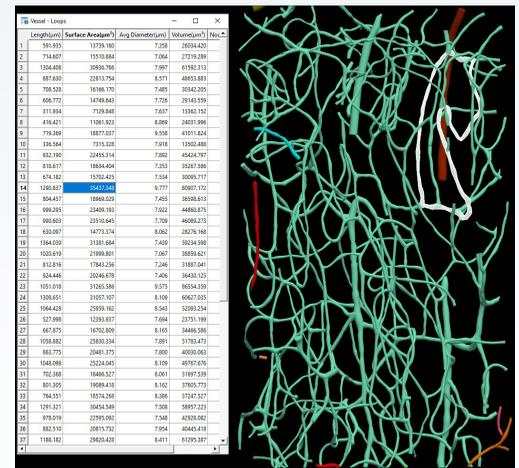
Perform dozens of analyses that calculate and graph hundreds of metrics.

- Characterize structure, distribution, orientation, and colocalization of features in your microvasculature models
- Generate comprehensive quantitative data tables that you can easily export for use in statistical and spreadsheet software
- Create graphical displays to visualize quantitative results in intuitive ways and use these high-quality displays in figures for publications and presentations

### A Mountain of Data without the Data Overload

Find answers in your data and present to others using Vesselucida 360 and Vesselucida Explorer.

- Load multiple data sets into Vesselucida Explorer for rapid, batch quantitative data analyses with no scripting or programming skills needed.



Learn more: [mbfbioscience.com/products/vesselucida-explorer](http://mbfbioscience.com/products/vesselucida-explorer)



## About MBF Bioscience

A rich history of creating the future of neuroscience.

MBF Bioscience develops advanced tools for collecting and analyzing accurate, reproducible data from histological specimens, 2D and 3D microscope images, and freely moving *C. elegans* so that scientists can better understand brain diseases and processes at a cellular level.

Our products have helped researchers publish over 17,000 peer reviewed papers.

## What our customers say

“After examining different vessel quantification tools for use with neovessel formation in the heart, we chose Vesselucida 360 as it offered us flexibility in viewing & adjusting vessel geometries in 3D to accurately represent our microCT dataset.”

Dr. Kareen Coulombe  
Brown University

“Vesselucida 360 enables us to visualize and quantify microvascular networks in a way that has not been done before. The ability to render 3D maps is integral to analysis of network structure, remodeling and blood flow distribution.”

Dr. Nicole Jacobsen  
University of Missouri

