

Quantitative Image Analysis and 3-D Digital Reconstruction of Aortic Valve Leaflet

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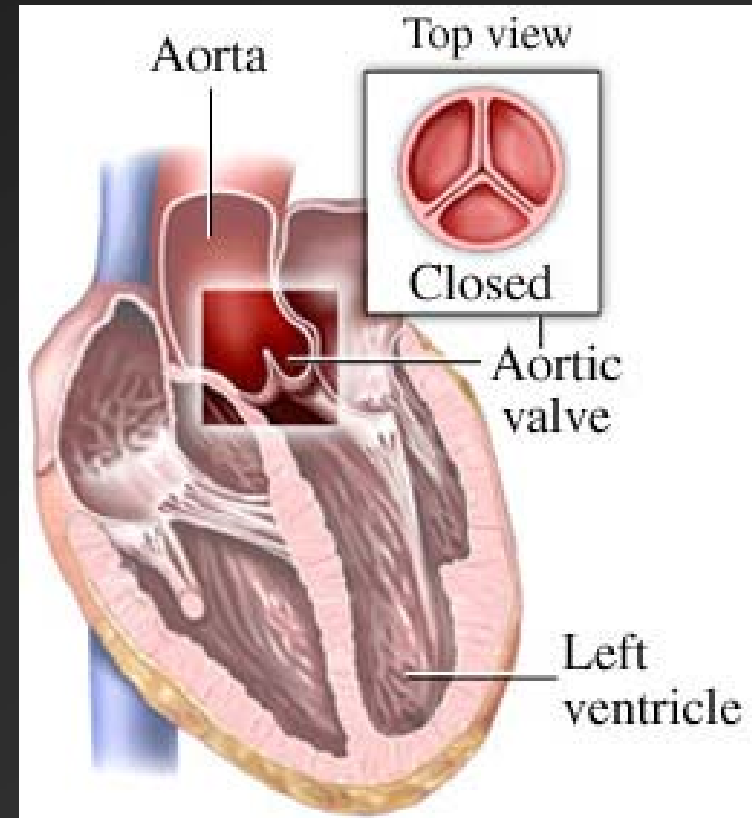
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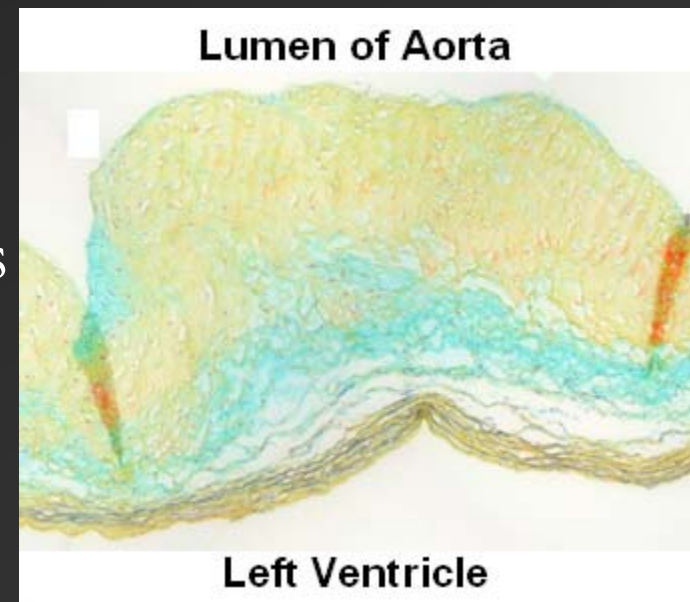
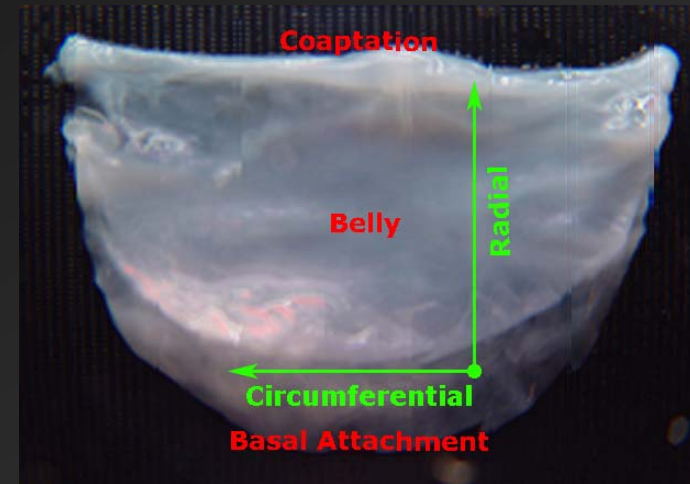
Introduction

- Aortic valve (AV): most commonly replaced valve
- Tissue engineering (TE) valve offers customized availability, growth potential, durability, and biocompatibility.
- Must first understand structure of the valve to replicate its function by TE



Background

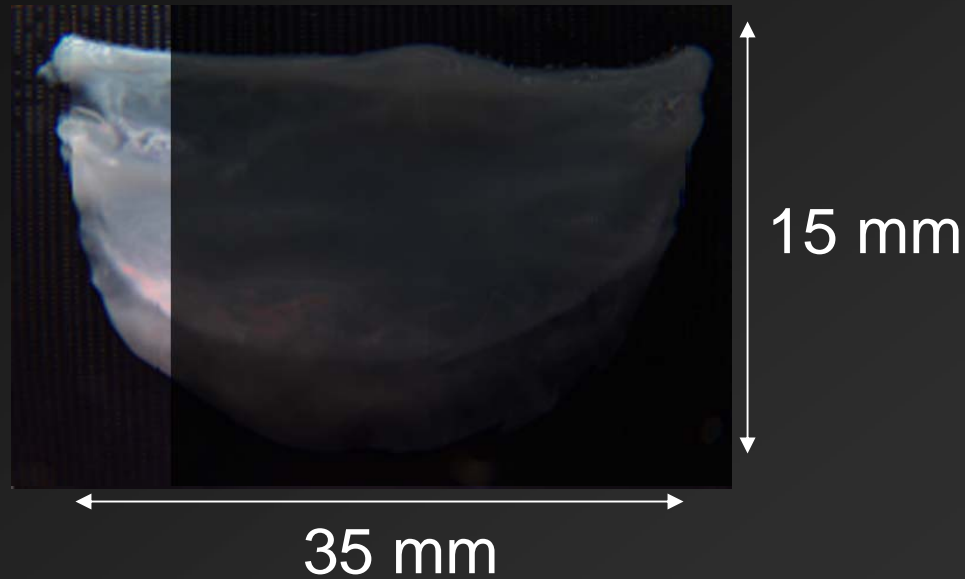
- Each valve is composed of three leaflets (right-coronary, left coronary, non-coronary)
- Each leaflet is composed of 3 cell layers
 - Fibrosa: Collagen and little elastin
 - Spongiosa: Glucosaminoglycans (GAGs)
 - Ventricularis : Collagen and elastin



Objectives

- Original objective: Construct 3D representation of a porcine right coronary AV leaflet containing...
 - Cell count and distribution
 - Layer thickness variations
 - Track collagen and elastin fibers
- Technological limitations bifurcated the objective
 - Quantification
 - 3D reconstruction

Status



- Total histological sections: ~ 300 slides
- Imaged, quantified, and 3D reconstructed 50 slides

Histology and Image Acquisition

- 5 μm thick circumferential slices fixed in formalin stained with Movat's pentachrome
 - Collagen = Yellow
 - GAGs = Blue
 - Nuclei and Elastin = Dark Purple
- Acquisition
 - Quantification: 17 slices, spaced 90 μm apart, were digitally captured using bright field microscopy at 20x and montaged
 - 3D Reconstruction: 50 slices, spaced 10-15 μm apart were scanned individually using slide scanner

Principles of Image Analysis

- Contrast
 - 256 shades of gray in a 8-bit monochrome image
 - Defines edges/borders of objects

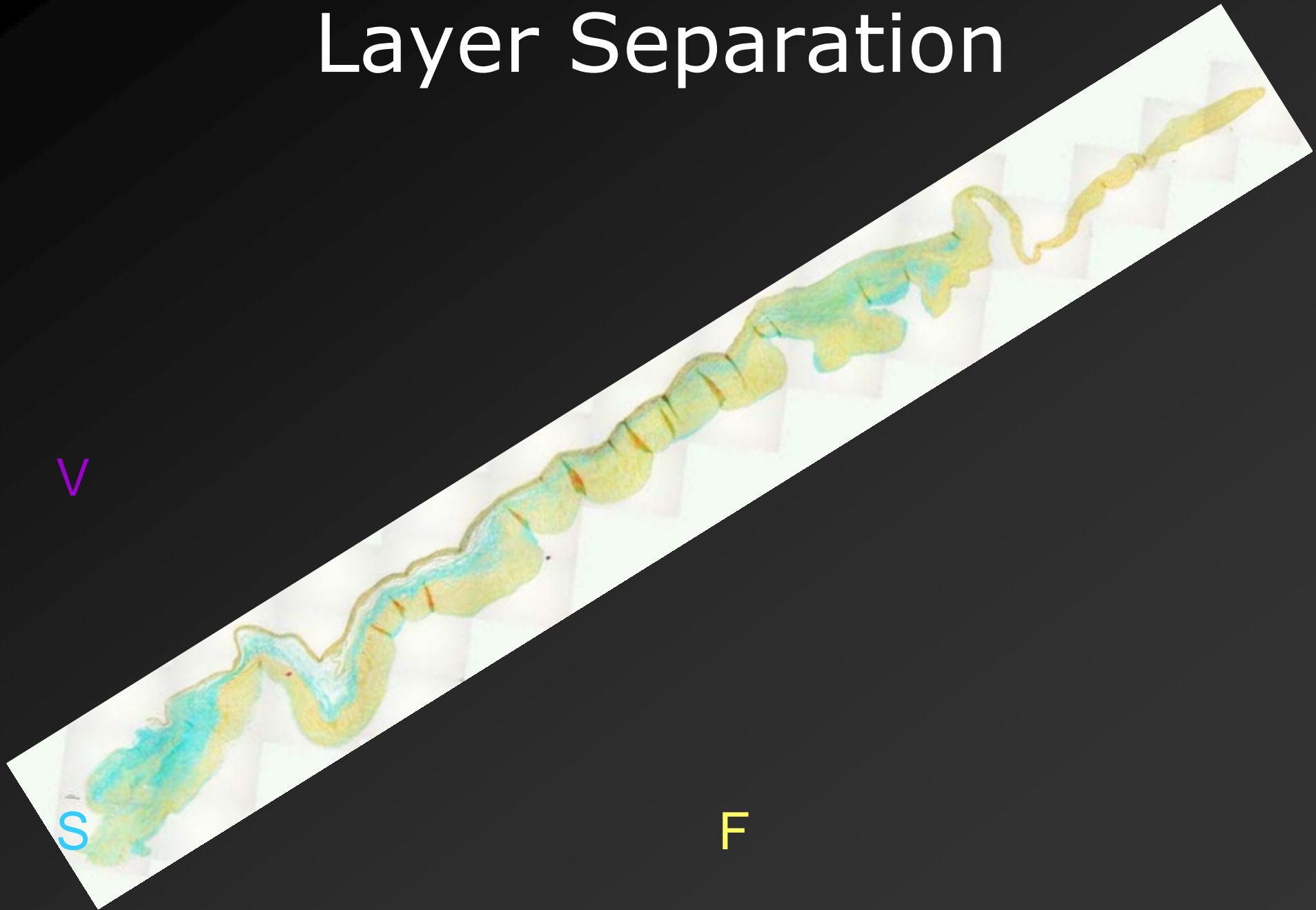


- Threshold
 - Enhances contrast by dividing image into two category
 - Objects within the image (subsets of image data) defined by contiguous pixels similar in intensity
 - Automatic threshold selection uses mode pixel intensity

Color Separation



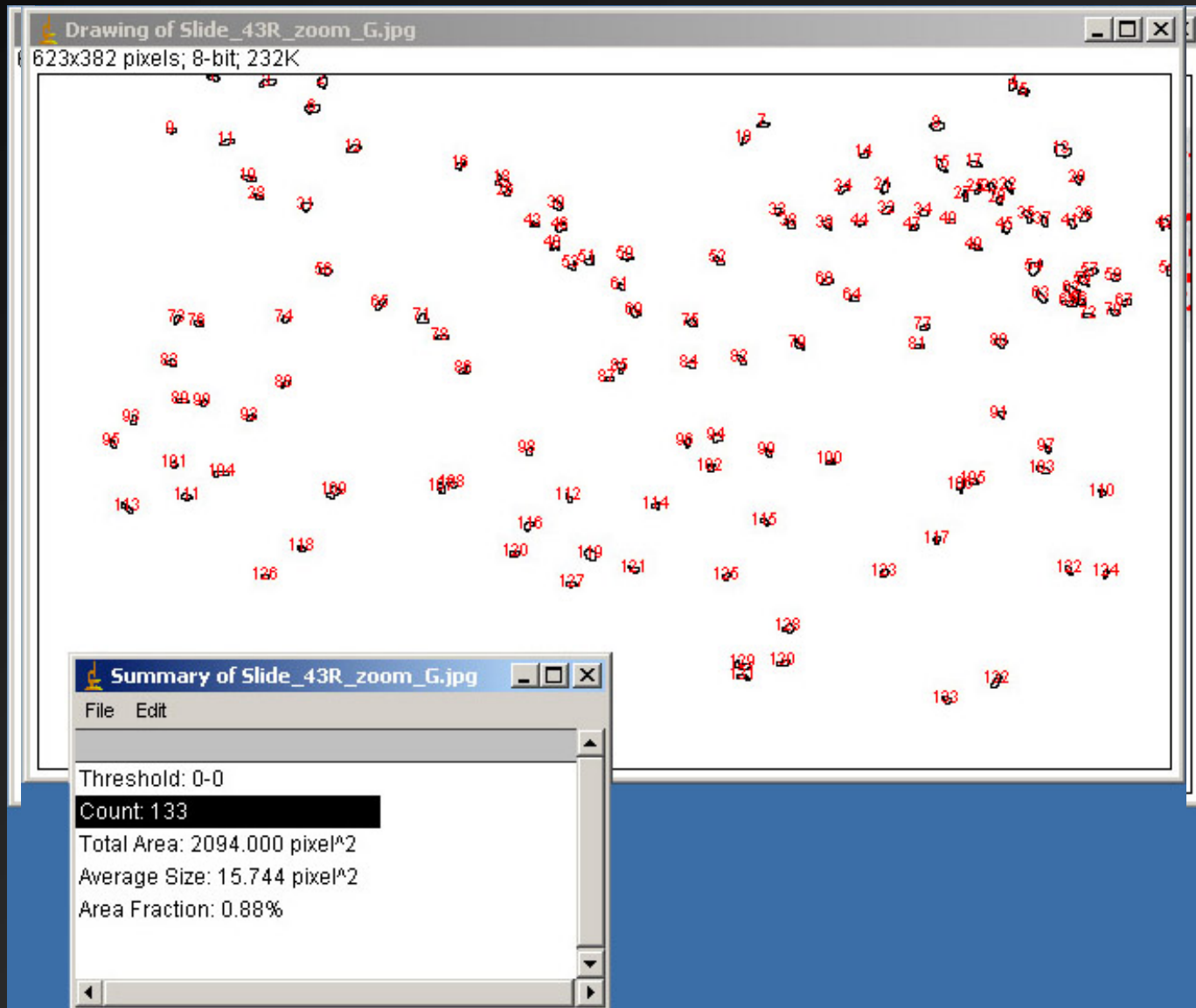
Layer Separation



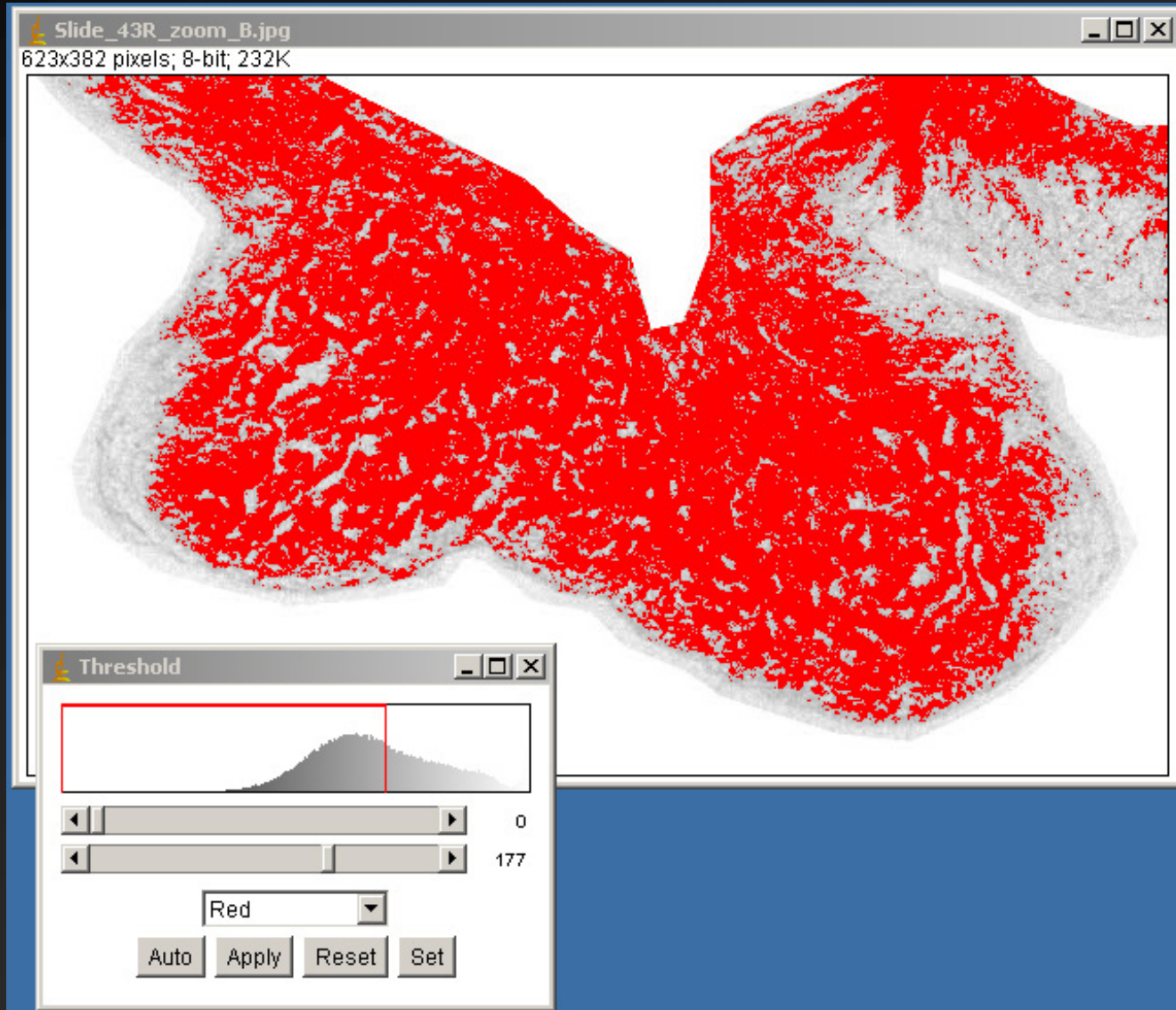
Cytometry

- Cell nuclei counting completed with particle analysis function in NIH's Image J
- RGB image converted to monochrome on the green channel
- Threshold is user defined
- Definition of a cell
 - size: 9-100 pixel²
 - circularity: 0.6-1.0

Particle Analysis of Nuclei



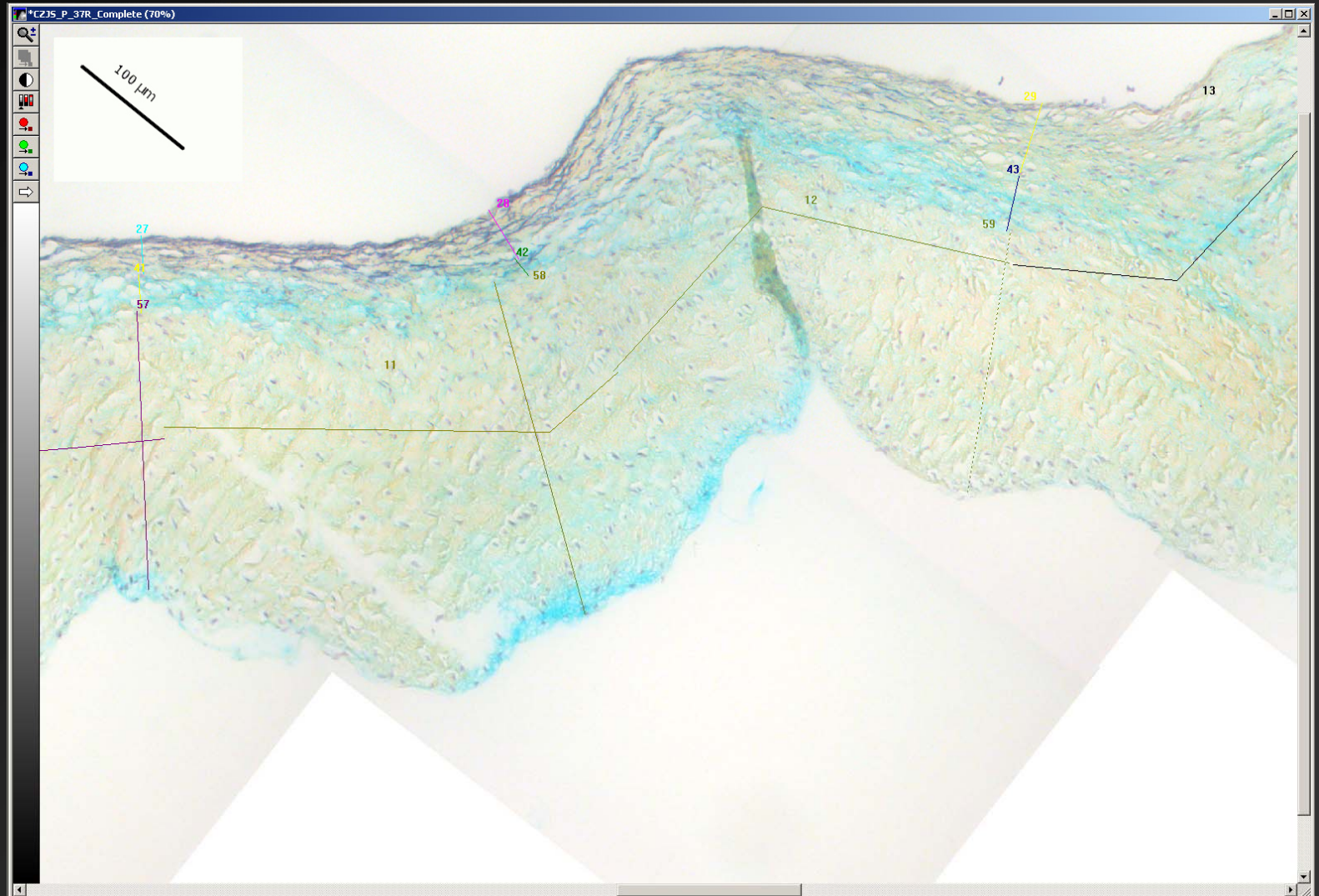
Collagen and Elastin Content



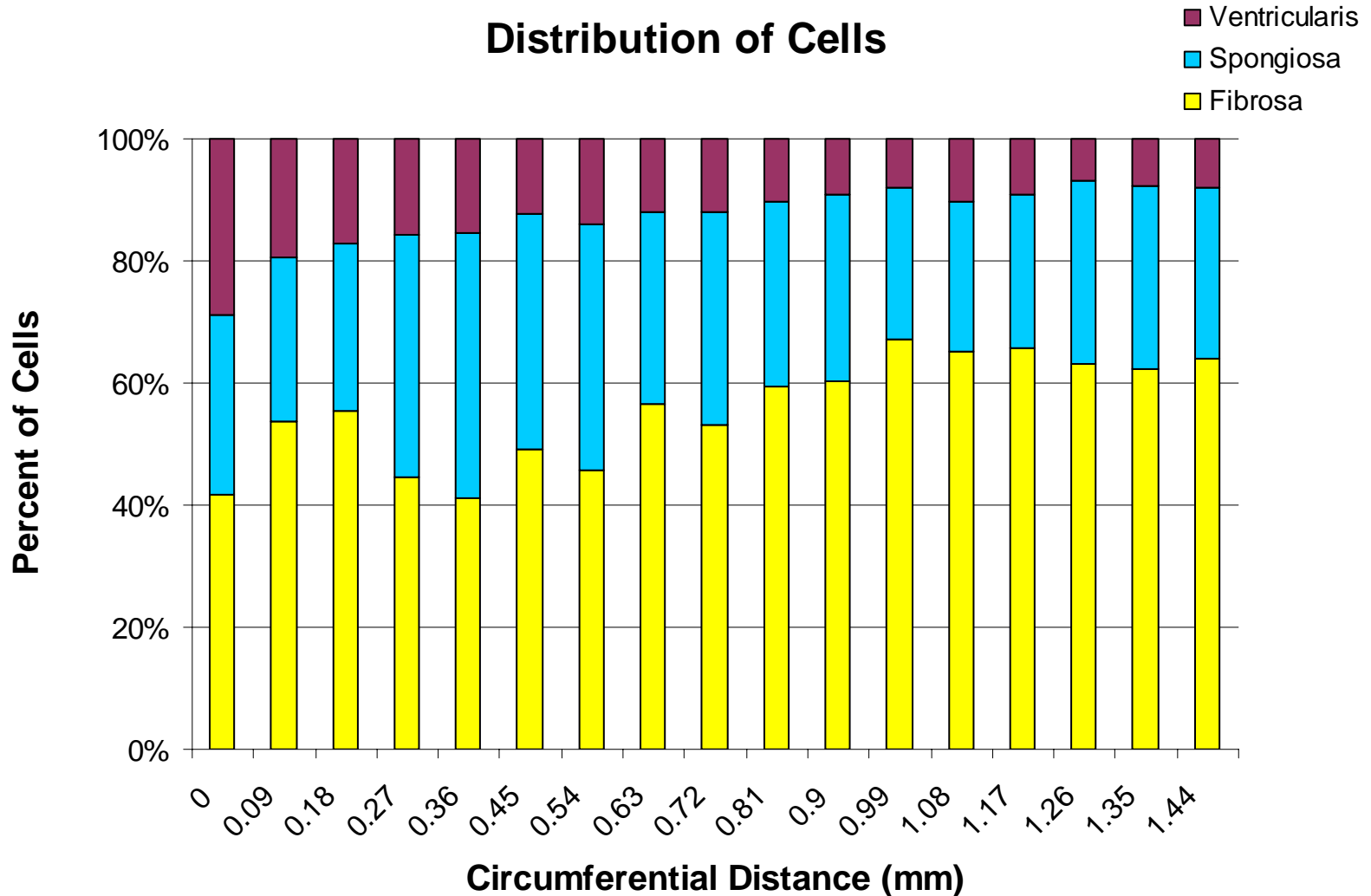
Layer Thickness Measurement

- Scale bar added during image acquisition used to establish calibration between pixels and μm in MetamorphTM
- Take representative samples, spaced apart by $500 \mu\text{m}$, along the radial length of each slice
- Results indicate
 - Average layer thickness
 - Topographic representation
 - Local variations

Measuring Thickness



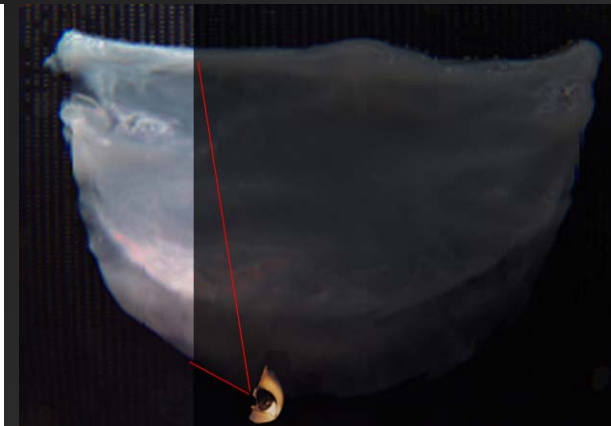
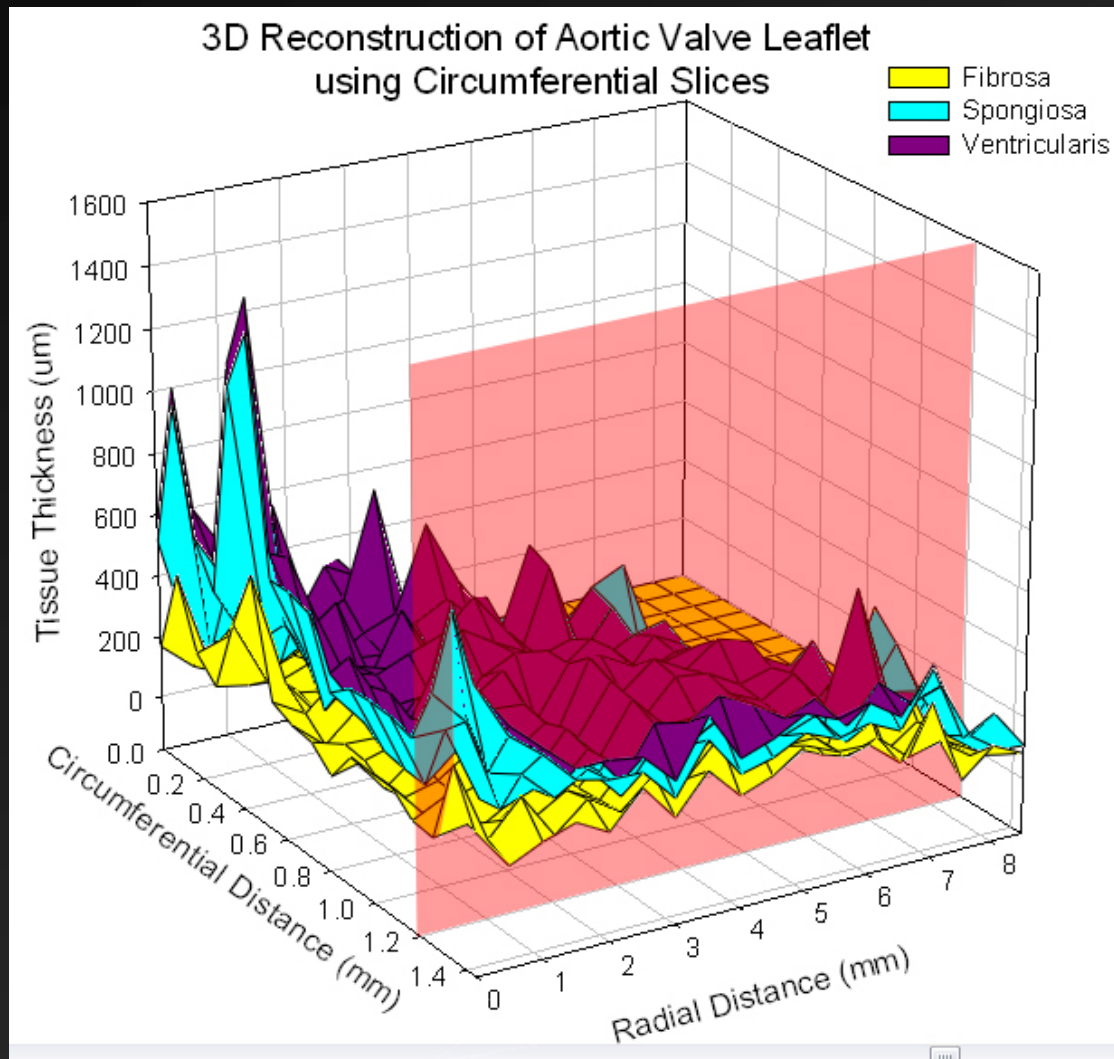
Quantification Results



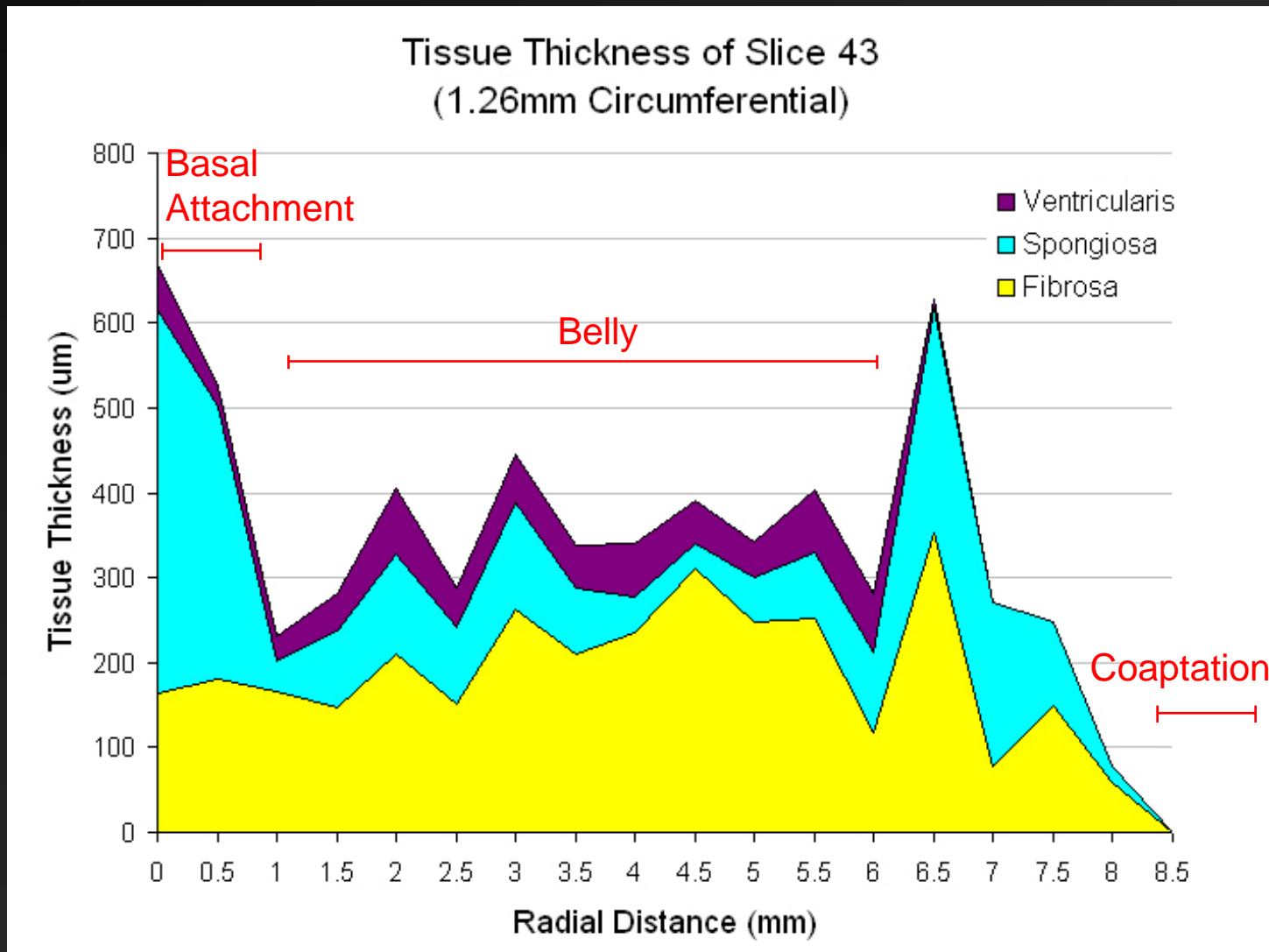
Quantification Results (cont'd)

- Collagen and Elastin Content
 - Fibrosa: $48.2 \pm 6\%$ area occupied by collagen
 - Ventricularis: $54.3 \pm 6\%$ collagen
 $39.3 \pm 5\%$ elastin
 - Need for comparison with literature values (entire tissue composition: 13% elastin and 50% collagen by dry weight)
- Average thickness
 - Fibrosa: 150-230 μm (~ 100 -350 μm)
 - Spongiosa: 110-200 μm (~ 70 -250 μm)
 - Ventricularis: 40-60 μm (~ 50 -150 μm)

Topography of Layer Thickness



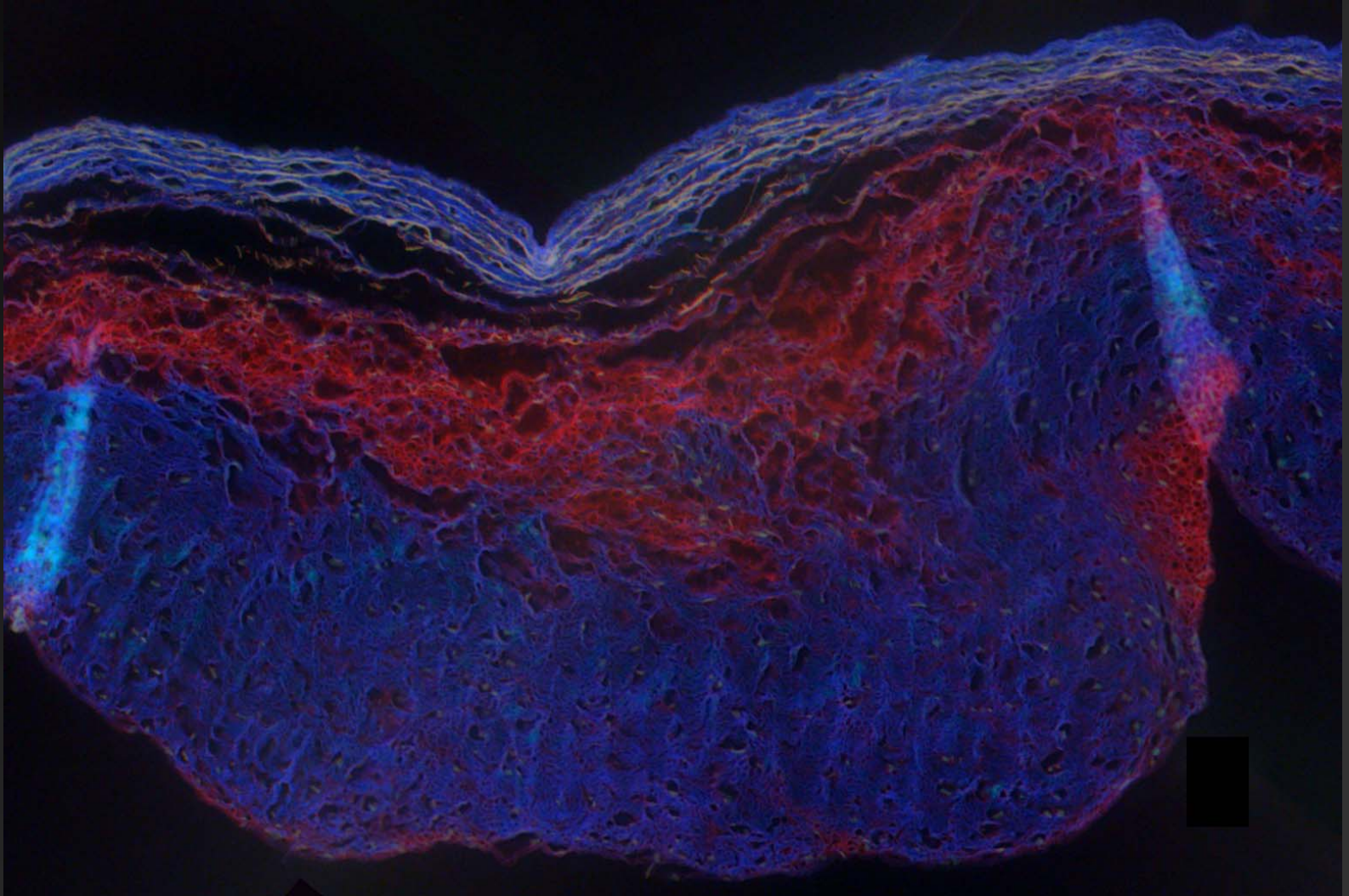
Regional Variation



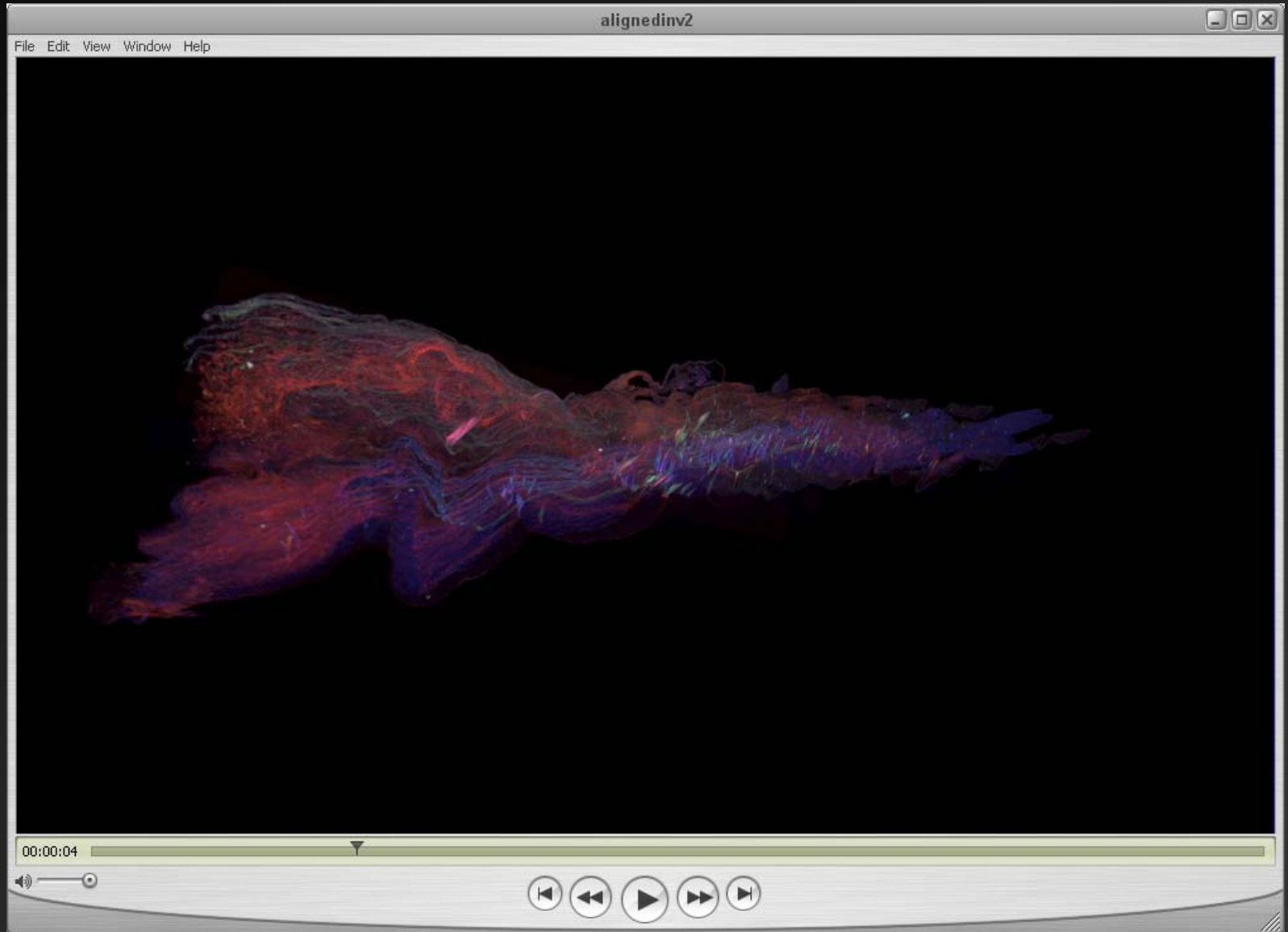
3D Reconstruction

- Images of 50 slices, spaced 10-15 μm apart, were digitally aligned by morphology and stacked to construct a 3D representation
- Software designed for fluorescent microscopy so image colors had to be inverted
 - Collagen = Blue
 - GAGs = Red
 - Nuclei and Elastin = Yellow
- 3D representation allows user to visualize the leaflet better than topography

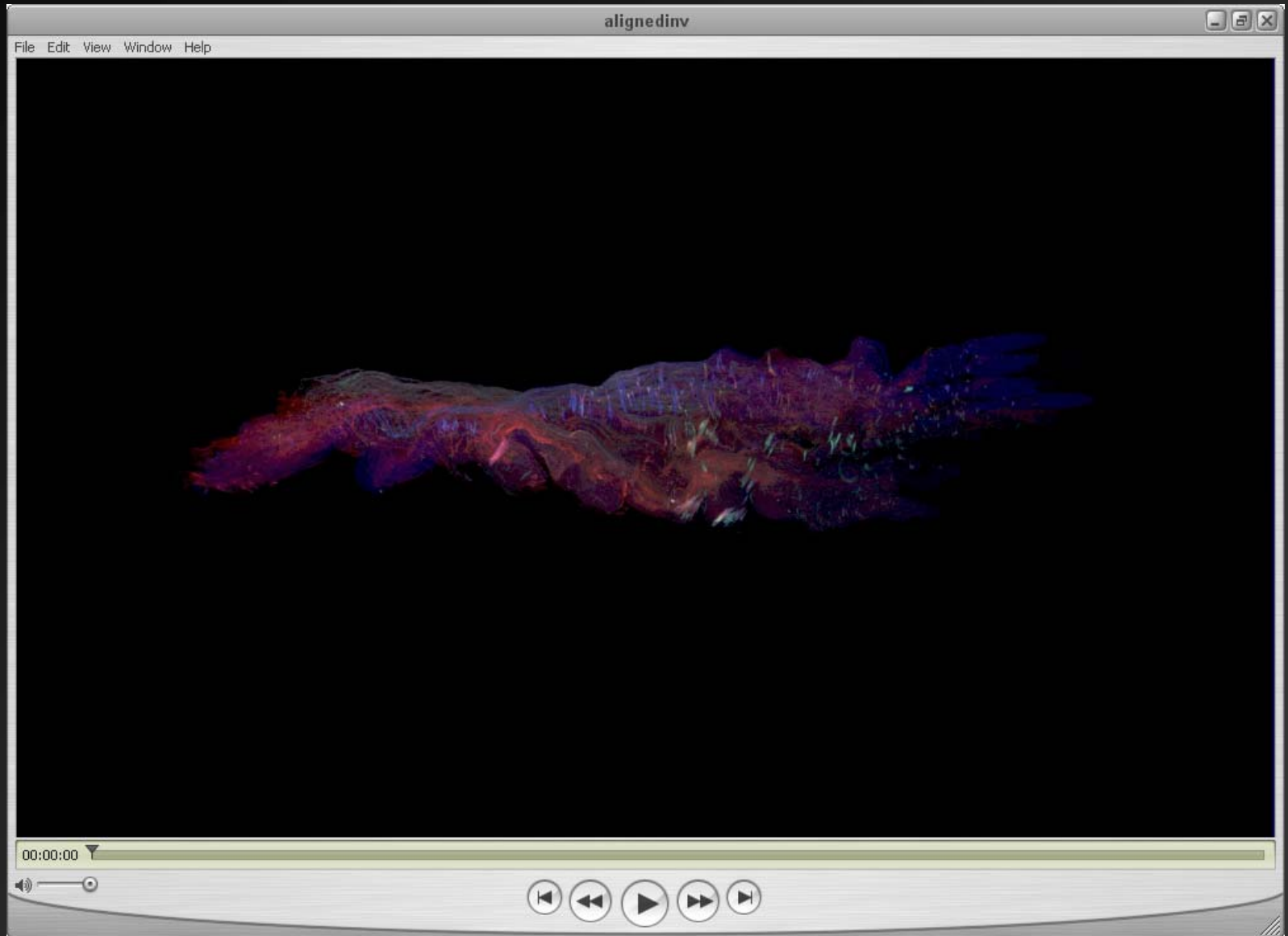
Inversion



Ooo....



Ahh...



Conclusion and Future Work

- Preliminary results calls for completion of the entire leaflet
- Statistical validation of quantification upon completion
- Explore other imaging techniques (fluorescent microscopy, X-ray, ultrasound, acoustic microscopy, SEM)
- Construct a 3D representation containing quantitative information
- Use 3D reconstruction to simulate and visualize dynamic response to applied load

Acknowledgements

- Engineered Tissue Mechanics Laboratory, Department of Bioengineering, University of Pittsburgh (Dr. Michael Sacks, John Stella, et al.)
- Center for Biologic Imaging, University of Pittsburgh (Dr. Simon Watkins, Jason Devlin, Stuart Shand, et al.)
- Department of Computational Biology, University of Pittsburgh
- Developers of NIH's Image J

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¿Questions?

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