

High-Level Sprout Geometry Extraction and Analysis of In Vitro Angiogenesis

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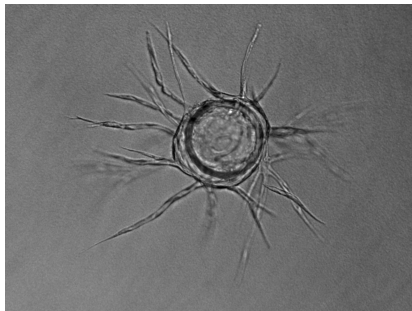


Figure: Before

Spreadsheet Report
Sprout Counts
:
Branching Factor

Figure: After

- ① Motivation
- ② Fibrin Gel Bead Sprouting Assay (FGBSA)
- ③ Methodology
- ④ Results

Solid tumors have an avascular (no nearby blood vessels) growth phase that allows for an approximate maximum size of 1-2mm in diameter¹.

¹Robert S. Kerbel. Tumor angiogenesis: past, present and the near future. *Carcinogenesis*, 21(3):505-515, 1999.

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Think of the size of *very coarse sand*.

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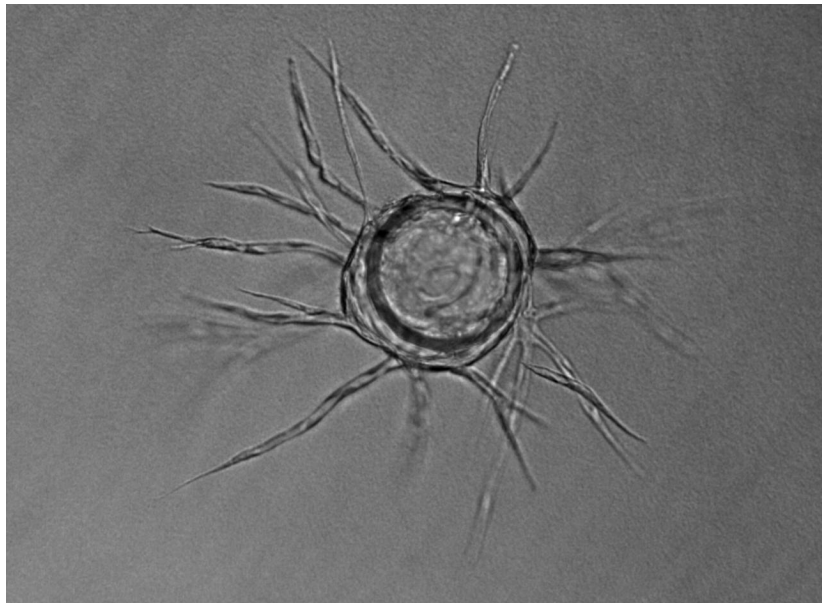
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Tumor angiogenesis enables relentless tumor growth and metastasis.

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FGBSA Image

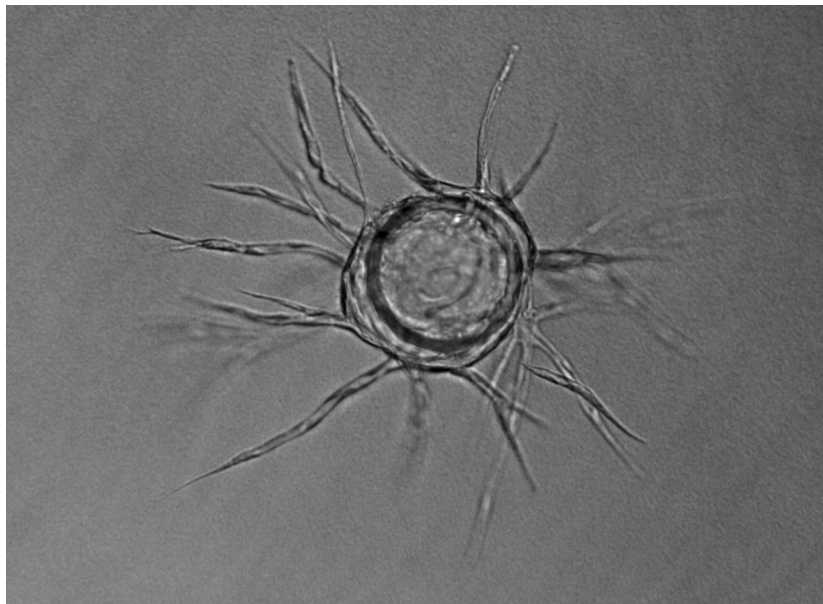


- ① Sprout Restoration
- ② Sholl Analysis

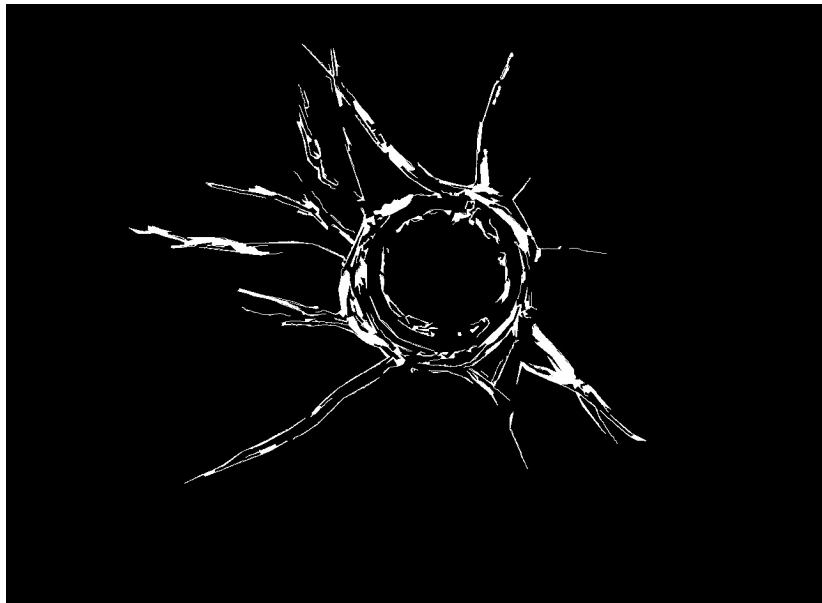
- 1 Edge Detection and Polygon Approximation
- 2 Bead Detection (Hough Transform)
- 3 Non-Sprout Detection
- 4 Dilation for Approximate Centerline
- 5 Thinning and Pruning
- 6 **Sholl Analysis**

Expanded sprout restoration methods

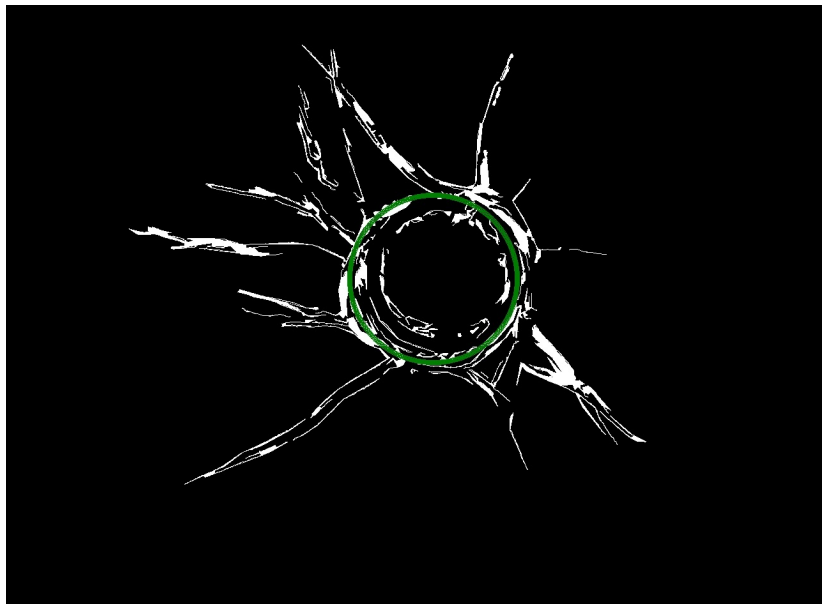
Original Image



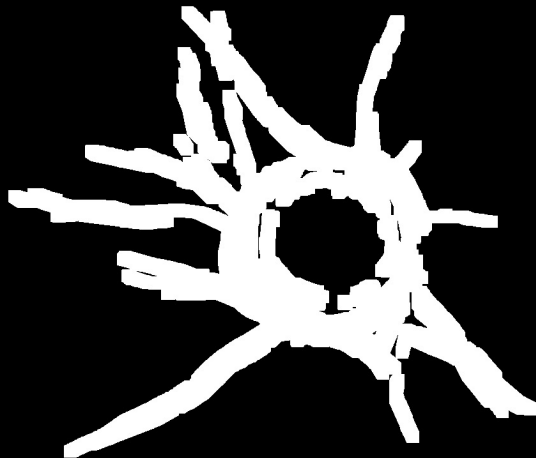
Edge Detection and Polygon Approximation



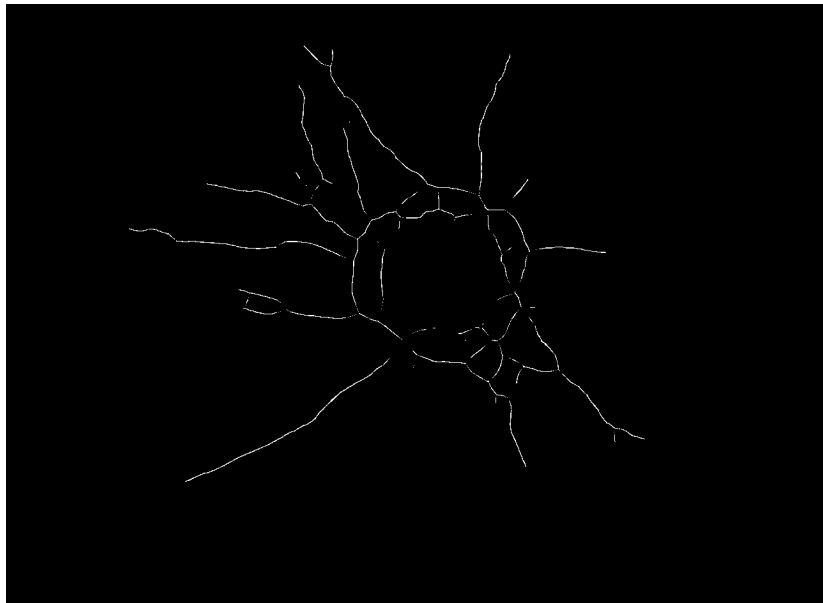
Bead Detection (Hough Transform)



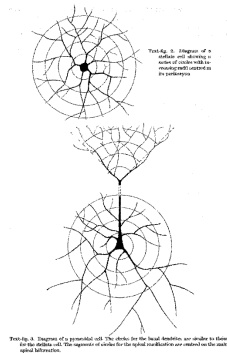
Dilation for Approximate Centerline



Thinning and Pruning



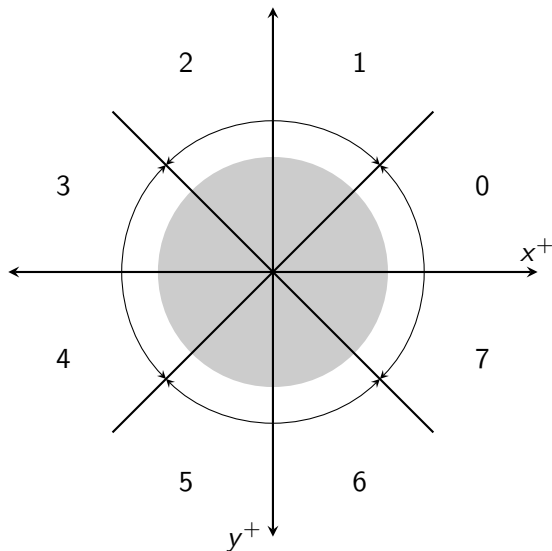
Sholl Analysis: Why



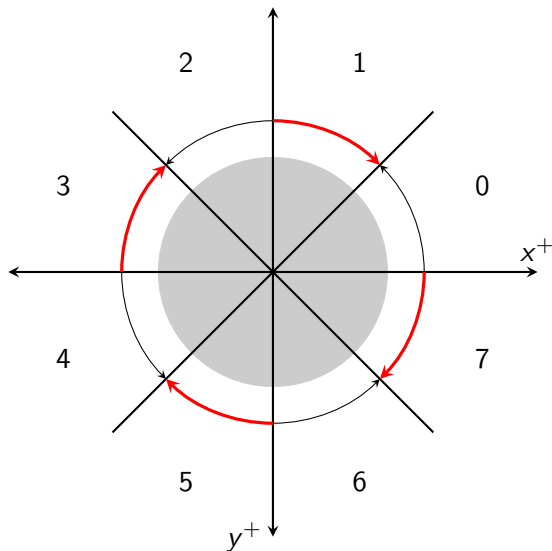
- No tracing required
- Morphometric descriptors can be obtained

Figure: D. A. Sholl. Dendritic organization in the neurons of the visual and motor cortices of the cat. *J Anat.*, 87(4):387406, 1953.

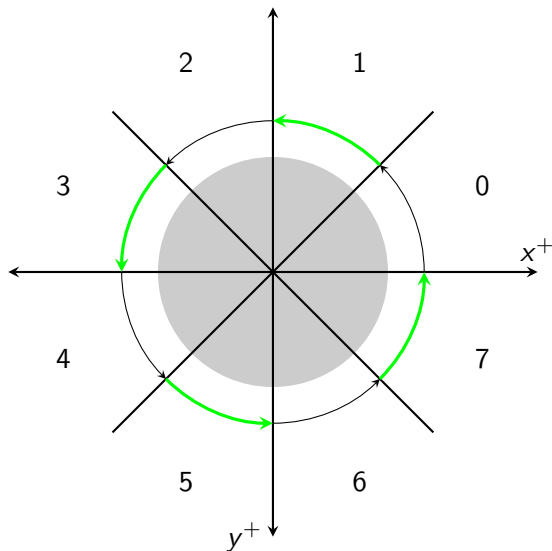
Sholl Analysis: Implementation



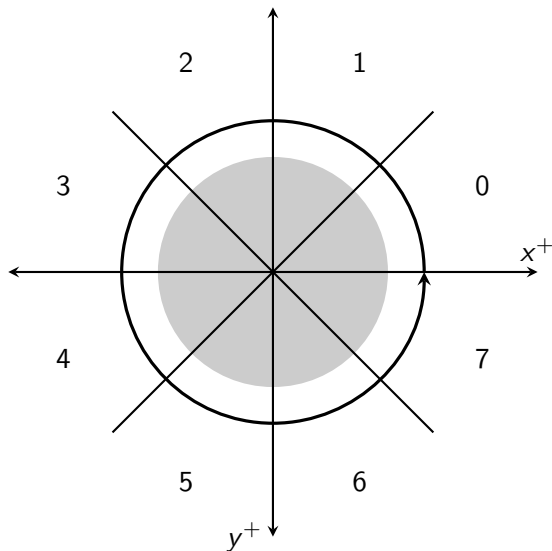
Sholl Analysis: Bresenham Circle Algorithm



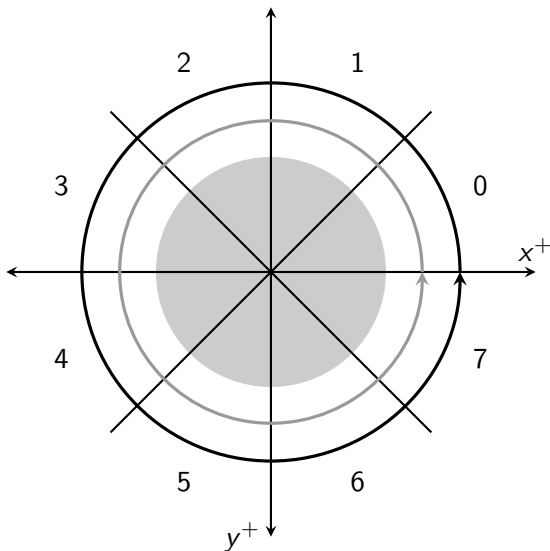
Sholl Analysis: Unordered Property of Circle Algorithm



Sholl Analysis: Ordered Bresenham Circle Algorithm



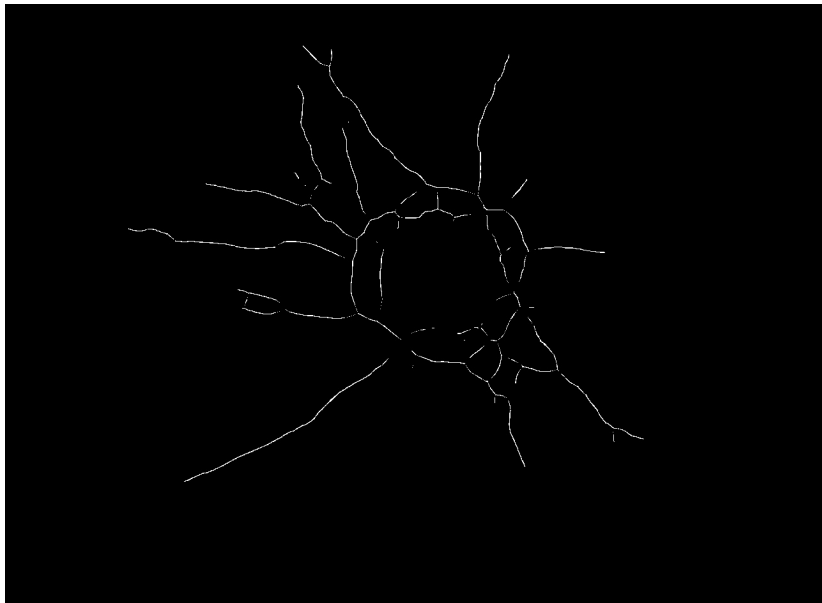
Sholl Analysis: Concentric Circles Analysis



Important descriptors

- ① Sprout enumeration
- ② Average sprout length
- ③ Branching factor (Shoenen Ramification Index)
- ④ Critical Value

Skeleton Reference



Sholl Analysis: Sprout Enumeration

Problem 1 Non-continuous sprouts

Problem 2 Bad initial radius

Solution Bounded crossings integration

- Mean of n crossings
- Median of n crossings

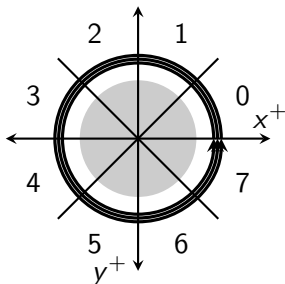


Figure: Crossing integration over intervals of size 3

Resulting sprout counts are compared with an expert observer using 15 images.

$$\text{RMSE} = \sqrt{\frac{1}{n} \sum_{i \leftarrow 1}^n (\hat{\mathbf{Y}}_i - \mathbf{Y}_i)^2} \quad (1)$$

Method Variant	RMSE
Median Integration Method	1.736
Ignoring isolated points	1.79
Mean Integration Method Benchmark	1.93
LIS on Large Image	3.45

Table: Results of Method Variants

- ① Gather results for median integration method
- ② Distinguish individual sprouts
- ③ Sprout tracing similar to neuron tracing; see Meijering E. Neuron tracing in perspective. *Cytometry A* 2010;77A: 693704.
- ④ Three-dimensional reconstruction using multiple image depths

Thank you. Questions?