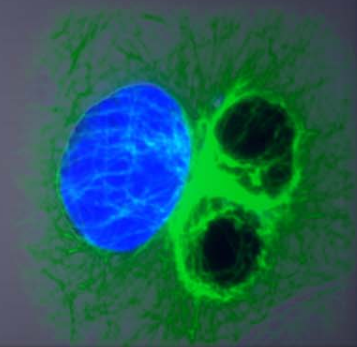


Analyzing COS1 Cells



for Studying Alzheimer's Disease

Vision Research Lab

Center for Bio-image Informatics

Ivan Villalba

Electrical Engineering

Cal Poly SLO

Oxnard College (2005)

Mentors

Jiyun Byun

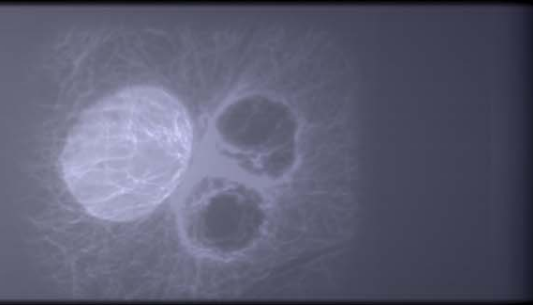
DeeAnn Hartung

Faculty Advisor

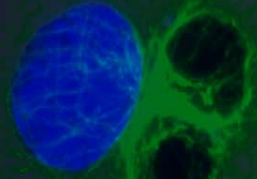
Dr. B.S. Manjunath



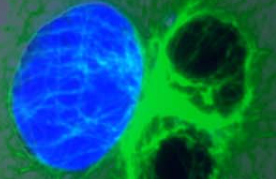
Overview



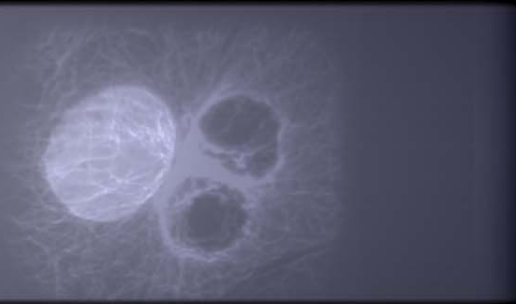
- Introduction
 - Alzheimer's
- Image Analysis
 - Properties
 - Procedure
- Experimental Results
 - Cell Segmentation
 - Cell Density
- Future Work



Introduction



Alzheimer's



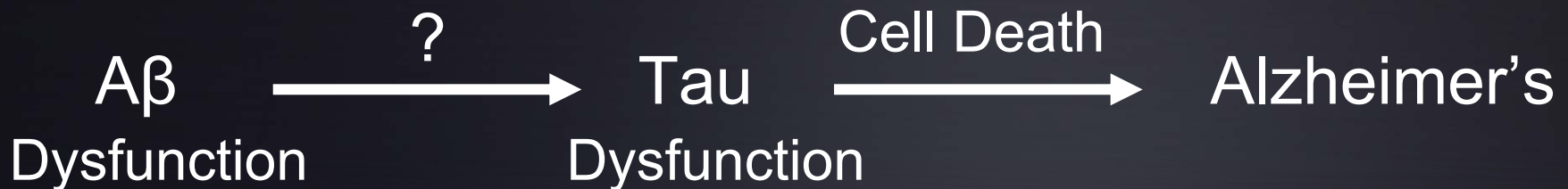
Characterized by two pathological hallmarks in the brain,

1. Extracellular Amyloid Plaques

➤ Amyloid-Beta ($A\beta$) protein

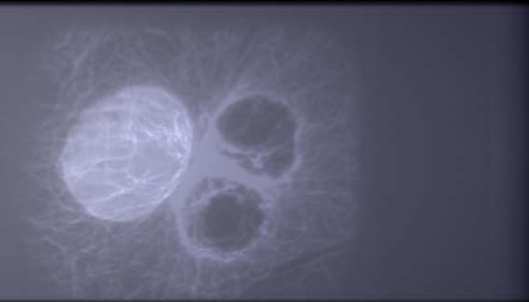
2. Intracellular Neurofibrillary Tangles (NFTs)

➤ Tau protein



- Big Picture: $A\beta$ and Tau

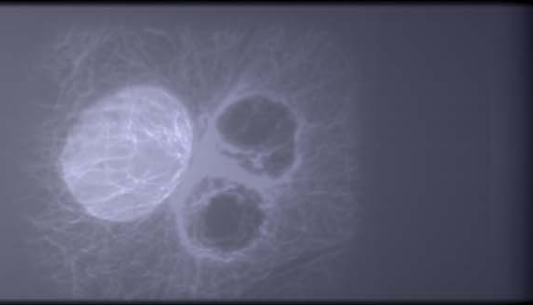
- Examining molecular mechanism of cell death.



Why COS1 Cells?

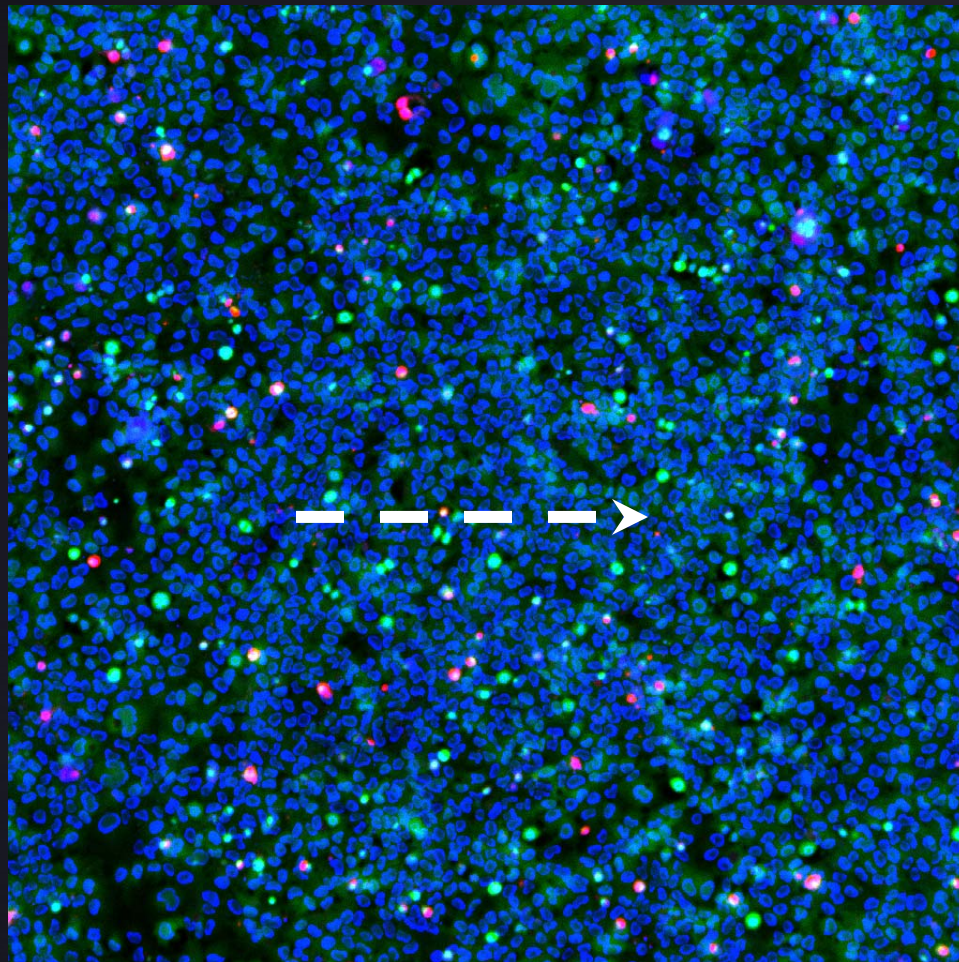
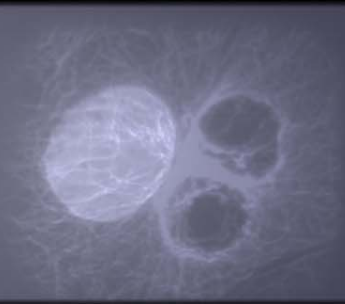
- Model System
 - Monkey Kidney Cells
 - Cancer cell line
 - COS1 Cells transfected with Tau
- Determine cell survival/death under various treatments and time points for Alzheimer's disease.

My Goals



- Cell survival/death ratio
 - Cell Count by segmentation
 - Live cells = total cells – dead cells
 - Cell shape: extract further information (e.g. Cell division ratio)

Motivation



Various ²⁷⁰4 treatments

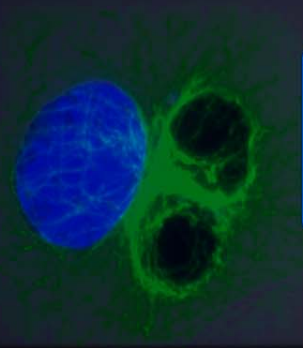
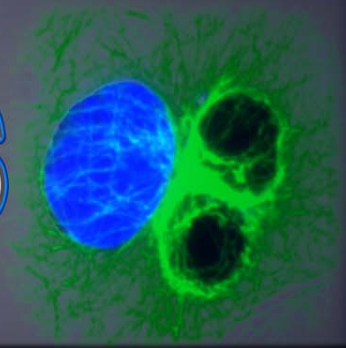
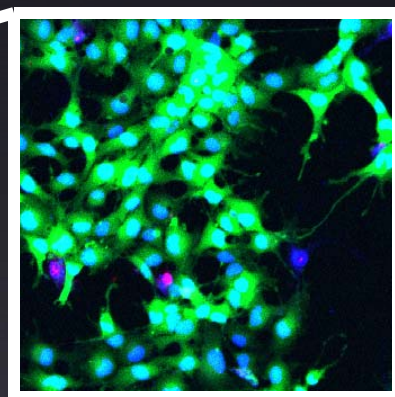
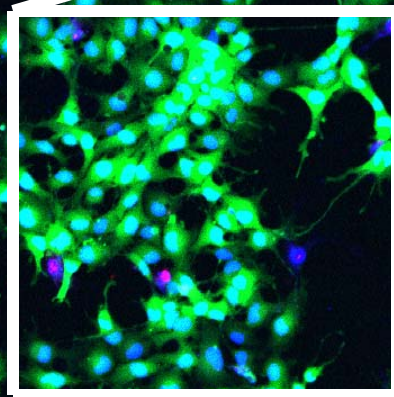
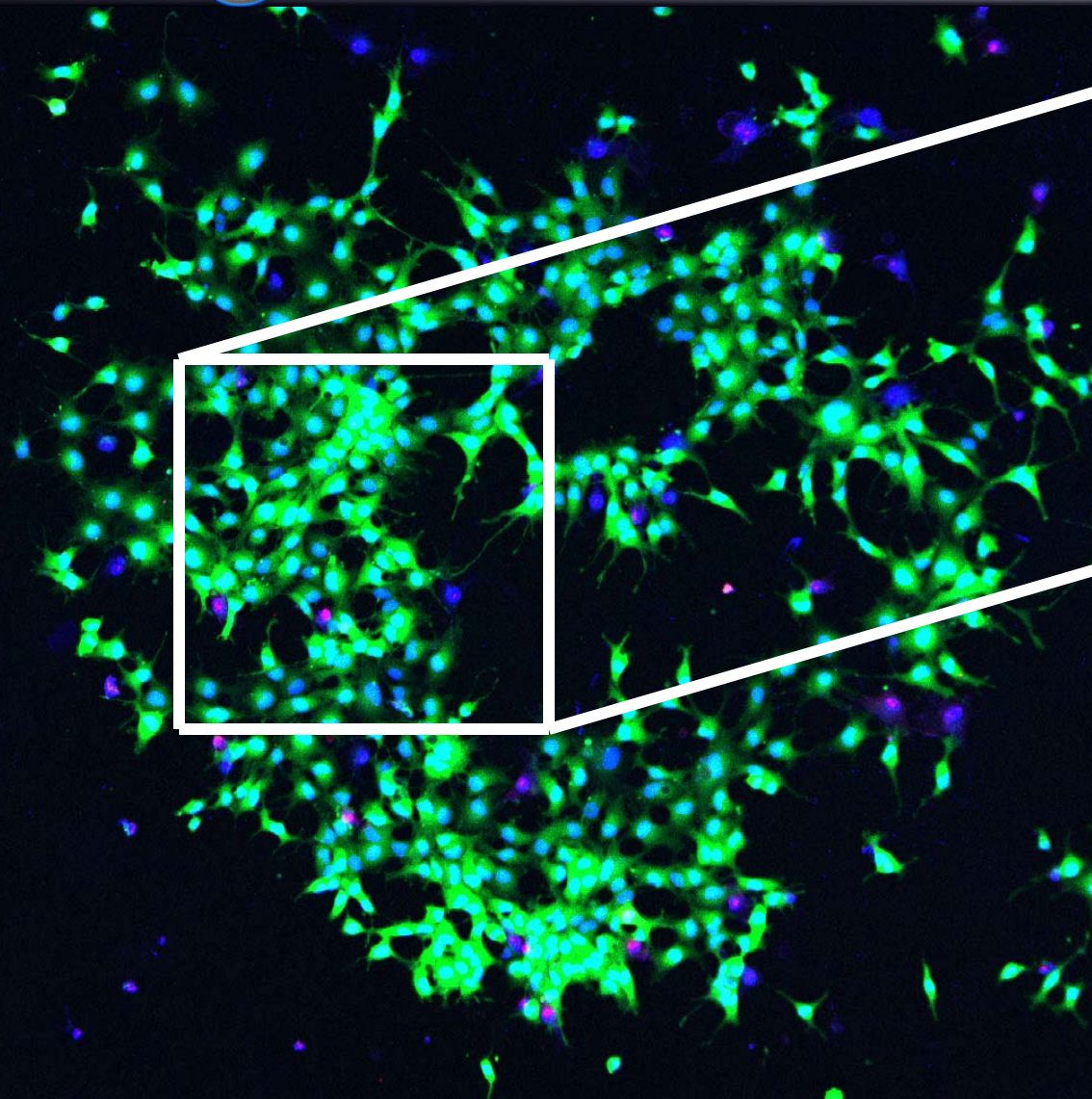
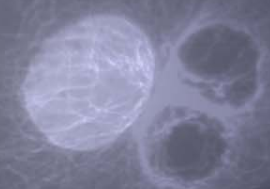


Image Analysis

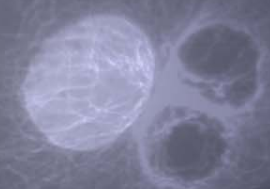


Image



10x
Confocal microscope
1024x1024 pix
1 pix = 0.1243 μ m

Image



 Hoechst

– Total Cells

 Propidium Iodide

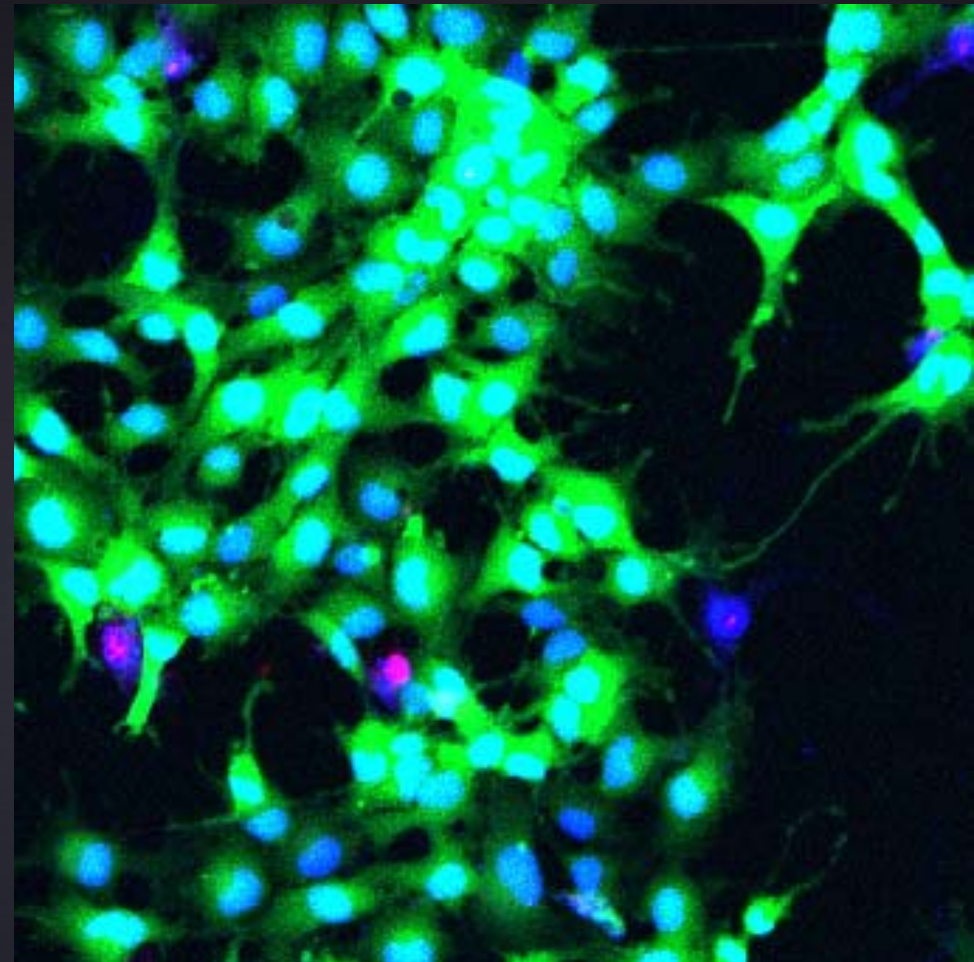
– Dead Cells

(Enters leaky membranes)

 Calcein AM

– Live Cells

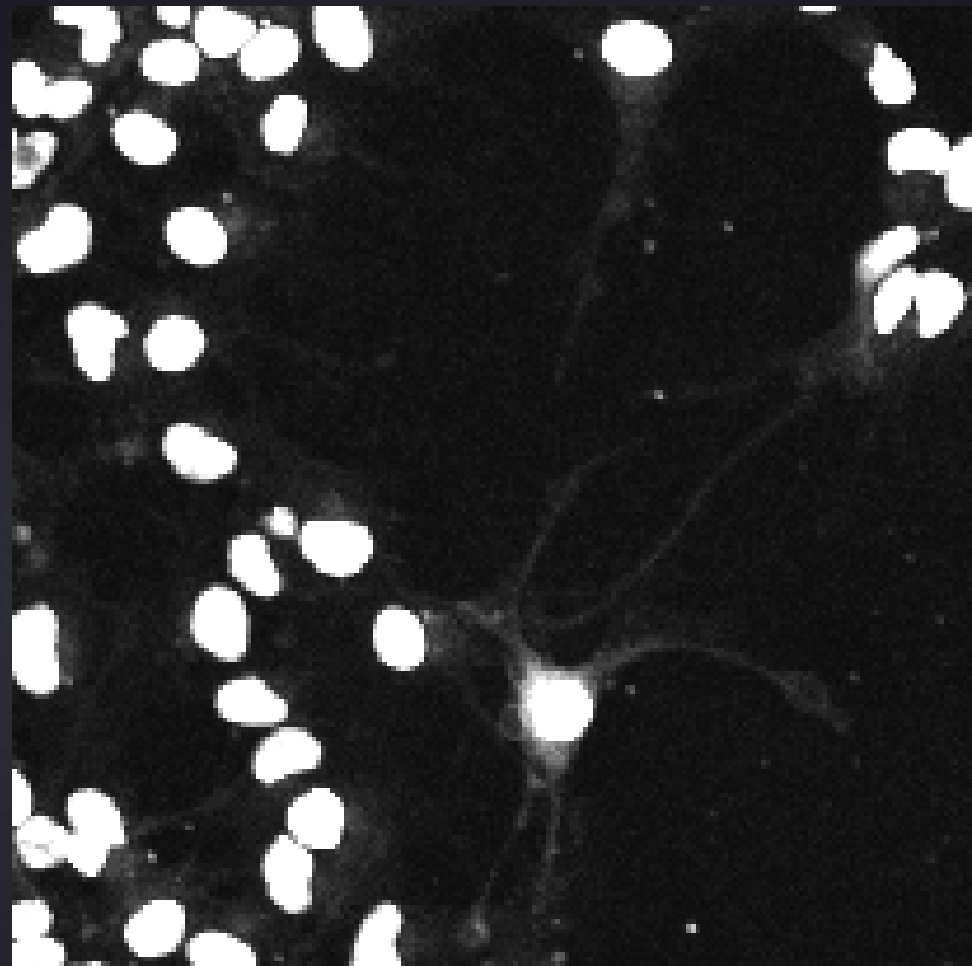
(Hydrolyzed by intracellular enzymes)



Hoechst Channel

Procedure

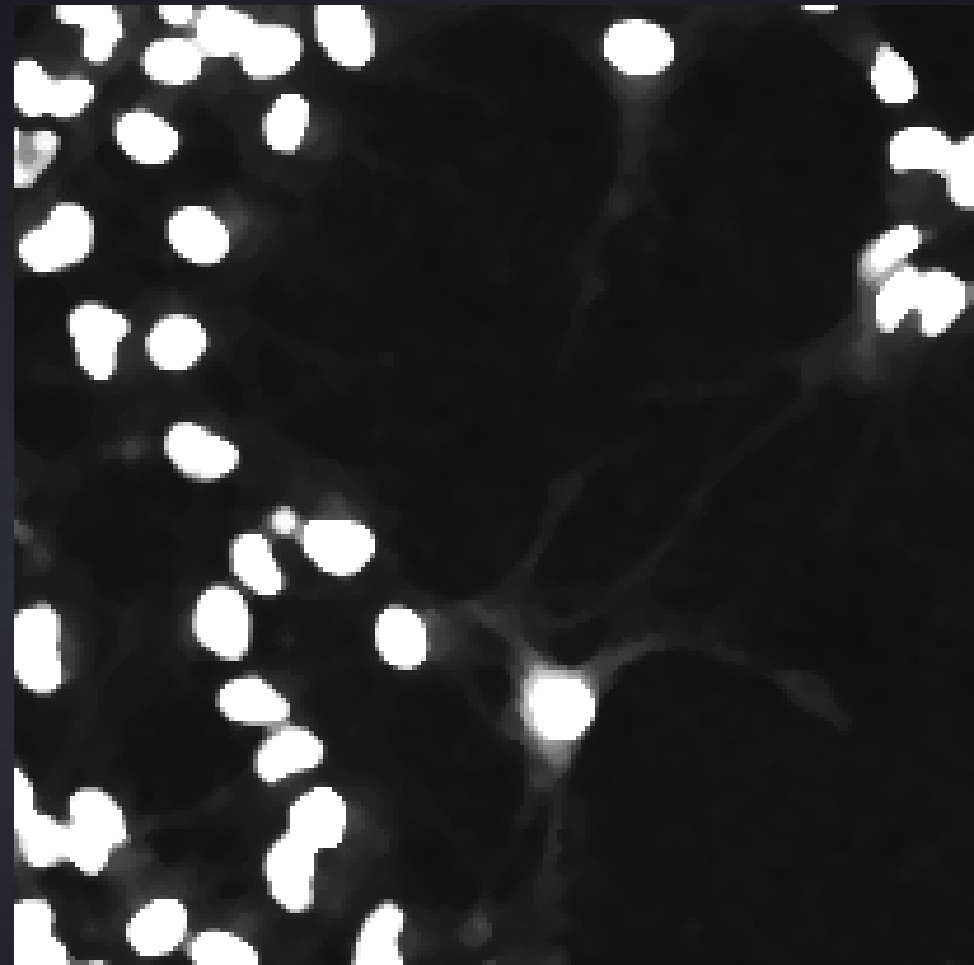
- Isolate Channels



Hoechst Channel

Procedure

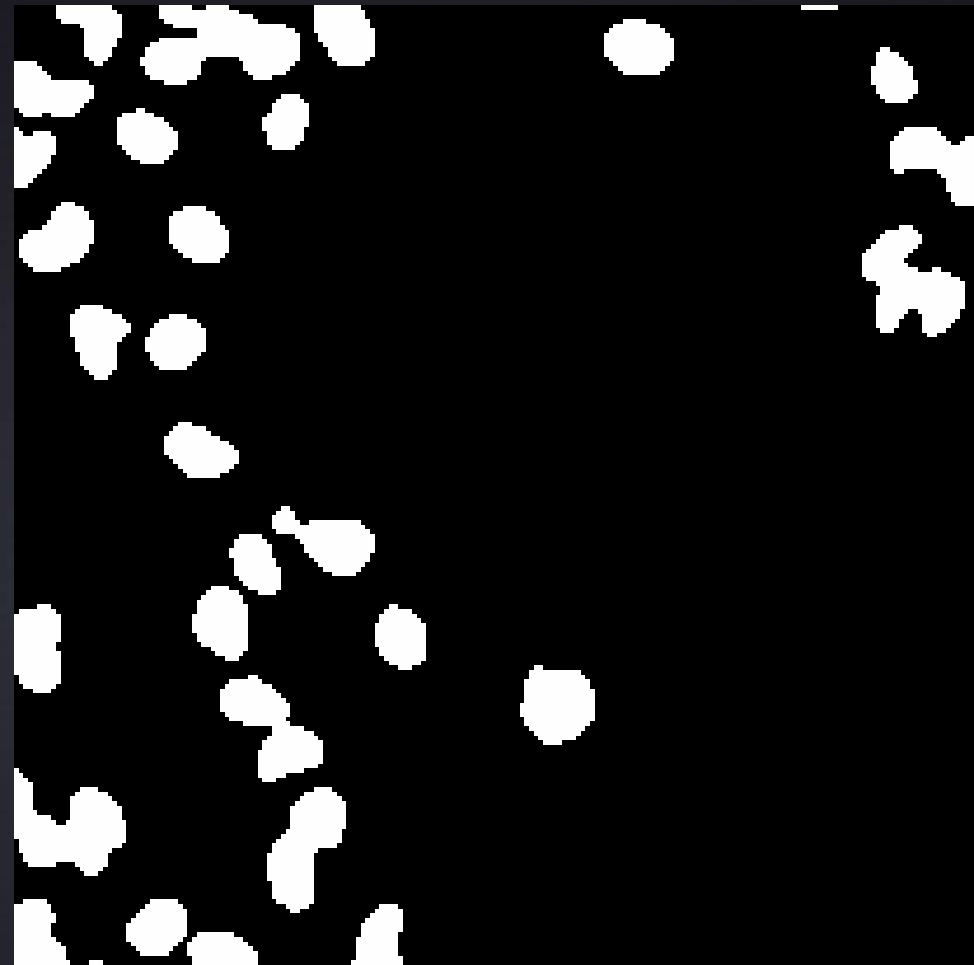
- Isolate Channels
- **Median Filtering**
 - **Noise Reduction**



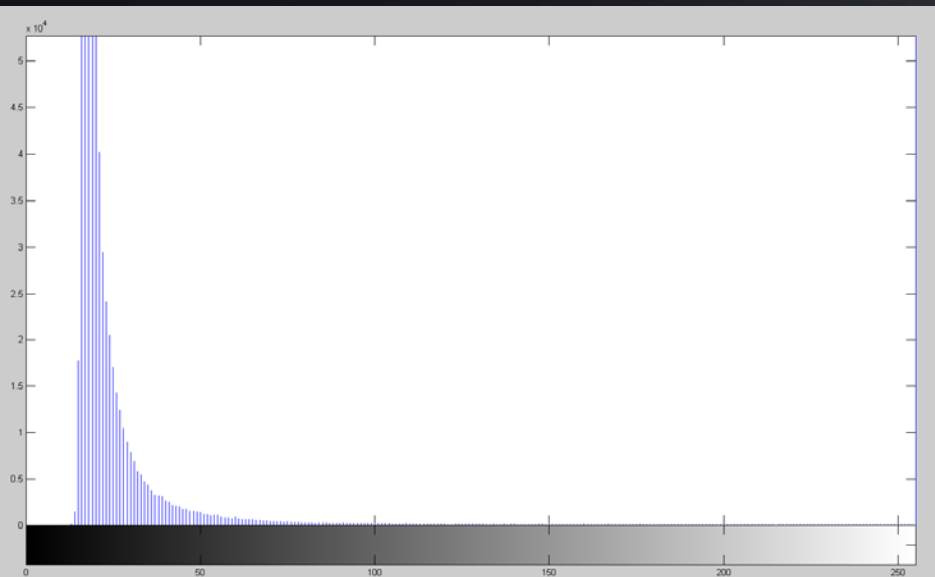
Hoechst Channel

Procedure

- Isolate Channels
- Median Filtering
- **Binary Image**
 - Threshold

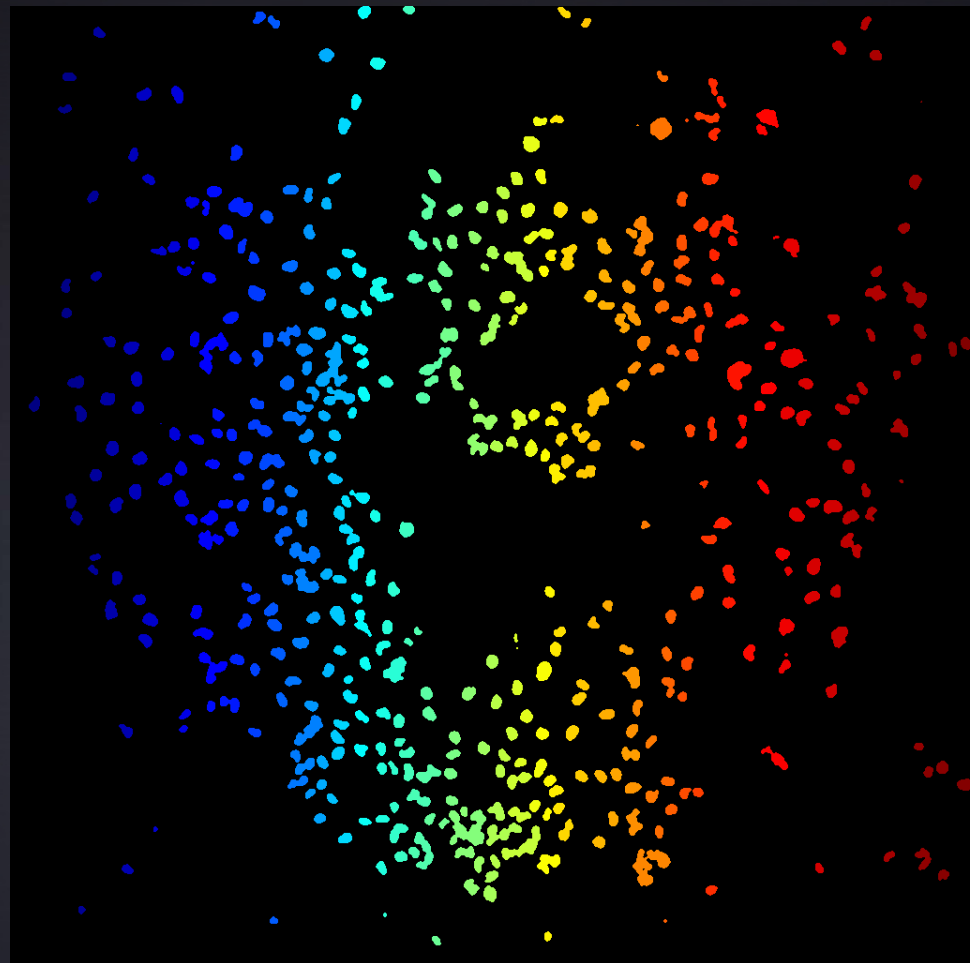


Hoechst Channel

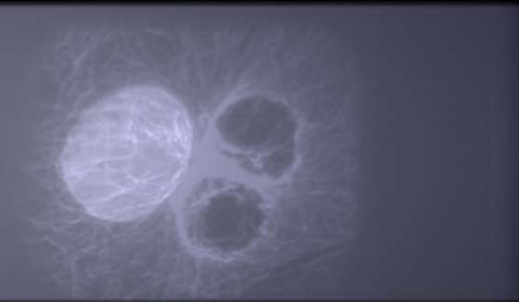


Procedure

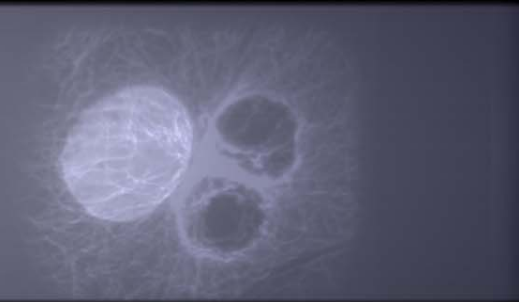
- Isolate Channels
- Median Filtering
- Binary Image
- **Label Image**



Hoechst Channel

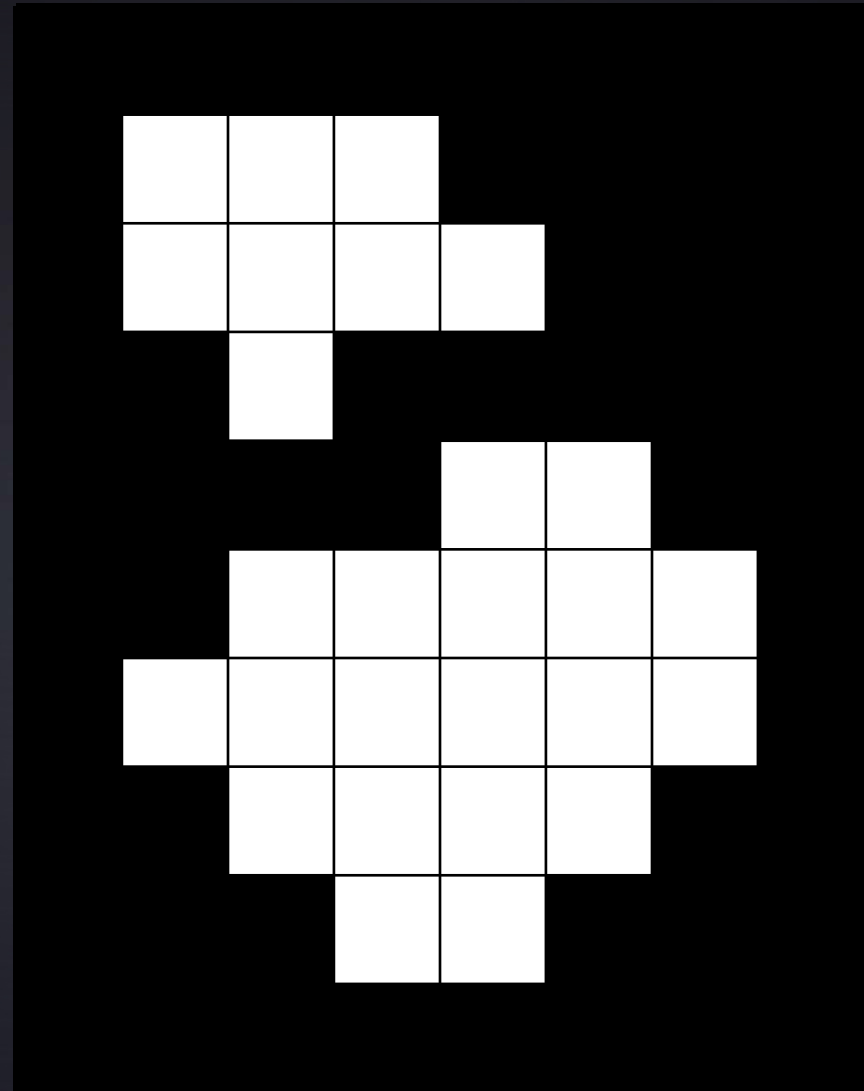


Procedure



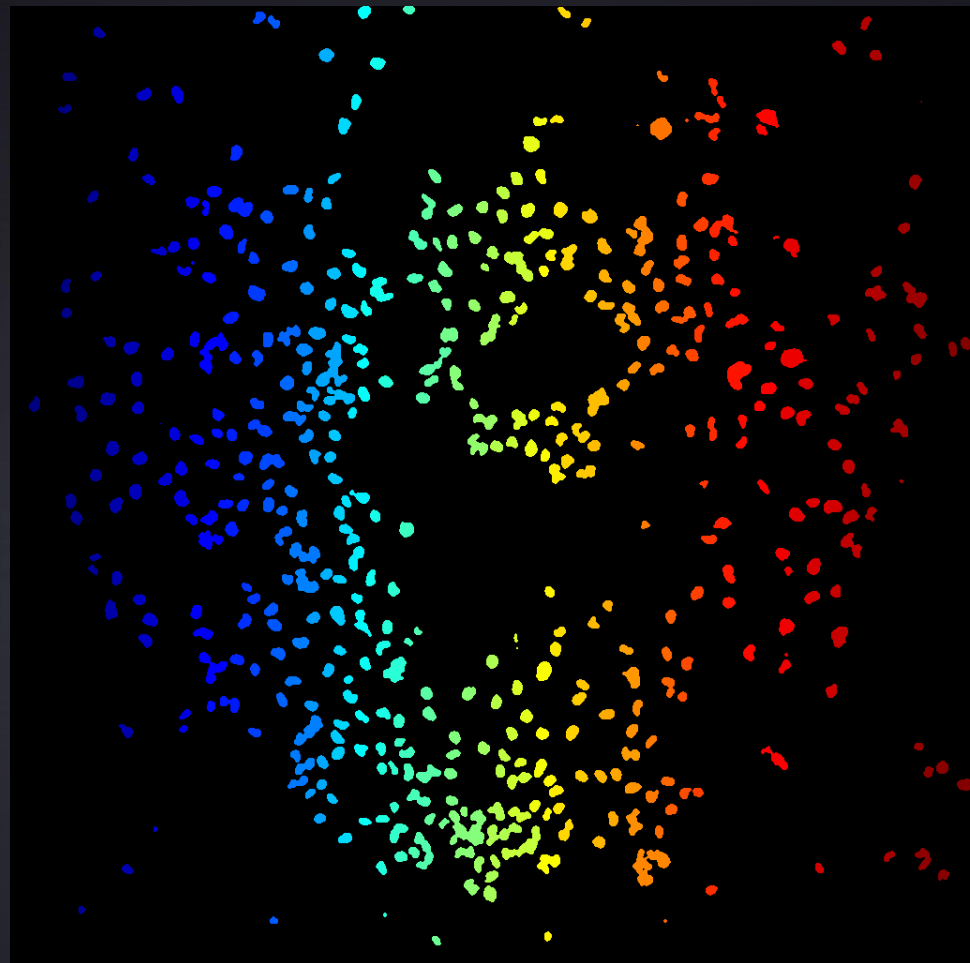
- Isolate Channels
- Median Filtering
- Binary Image
- **Label Image**

Example:
10x8 pixel
Binary image

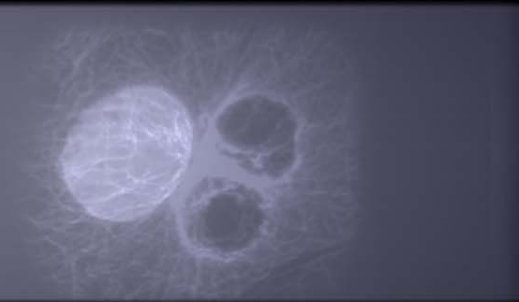


Procedure

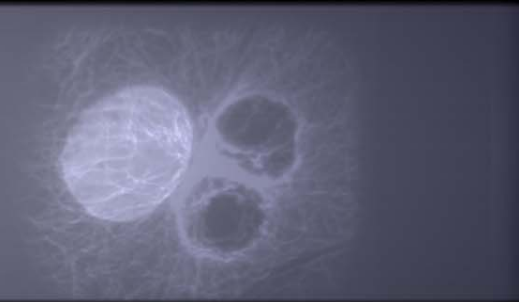
- Isolate Channels
- Median Filtering
- Binary Image
- **Label Image**



Hoechst Channel



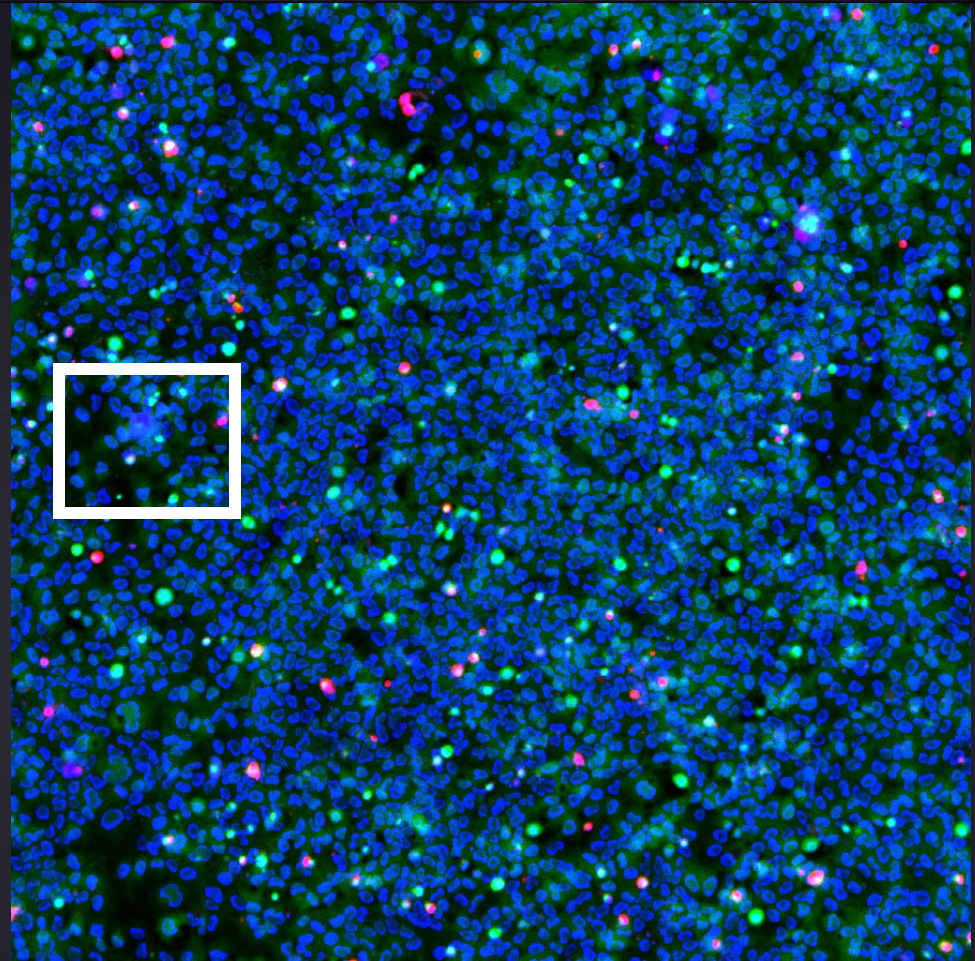
Procedure



- Isolate Channels
- Median Filtering
- Binary Image
- Label Image
- **Object Analysis**

Procedure

- Isolate Channels
- Median Filtering
- Binary Image
- Label Image
- **Object Analysis**

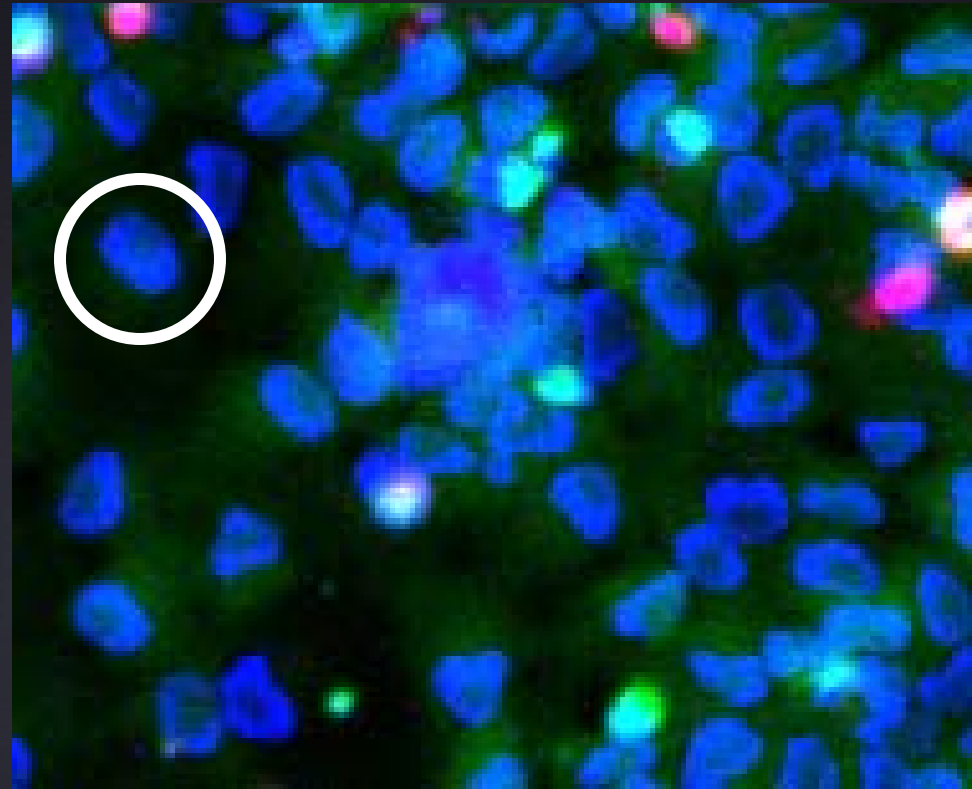


120 Hours Untreated



Procedure

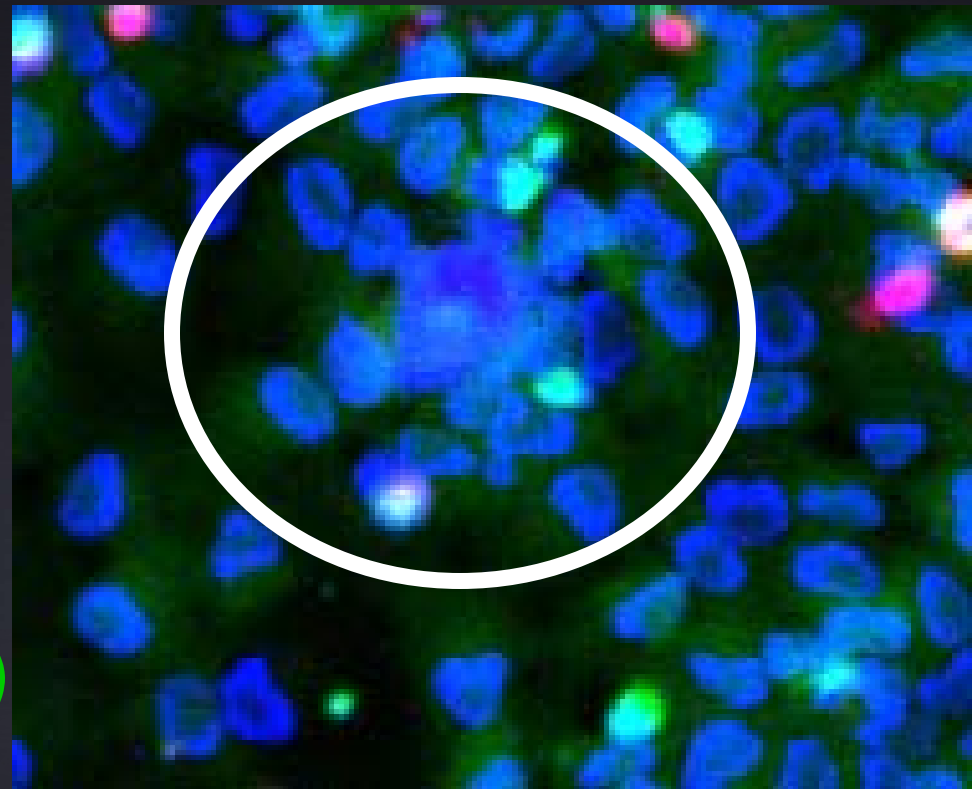
- Isolate Channels
- Median Filtering
- Binary Image
- Label Image
- **Object Analysis**
 - Normal (1 cell)



120 Hours Untreated

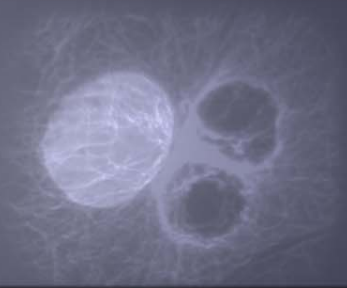
Procedure

- Isolate Channels
- Median Filtering
- Binary Image
- Label Image
- **Object Analysis**
 - Cluster (≥ 3 cells)



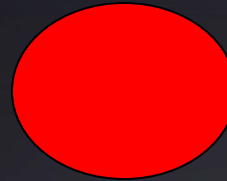
120 Hours Untreated

Procedure



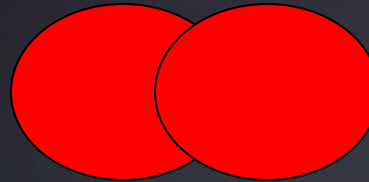
- **Object Analysis**

- Normal

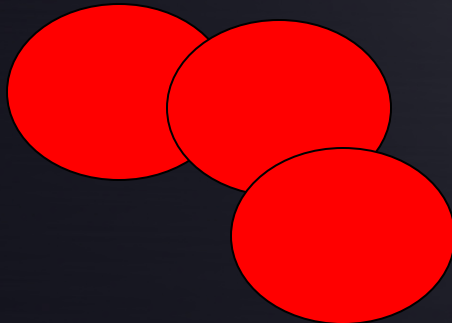


- Peanut

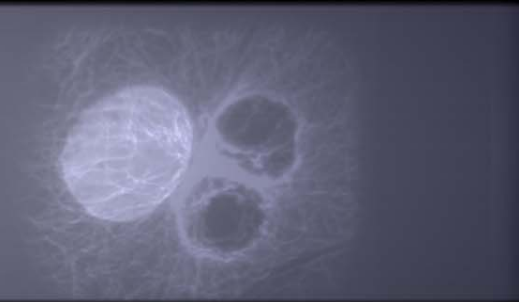
- (Dividing cell)



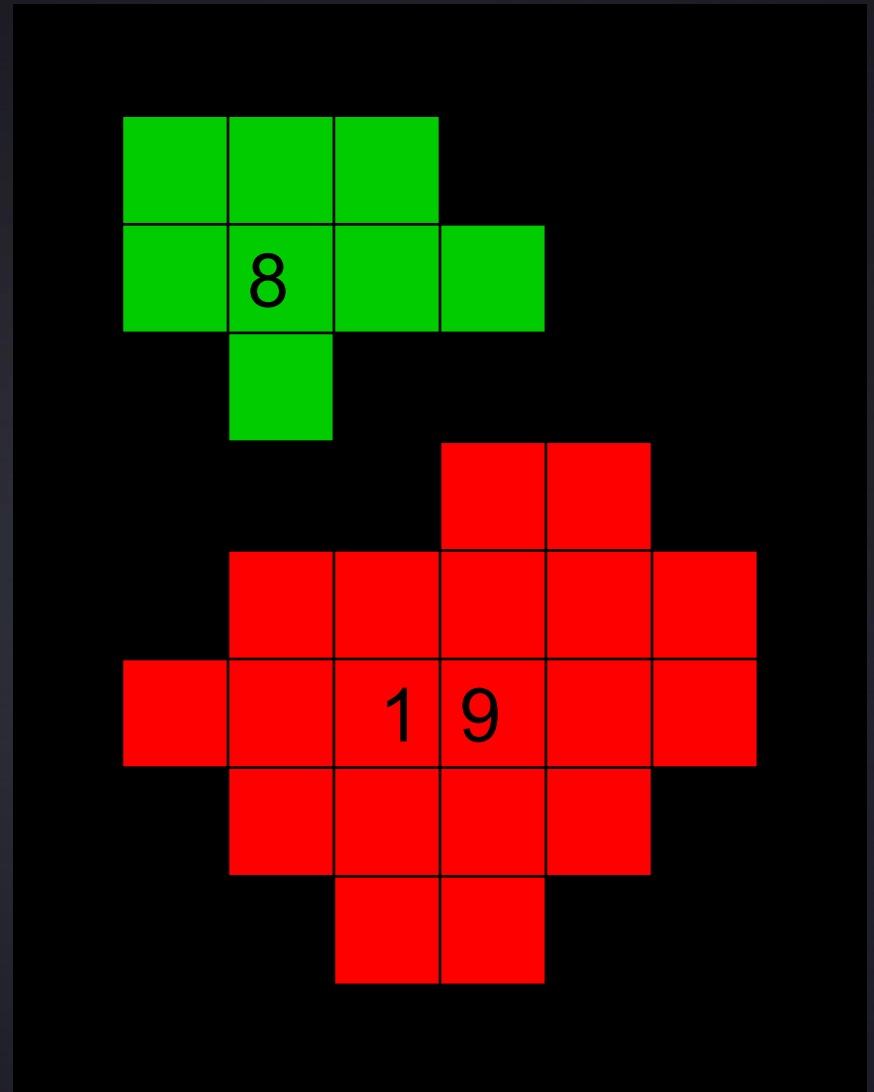
- Cluster (≥ 3 cells)



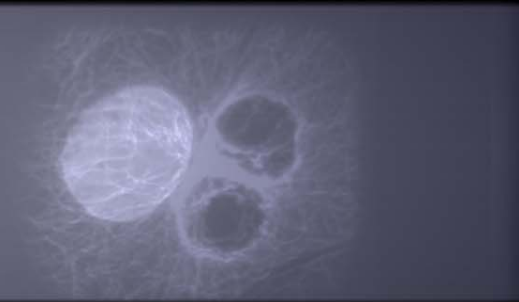
Statistics



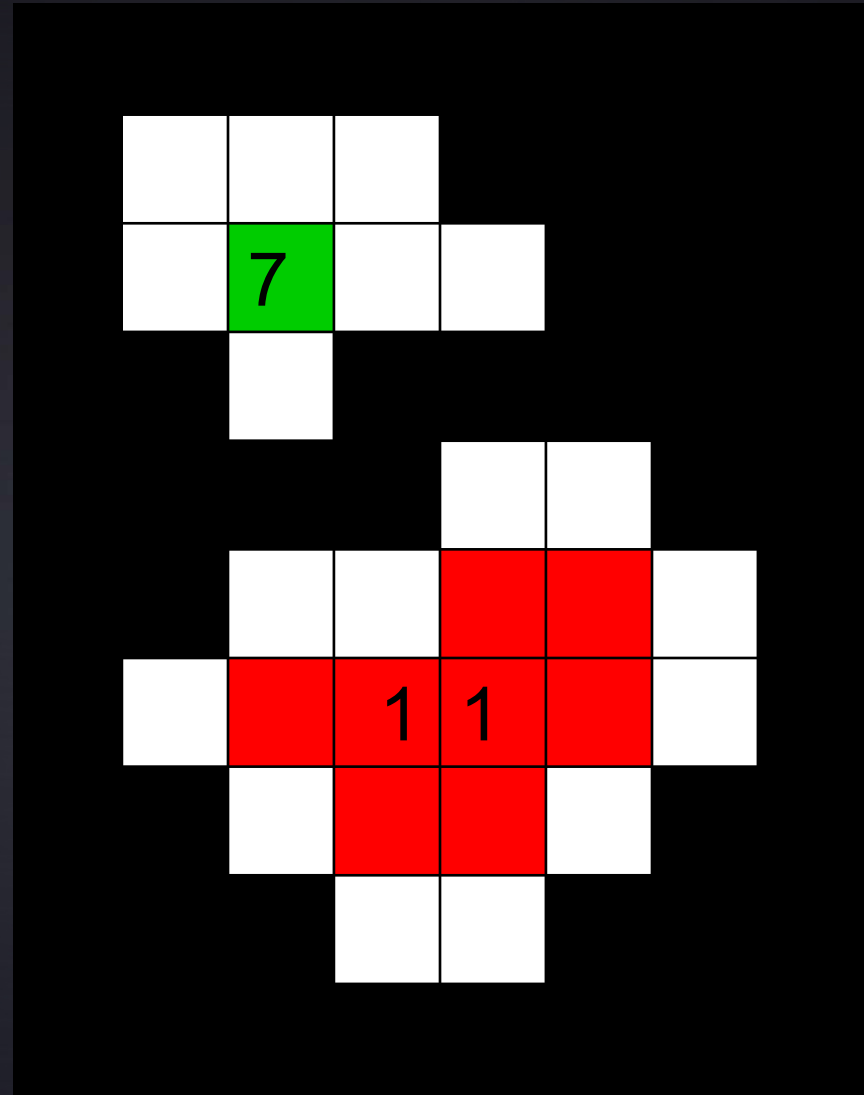
- Area



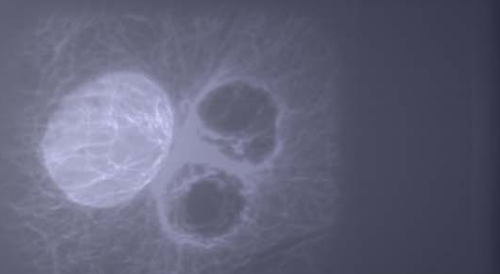
Statistics



- Area
- **Perimeter**

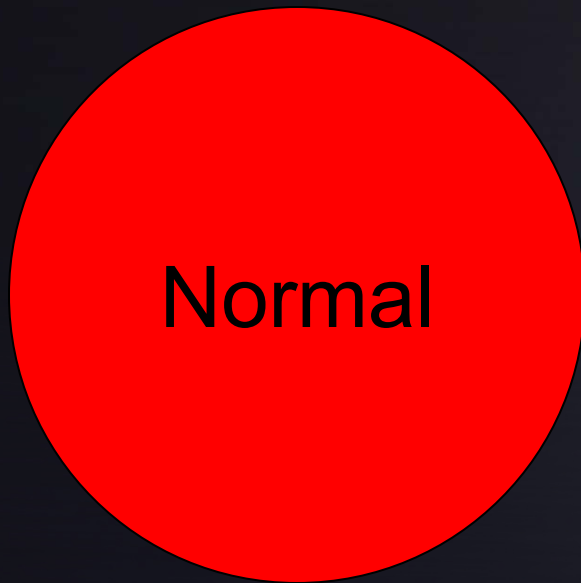


Statistics

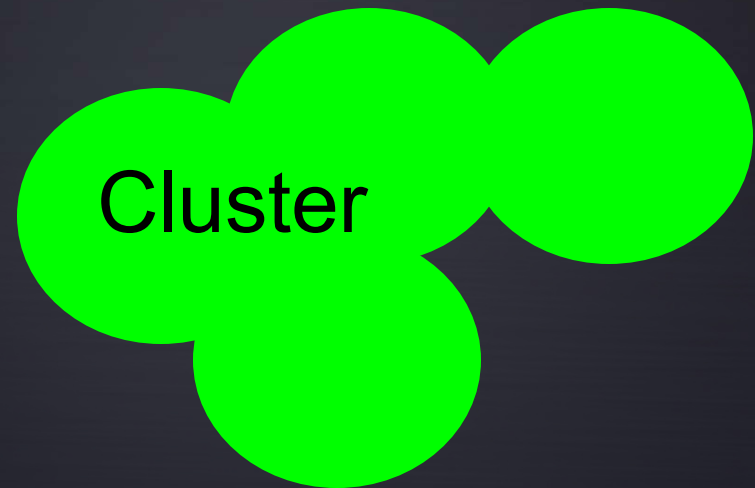


- Area
- Perimeter
- **Compactness**

$$\text{Compactness} = \frac{4 \times \pi \times \text{Area}}{\text{Perimeter}^2}$$

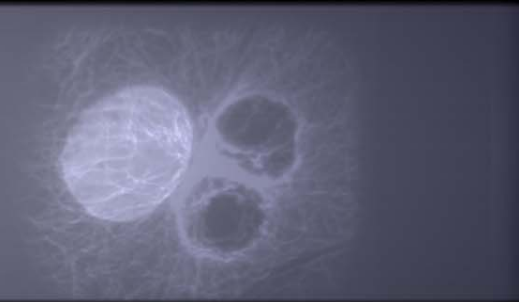


Compactness = 1

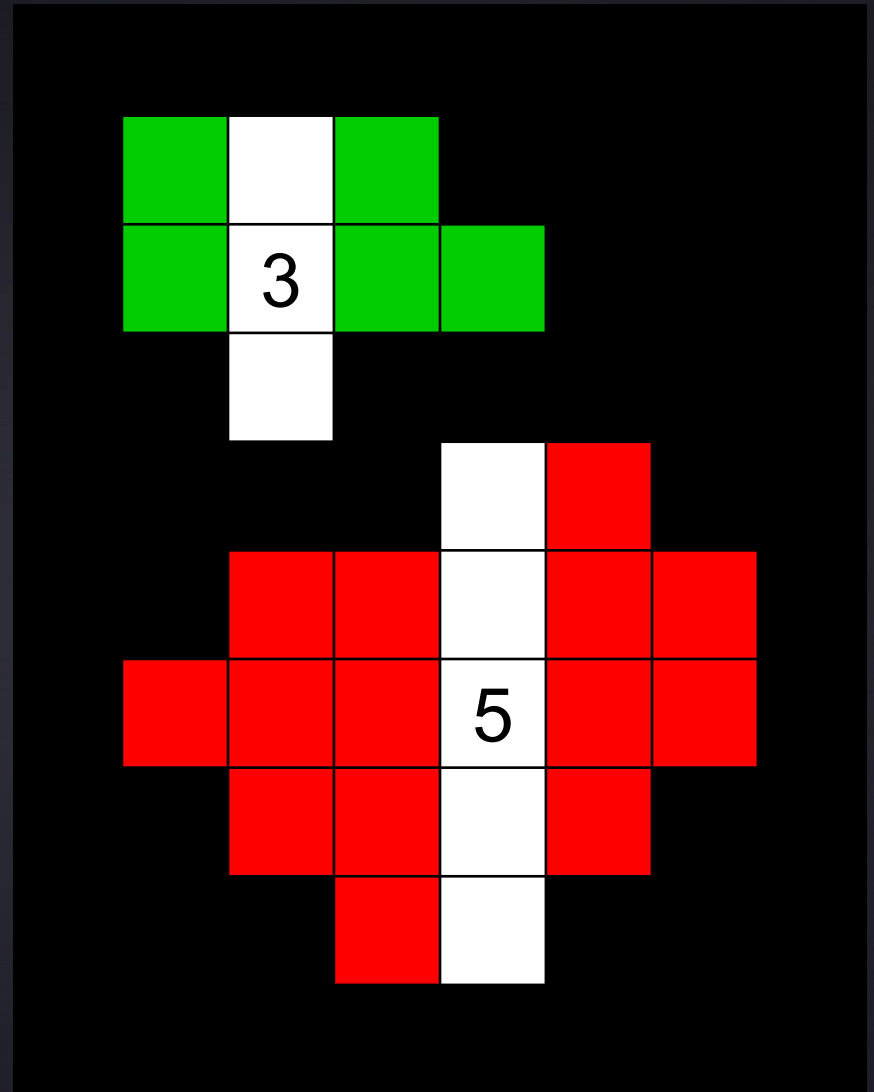


Low Compactness

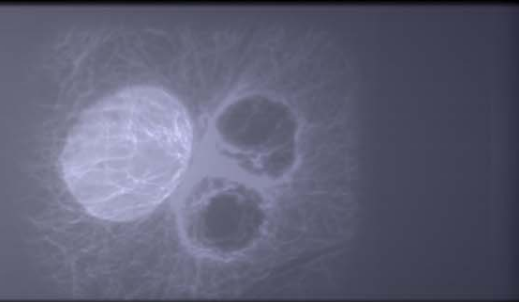
Statistics



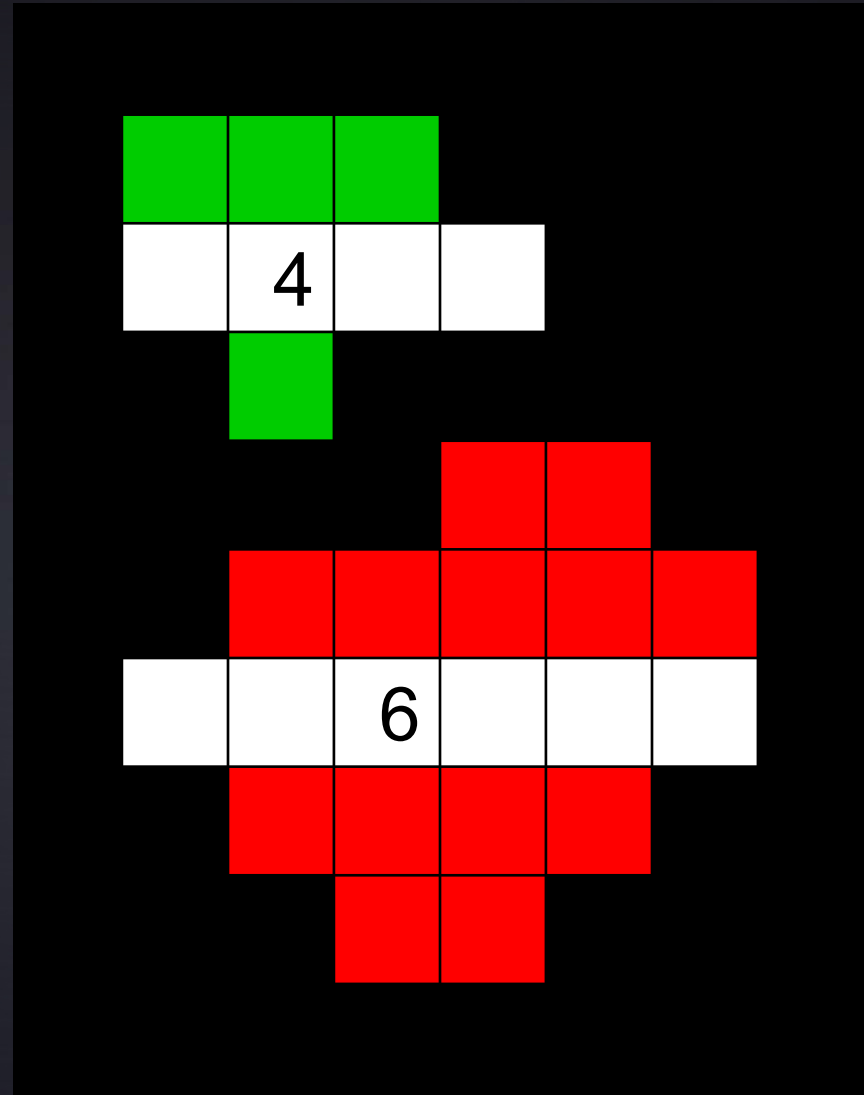
- Area
- Perimeter
- Compactness
- **Minor Axis**



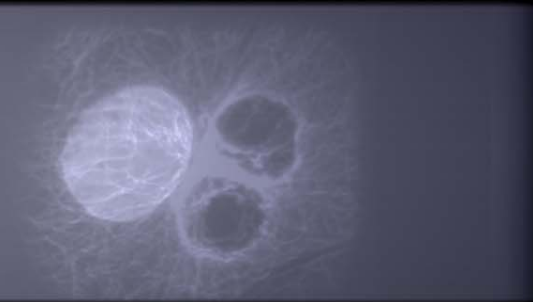
Statistics



- Area
- Perimeter
- Compactness
- Minor Axis
- **Major Axis**



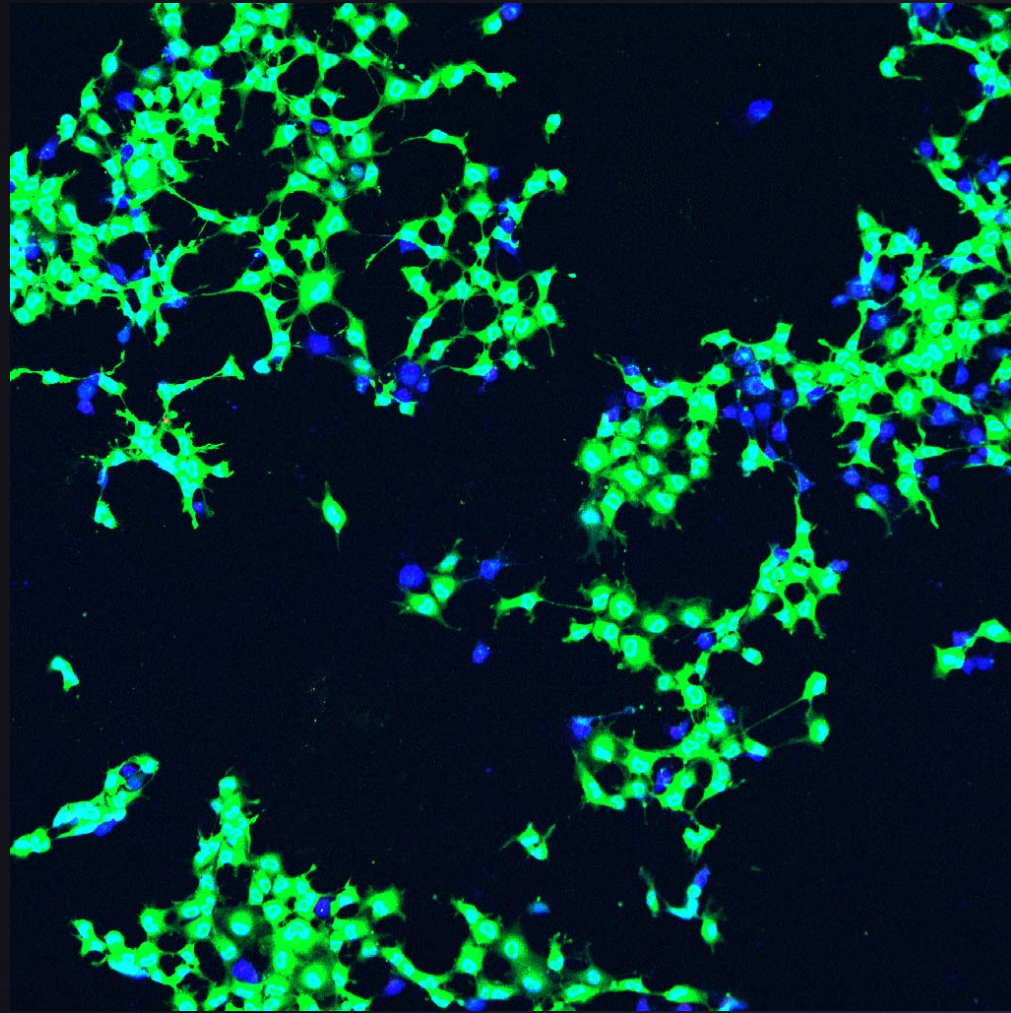
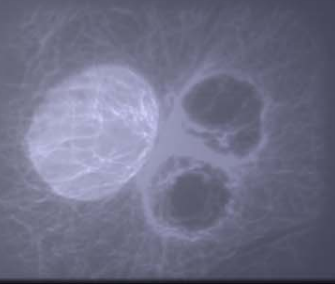
Statistics



- Area
- Perimeter
- Compactness
- Minor Axis
- Major Axis
- **Axis Ratio**

$$\text{Axis Ratio} = \frac{\text{Minor Axis}}{\text{Major Axis}}$$

Procedure

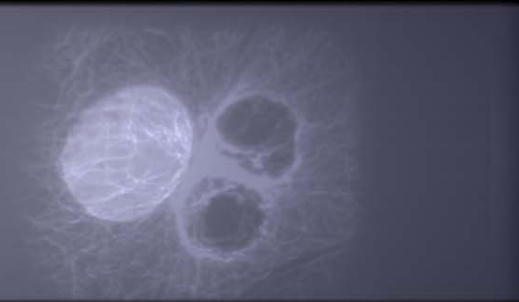


2hrs Untreated

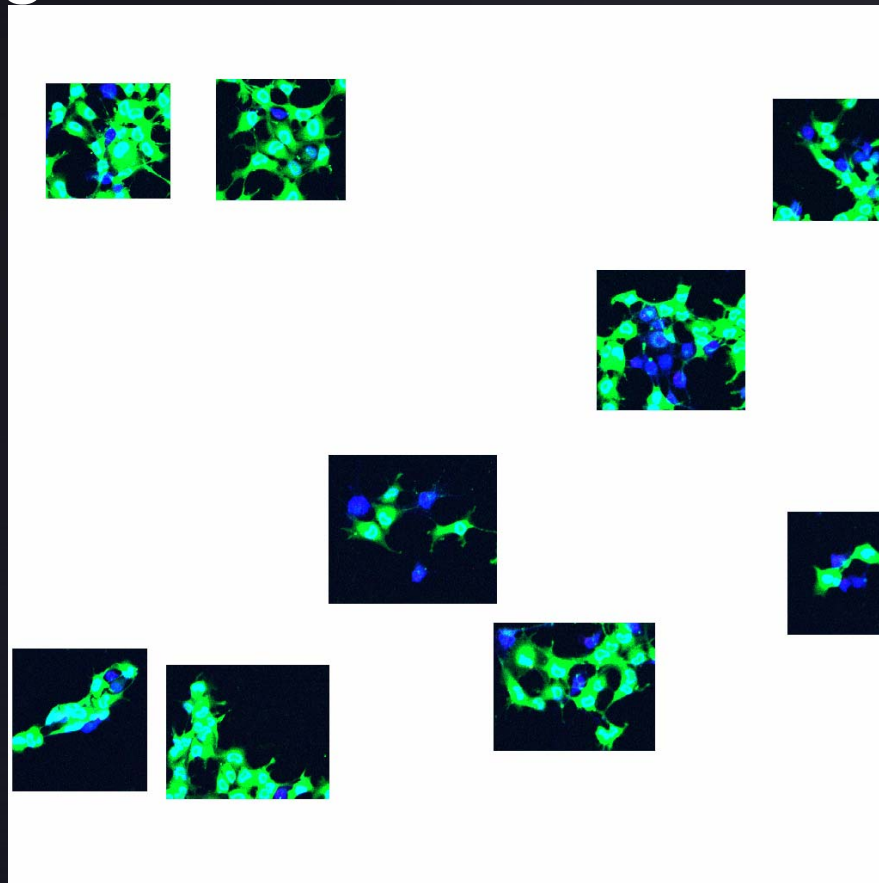
Class Type:

1. Normal + Peanut
2. Cluster

Procedure

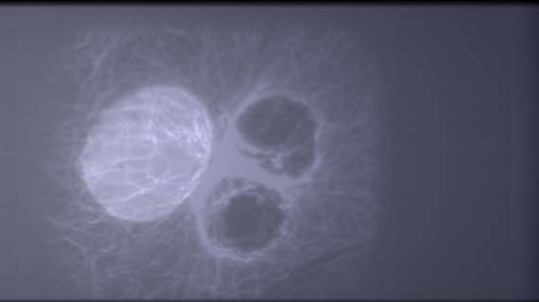


- Crop Regions

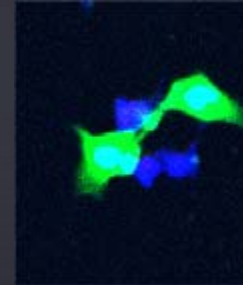
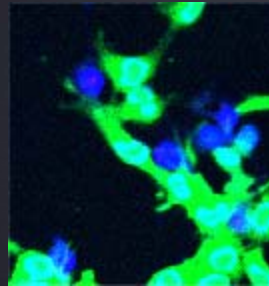
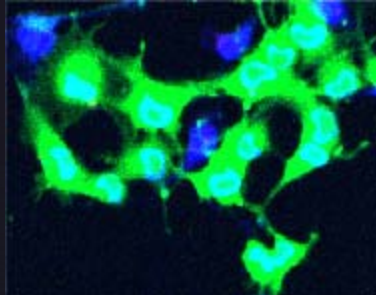
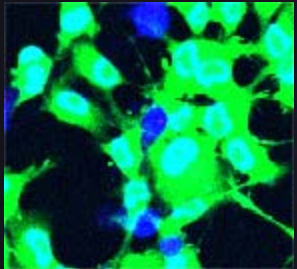
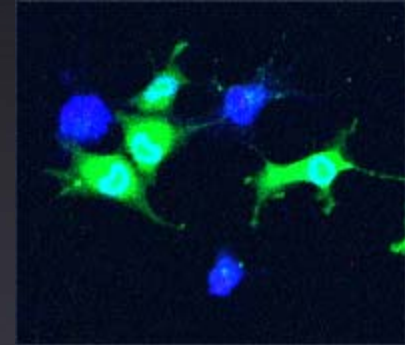
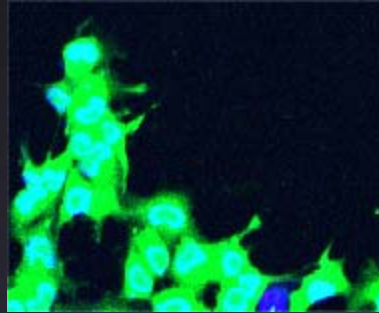
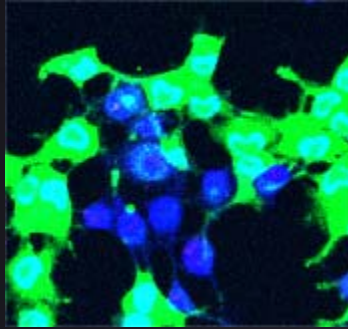
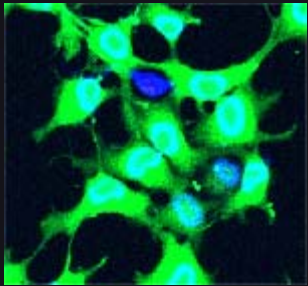


2hrs Untreated

Procedure

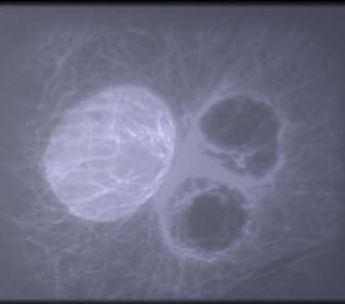


- Crop Images



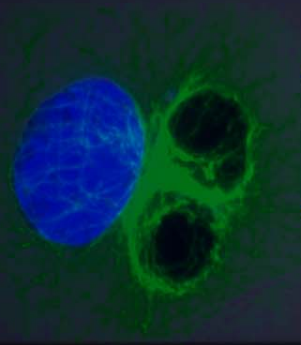
2hrs Untreated

Statistics

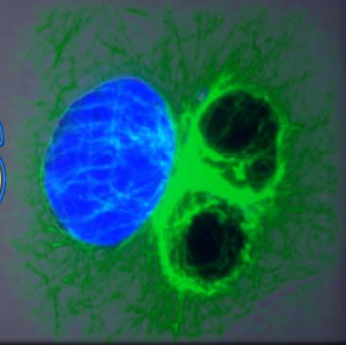


- Area
- Perimeter
- Compactness
- Minor Axis
- Major Axis
- Axis Ratio

- Clusters can be discriminated.
- Normal & Peanut are difficult to discriminate.

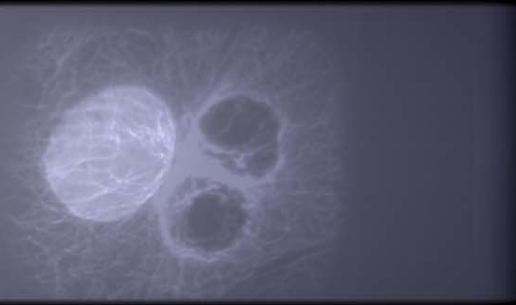


Experimental Results



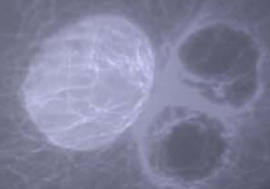
Dataset

Analyzed with MATLAB



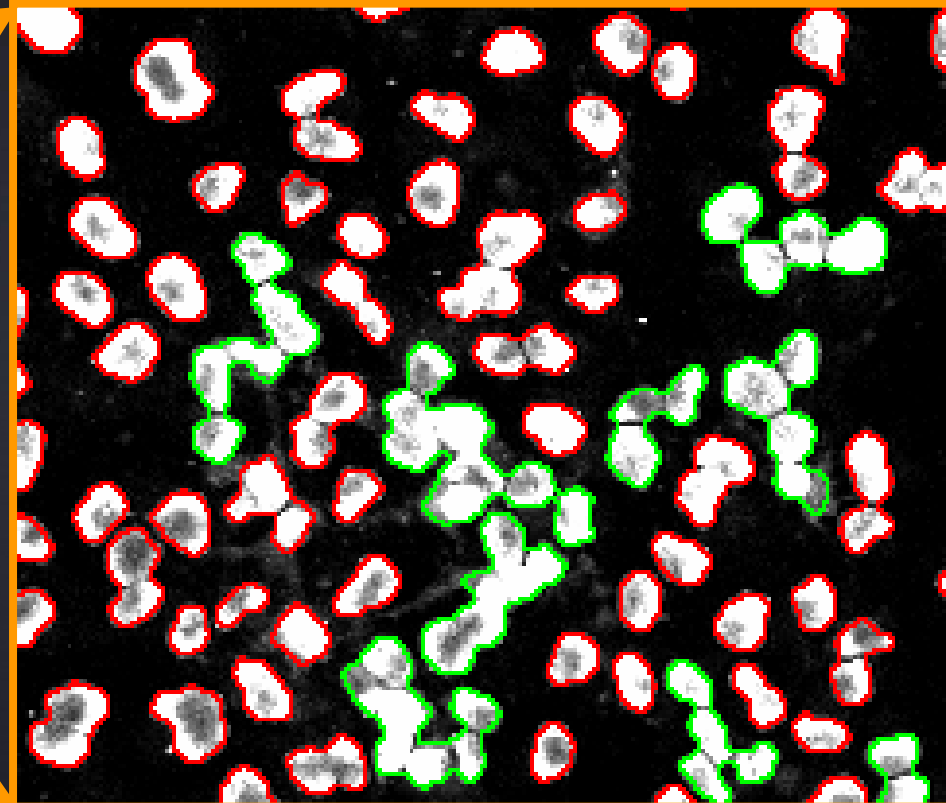
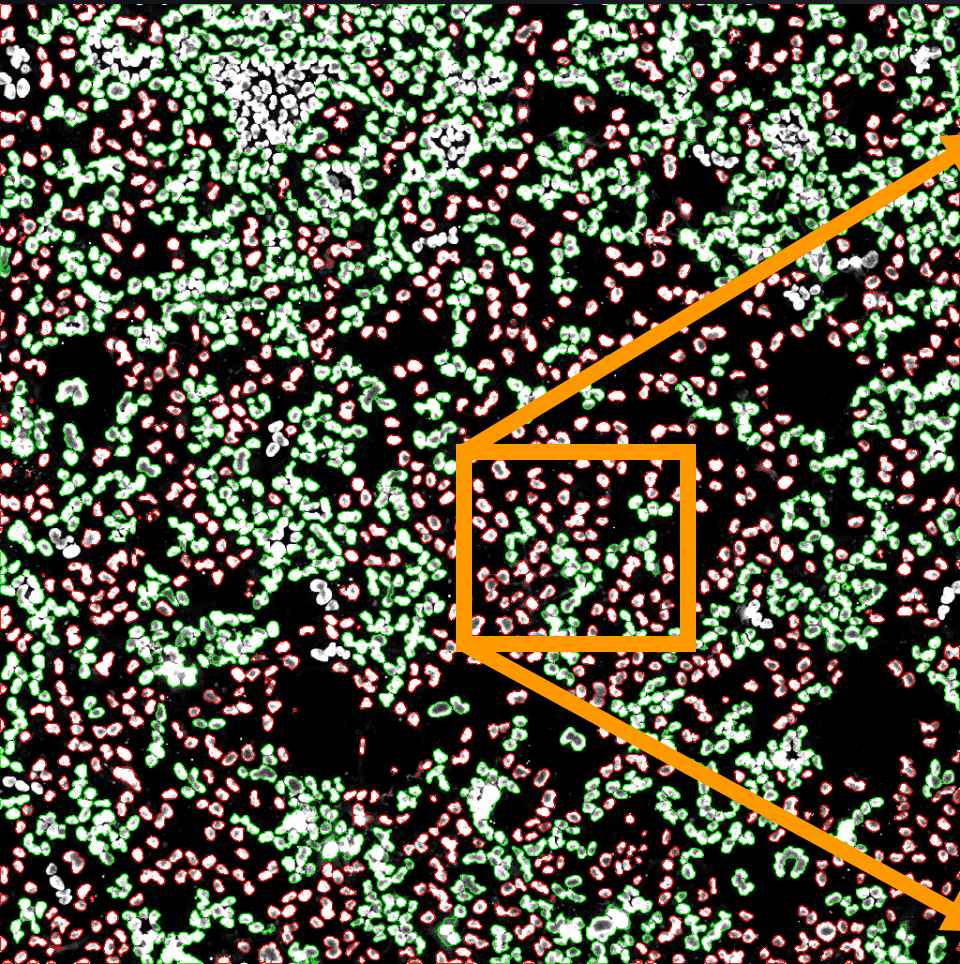
- 42 “Untreated” Images from 7 different time points
 - 2 hrs (7)
 - 6 hrs (5)
 - 12 hrs (6)
 - 24 hrs (5)
 - 48 hrs (8)
 - 72 hrs (6)
 - 120 hrs (5)

Experimental Results

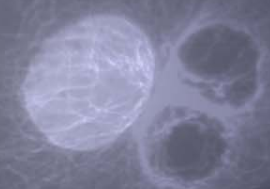


120 Hours

Red = Normal + Peanut
Green = Clusters

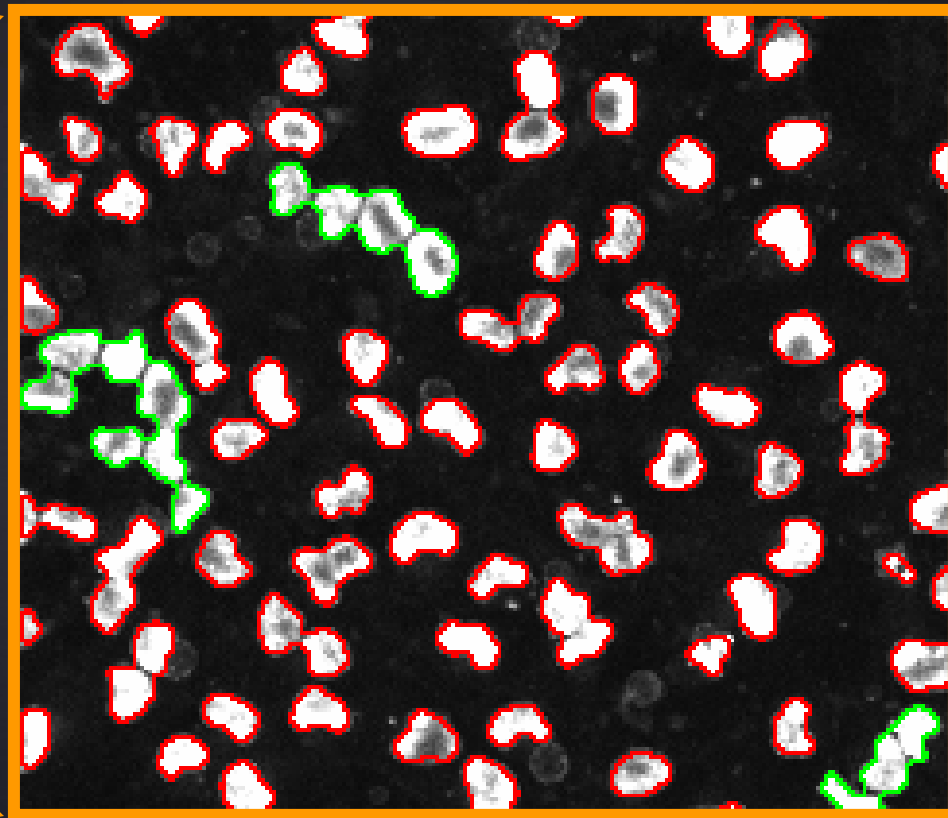
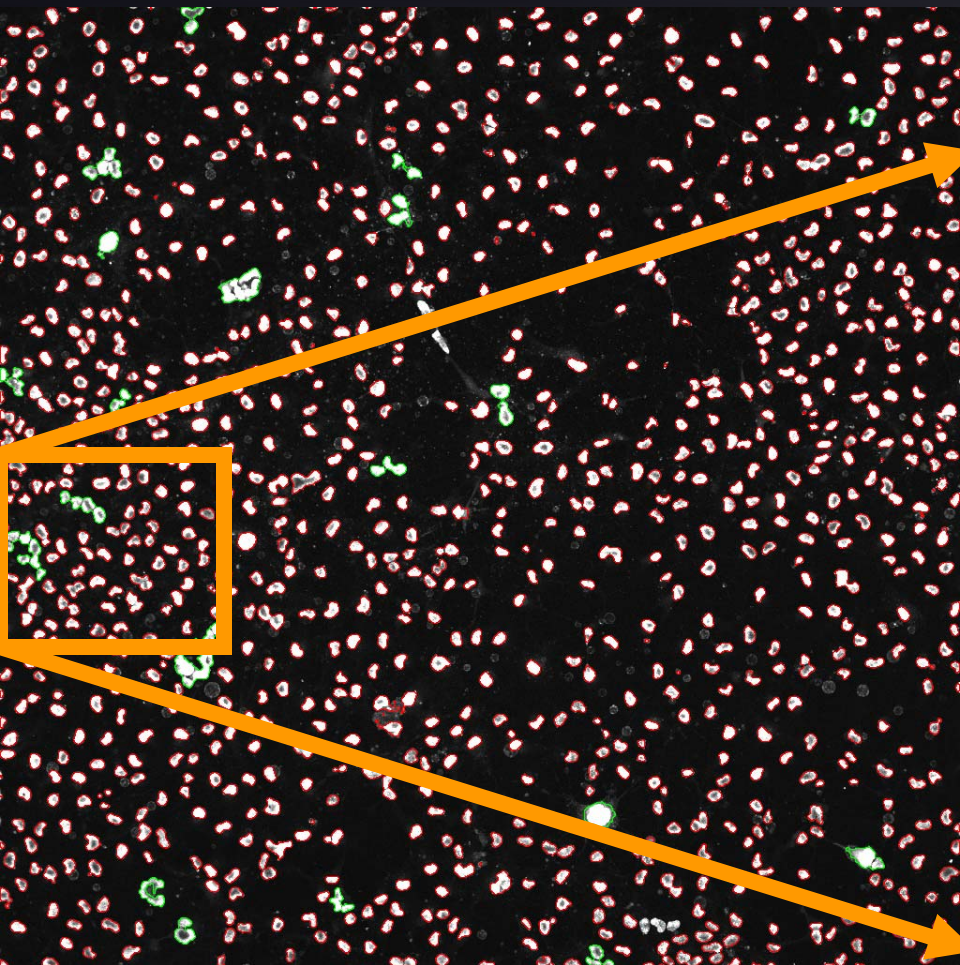


Experimental Results

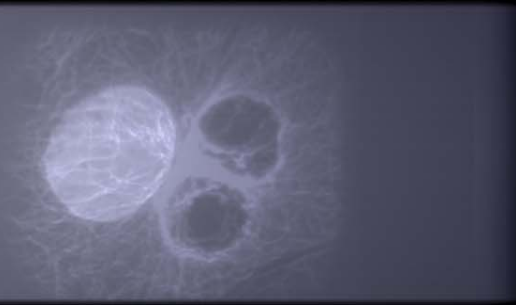


24 Hours

Red = Normal + Peanut
Green = Clusters



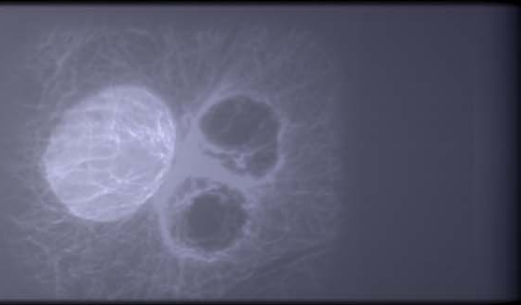
Cell Density



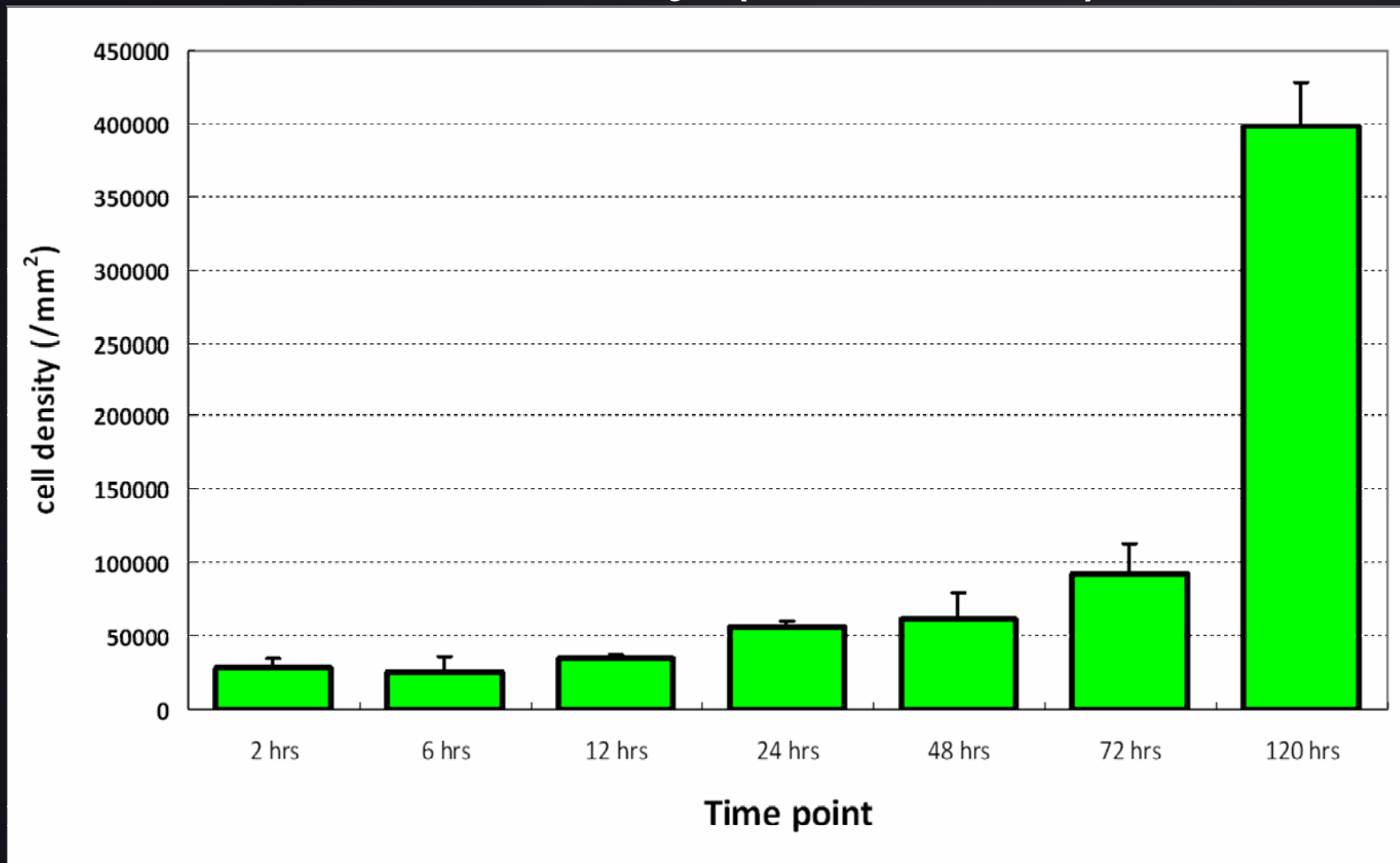
- Cell Density:
 - Cells = Number of “Normal & Peanut”
 - Area = Image Area – Cluster Area

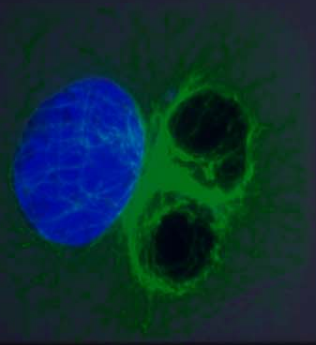
$$\text{Cell Density} = \frac{\text{Cells}}{\text{Area}}$$

Experimental Results

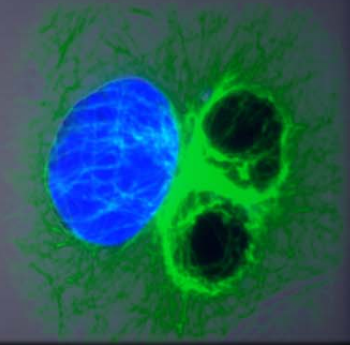


Cell Density (Untreated)

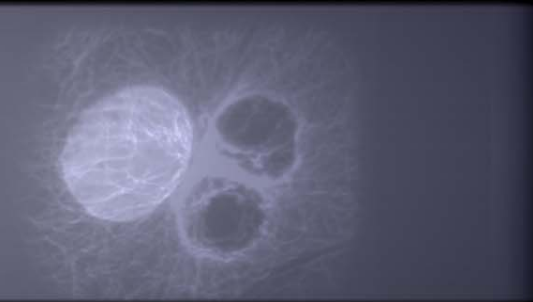




Future Work

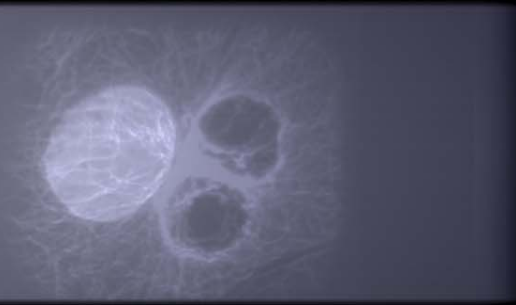


Future Work



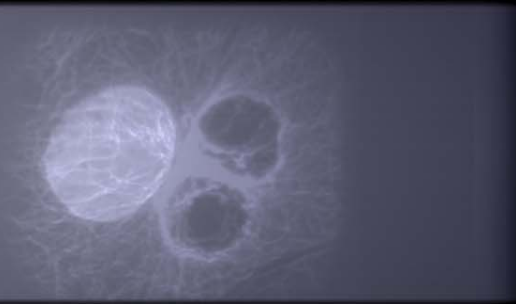
- Apply to various treatment
 - Untreated
 - 5 μM Staurosporine (stauro)
 - 10 μM Amyloid-Beta ()
 - 100 nM Taxol (Taxol)
 - 100 nM Taxol + 10 μM Amyloid-Beta (Tx-)
- Cell survival/death ratio.

Future Work

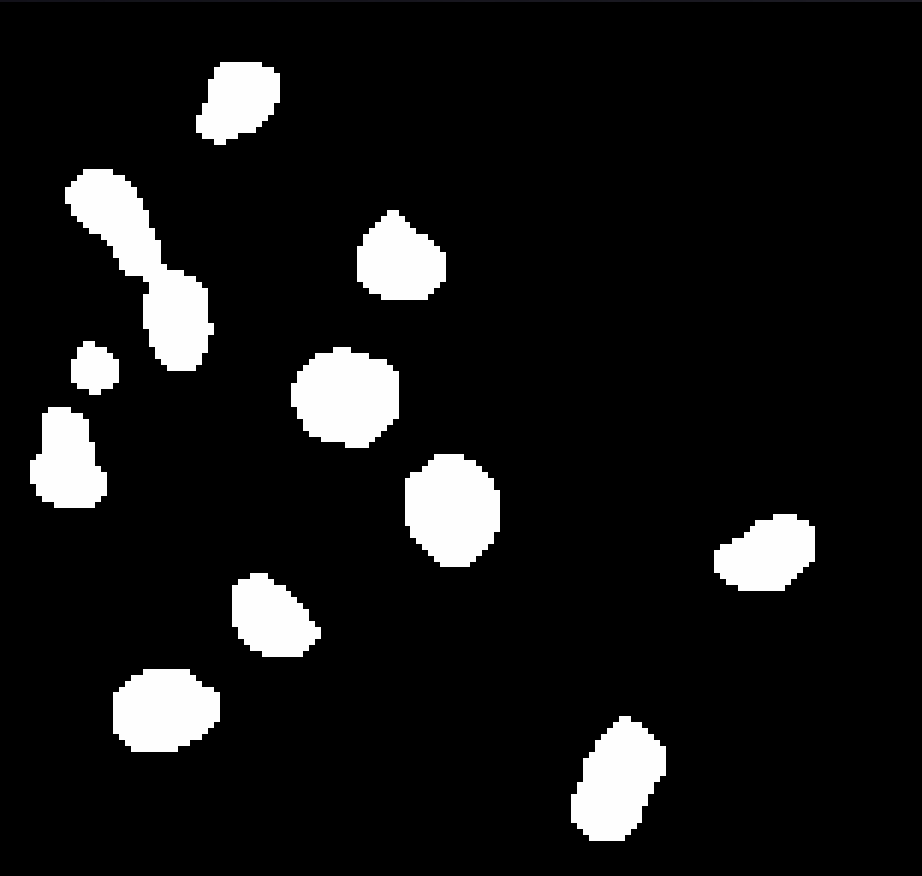


- Better Threshold
 - Grayscale Image \longrightarrow Binary Image
- Improve Segmentation

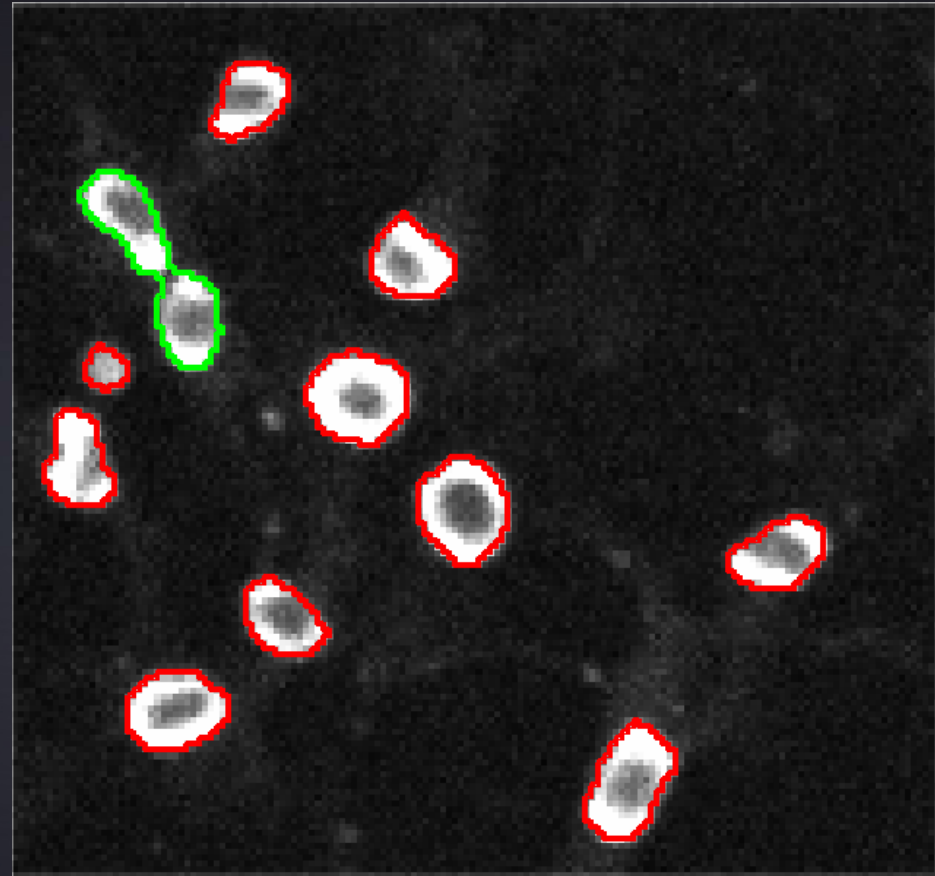
Future Work



Red = Normal

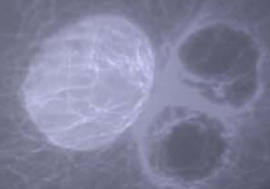


Green = Peanut



2hrs Hoechst Channel

Acknowledge



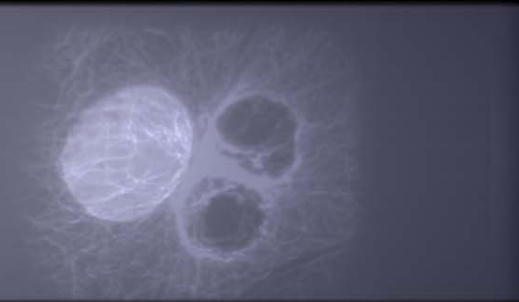
Acknowledge



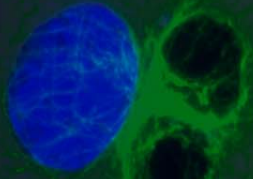
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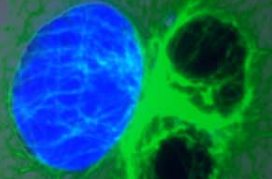
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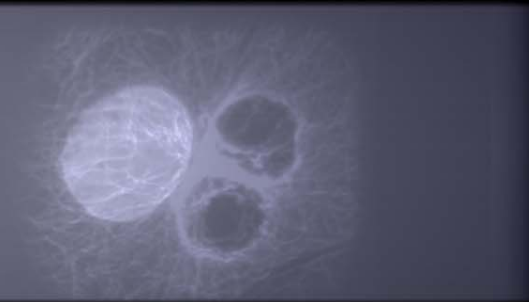
- Faculty Advisor
 - Dr. B.S. Manjunath
- Mentors:
 - Jiyun Byun
 - DeeAnn Hartung
- Center for Bio-Image Informatics



Questions?



Alzheimer's



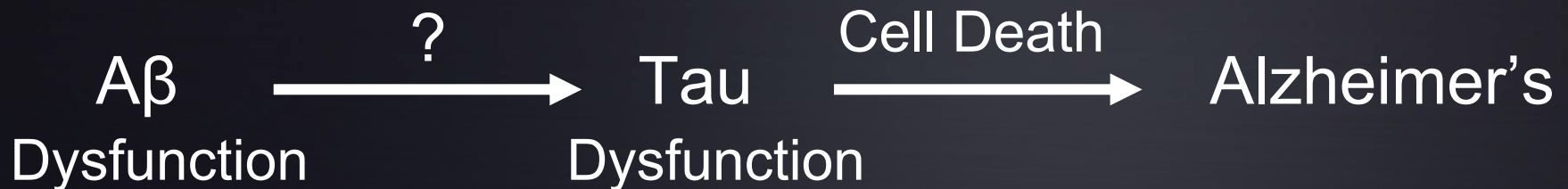
Characterized by two pathological hallmarks in the brain,

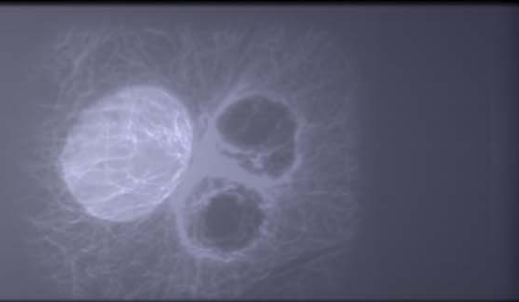
1. Extracellular Amyloid Plaques

➤ Amyloid-Beta ($A\beta$) protein

2. Intracellular Neurofibrillary Tangles (NFTs)

➤ Tau protein

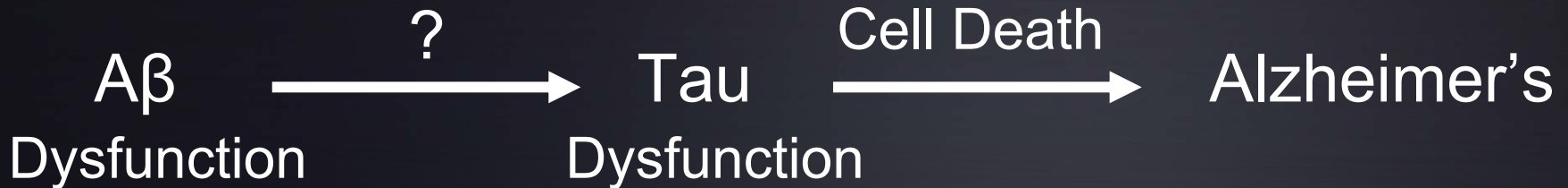




Alzheimer's

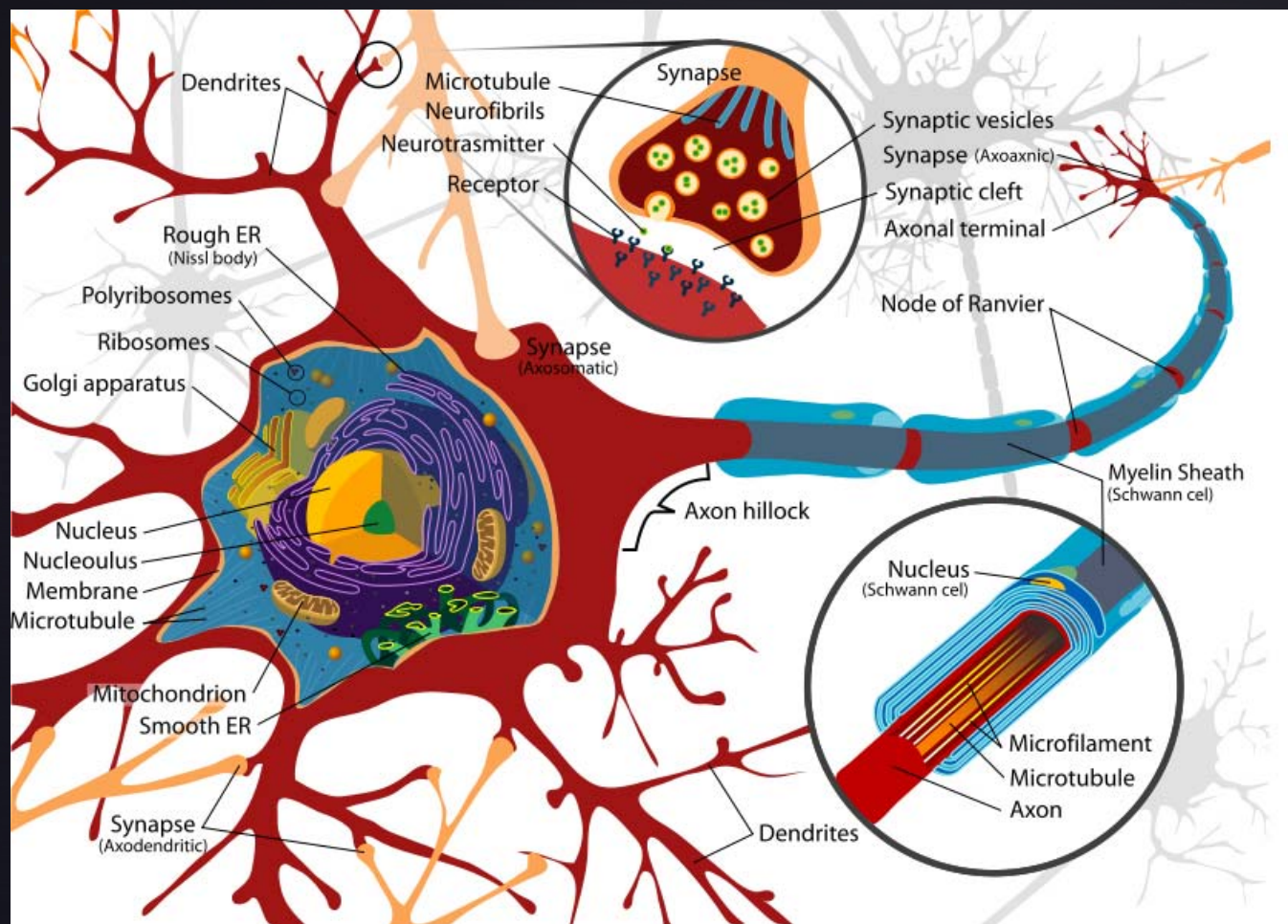
Amyloid-Beta ($A\beta$)

1. Amyloid Precursor Protein (APP)
 - a. Function: Undetermined
2. Can't cause disease without Tau



Alzheimer's

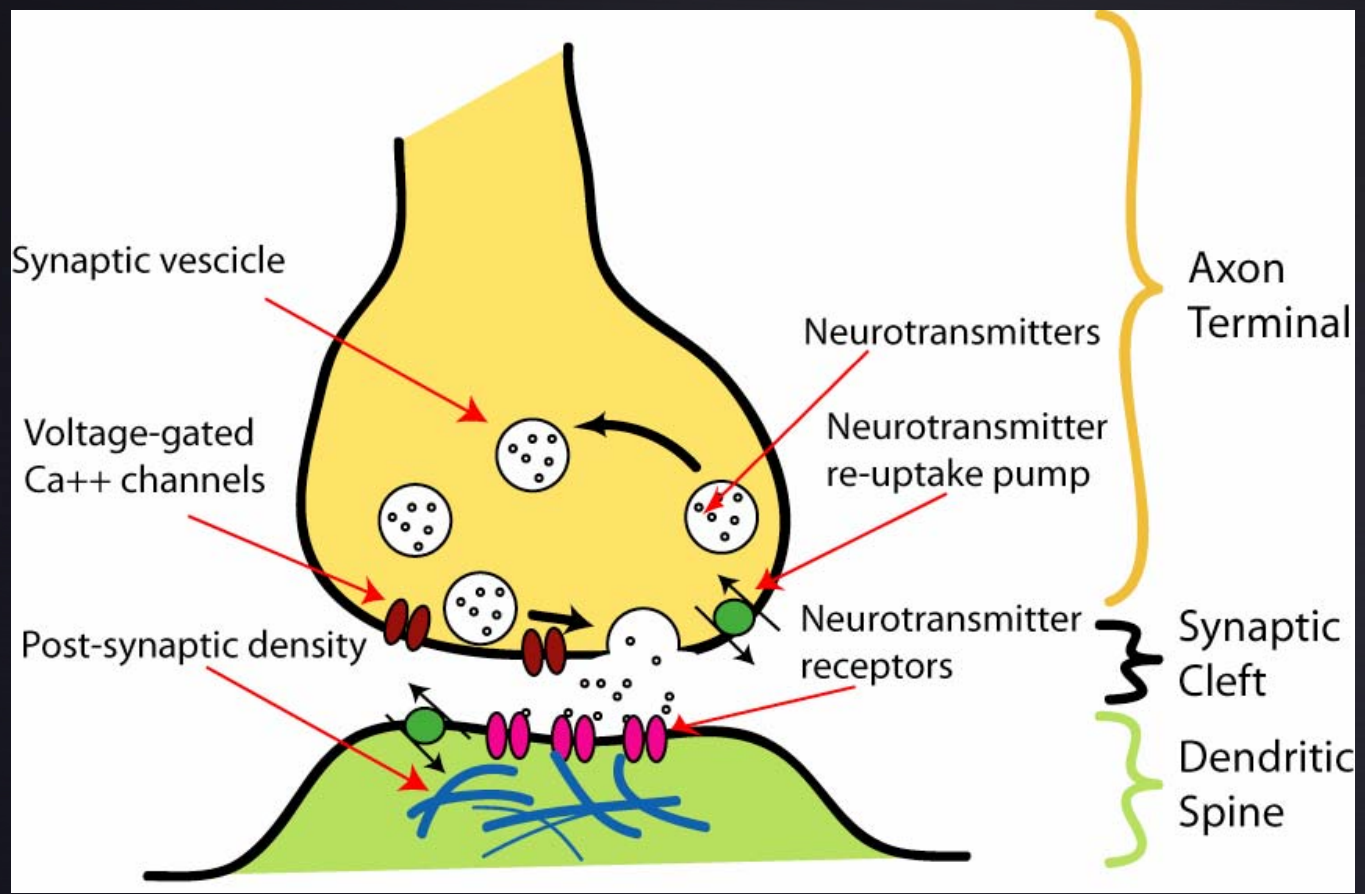
Amyloid-Beta ($A\beta$)



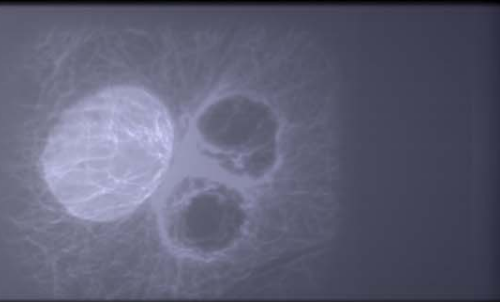
Alzheimer's



Amyloid-Beta ($A\beta$)



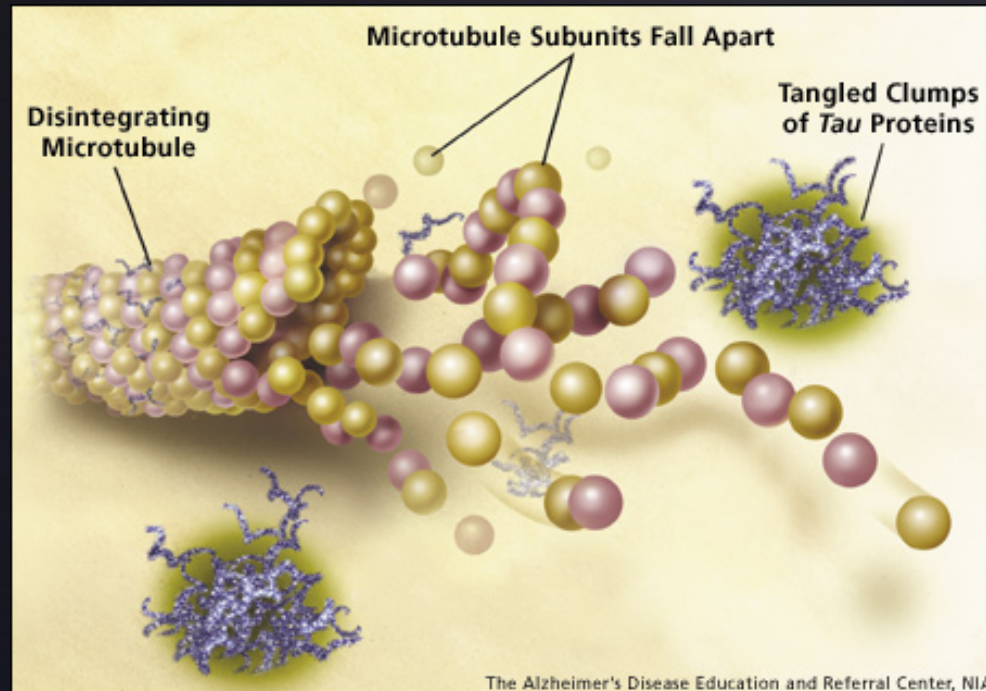
Alzheimer's



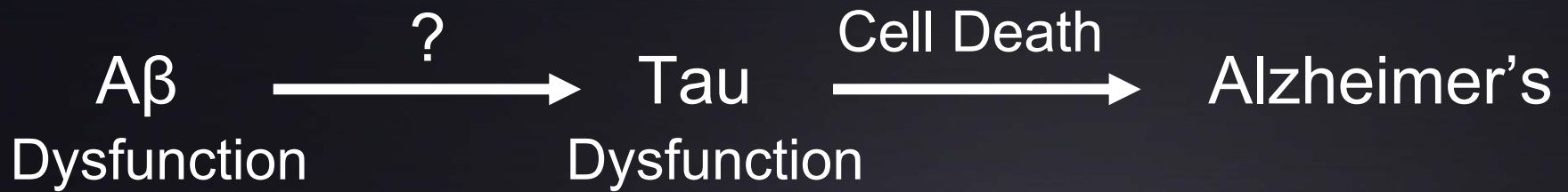
Tau Protein

1. Function:

Regulates neuronal microtubule dynamics



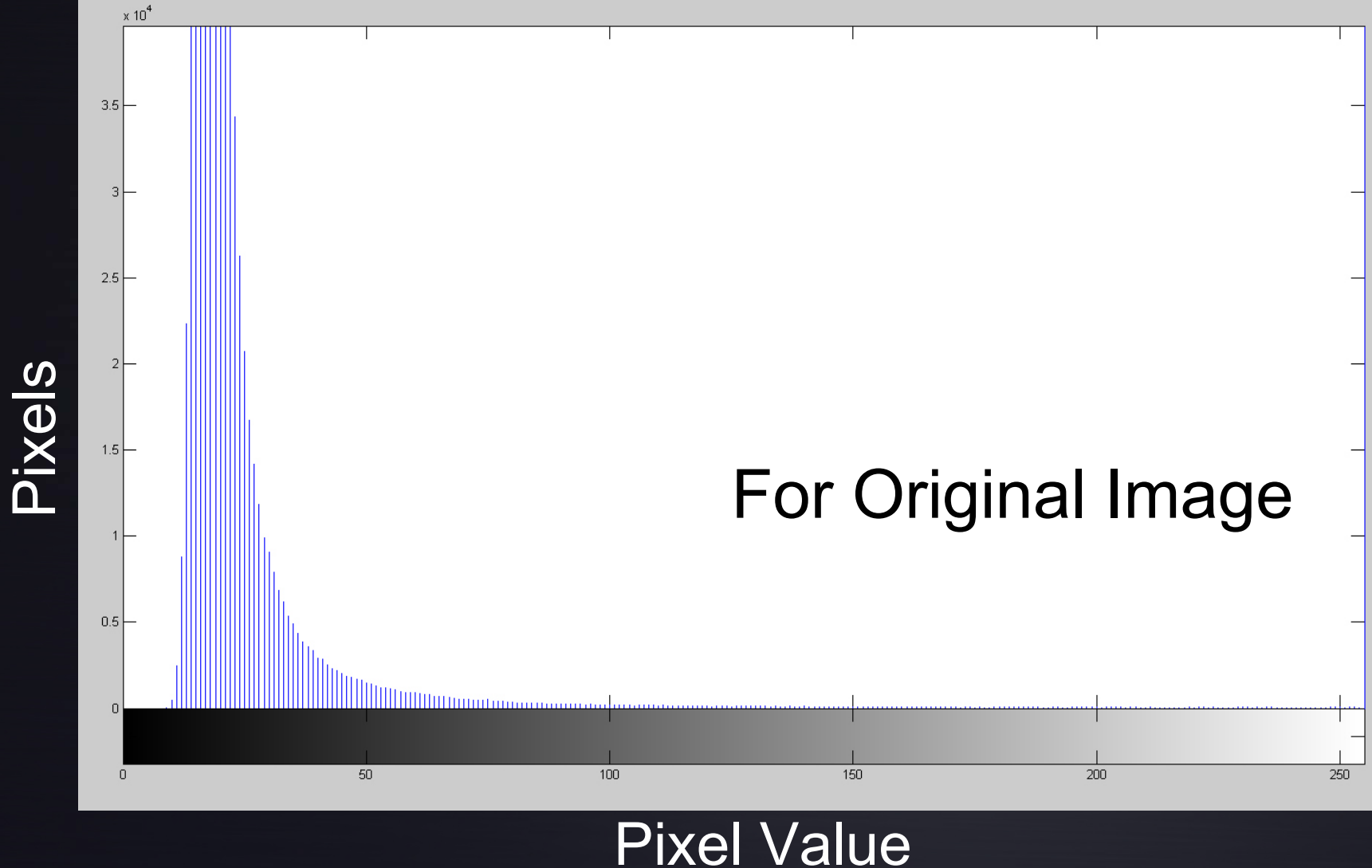
Alzheimer's



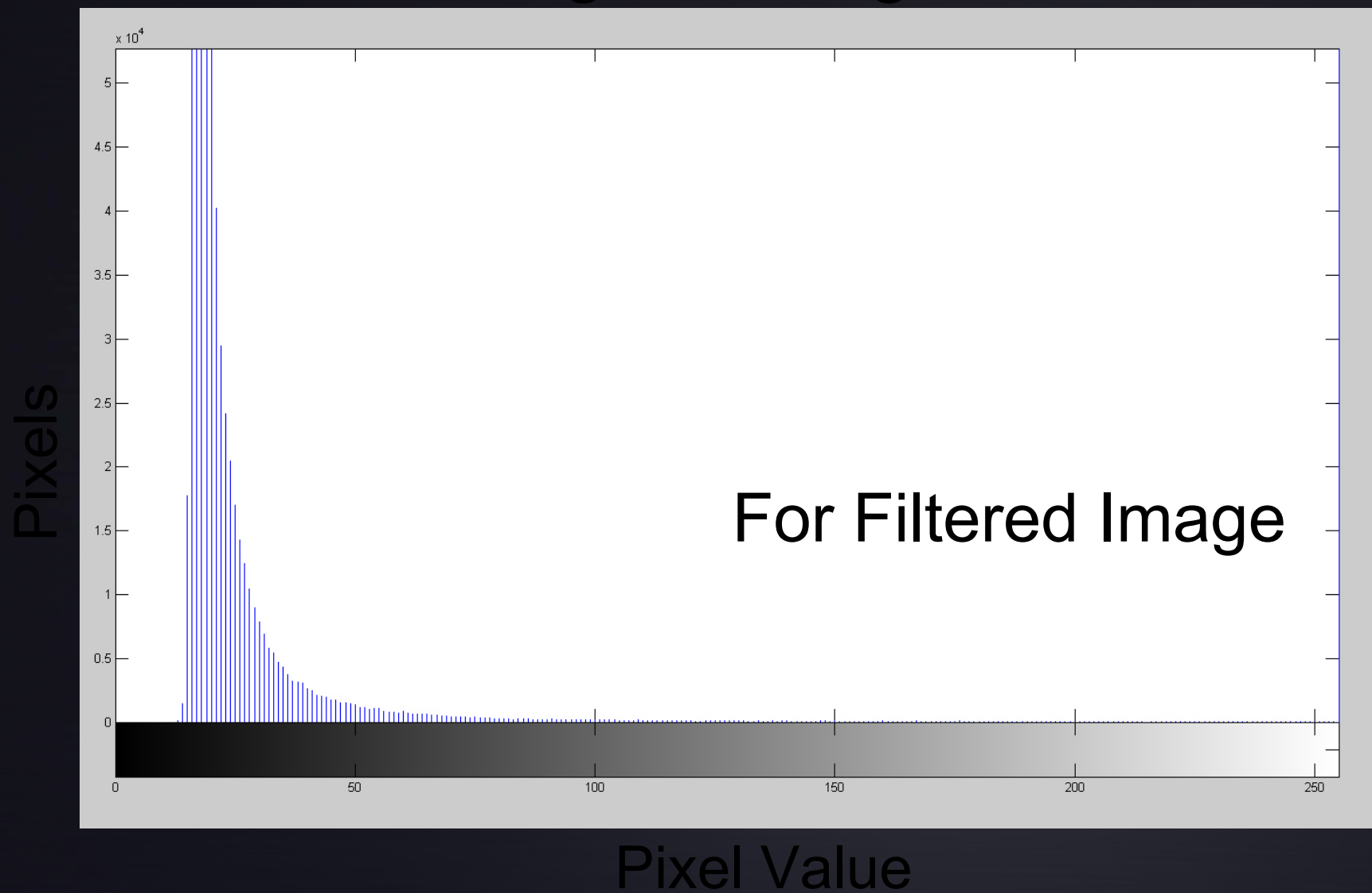
Experimental Setup

- Treatments
 - Untreated
 - 5 μM Staurosporine
 - 10 μM $\text{A}\beta$
 - 100 nM Taxol
 - 100nM Taxol +
10 μM $\text{A}\beta$
- Timepoints
 - 2 hours
 - 6 hours
 - 12 hours
 - 24 hours
 - 48 hours
 - 72 hours
 - 120 hours

Threshold Logical Image



Threshold Logical Image



Threshold Logical Image

