

# Spectral Unmixing and Segmentation of Biomedical Hyperspectral Images

Fields Undergraduate Summer Research Program Project 9

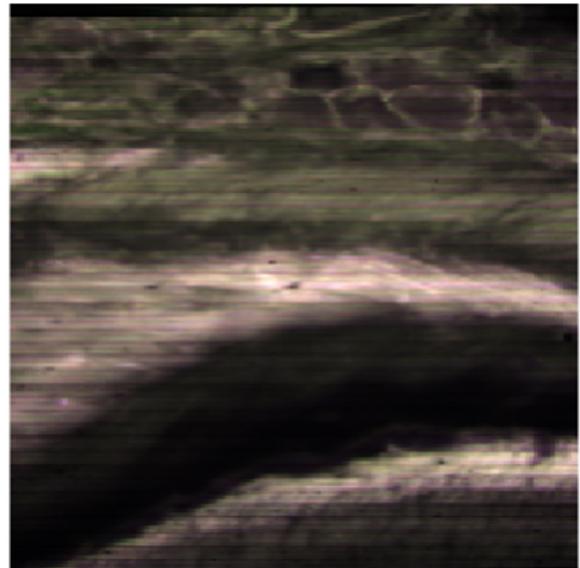


Xuanze (Charlie) Li, Aleksandar Popovic, Hannah Johnson

Supervised by Dr. Na Yu & Dr. You Liang

# Outline

- Project Introduction
- Hyperspectral Unmixing
- Hyperspectral Segmentation
- Dimension Reduction

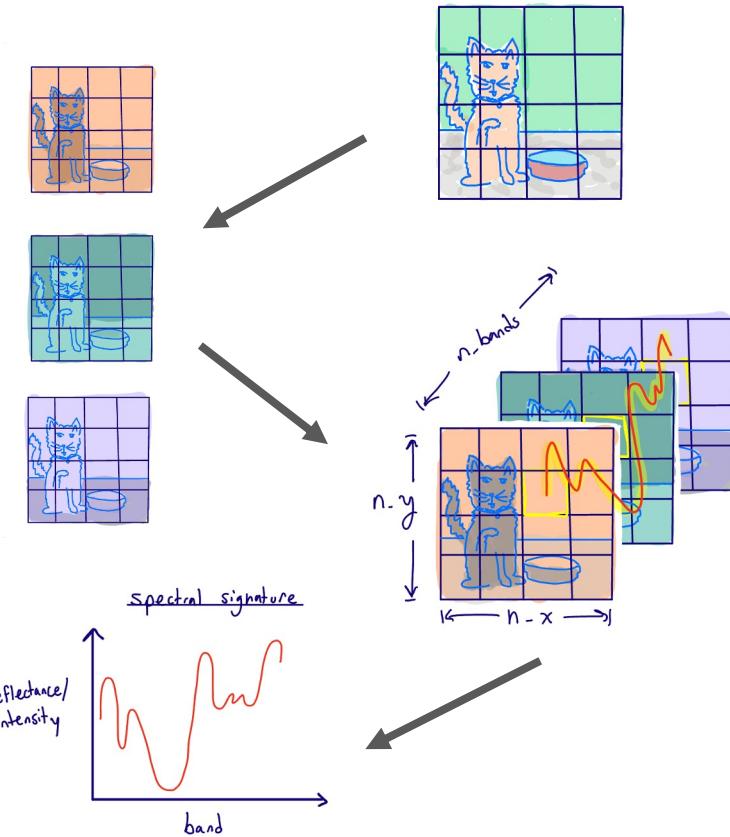


# Project Introduction

- ◆ What is Hyperspectral Imaging?
- ◆ Hypercube and Spectral Signatures
- ◆ Applications to Biomedicine
- ◆ Reference Dataset Information

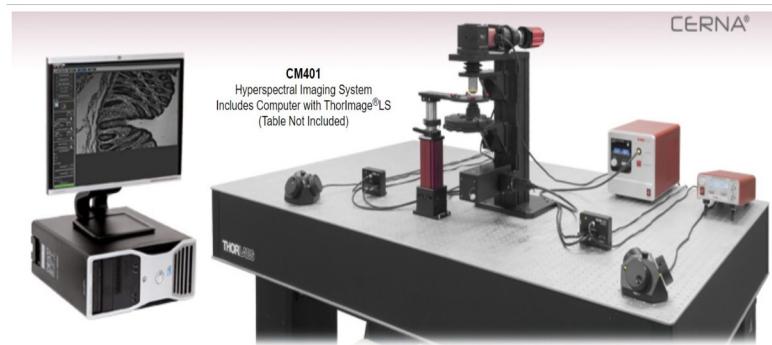
# What is Hyperspectral Imaging?

- Imaging camera measures light reflectance off of objects
- The camera separates the image by wavelength
- Different **endmembers**, or materials, in the image reflect different amounts of light at each wavelength
- Endmembers will have different **spectral signatures** which can be used to classify elements in the image
- Commonly applied to geospatial remote sensing - ex. vegetation covers



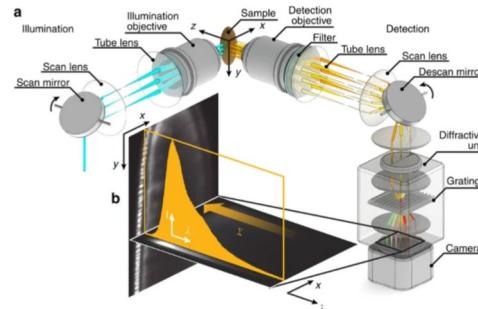
# Biomedical Hyperspectral Imaging

- Applying traditional geospatial HSI techniques to **biomedical** data
- Create **open source** python package that performs dimensionality reduction, unmixing, and segmentation
- Applying HSI techniques to eye slice samples
- A special microscope called a **spectrometer** is used to collect hyperspectral data from the samples



[https://www.thorlabs.com/newgroupage9.cfm?objectgroup\\_id=11095](https://www.thorlabs.com/newgroupage9.cfm?objectgroup_id=11095)

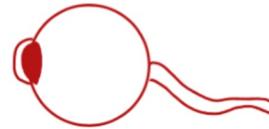
Figure 1: Hyperspectral SPIM setup and image formation.



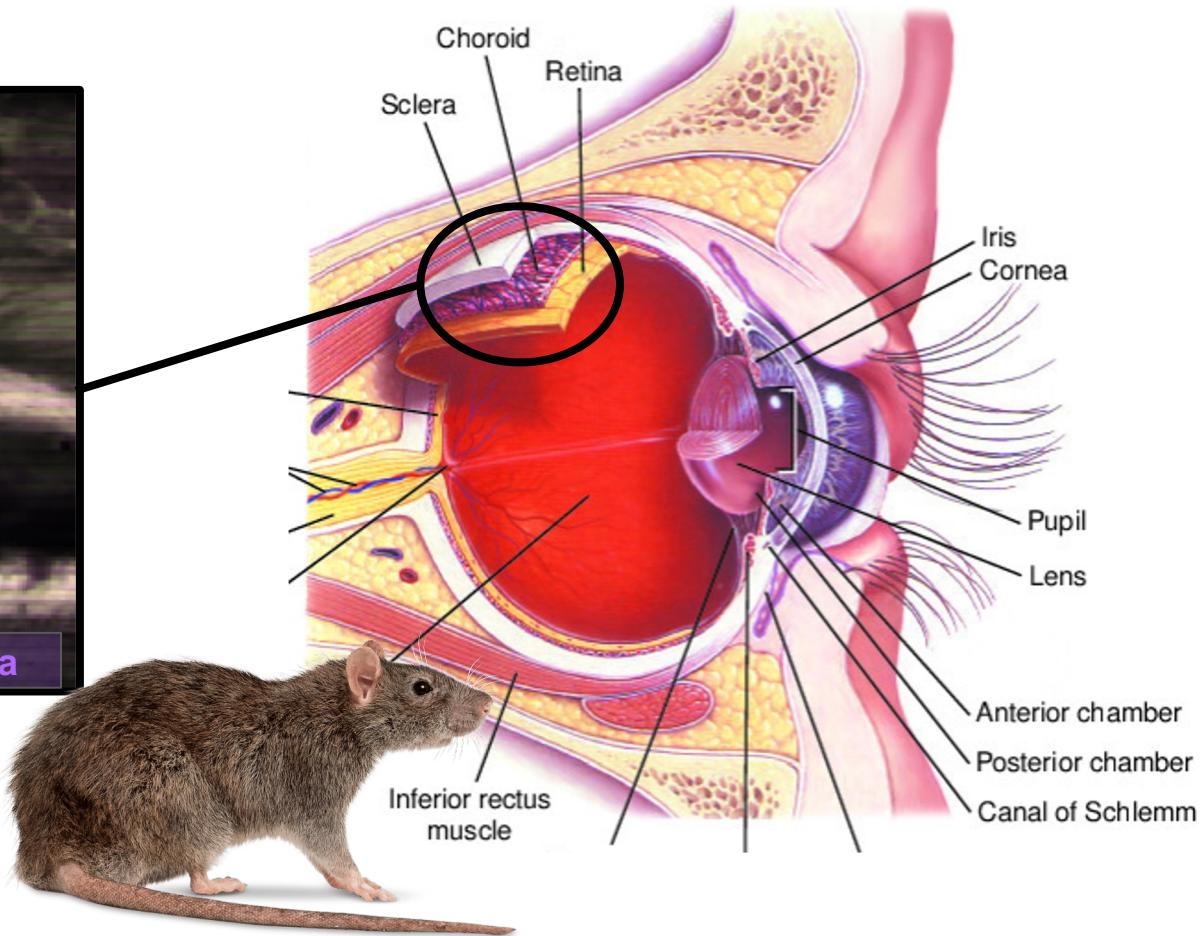
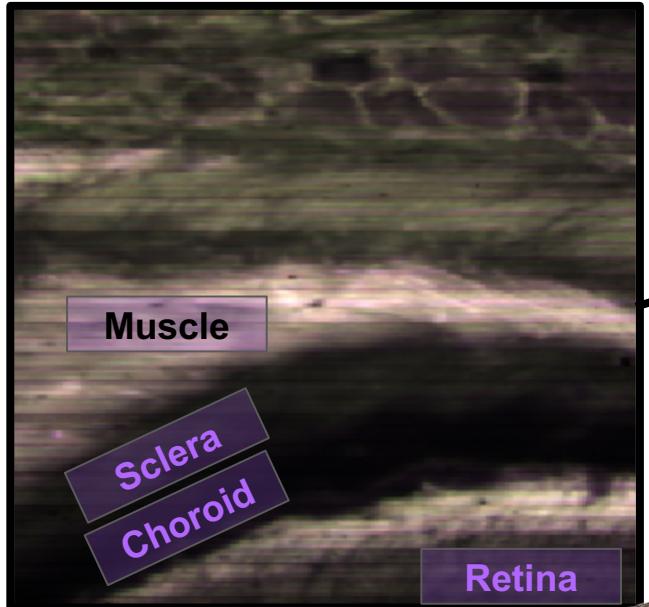
<https://www.nature.com/articles/ncomms8990>

# Applications of Biomedical HSI

- Dr. Yeni Yucel at Eye Pathology Lab
- Effect of going into space on the eyes
- SANS: Spaceflight-Associated Neuro-Ocular Syndrome
- In space, fluids in the body pool in the head
- Result in flattening of the back of the eye and retinal nerve fiber thickening



# Eye Anatomy



# Reference Dataset

For the sake of this summer project, we focused on the lower half of the image containing the **Muscle**, **Sclera**, **Choroid**, and **Retina**

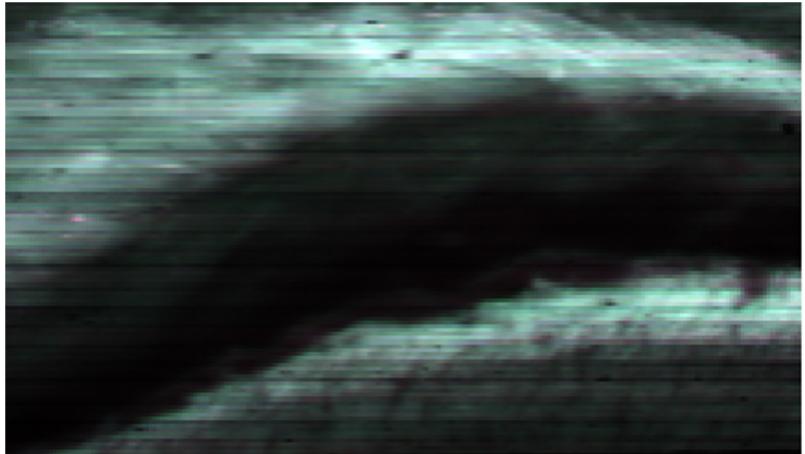
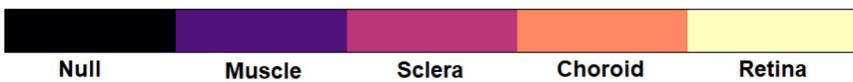
**Spectral Range:** 528 to 836 nm (NUV to NIR)

**Band Information:** 78 bands at width of 4.0 nm

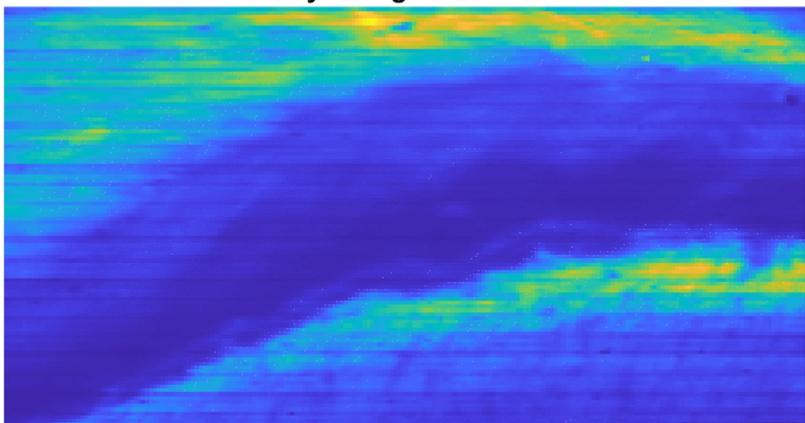
**Dataset Information:**

$$n_x = 210 \quad n_y = 120 \quad n_{\text{bands}} = 78 \quad n_{\text{end}} = 4$$

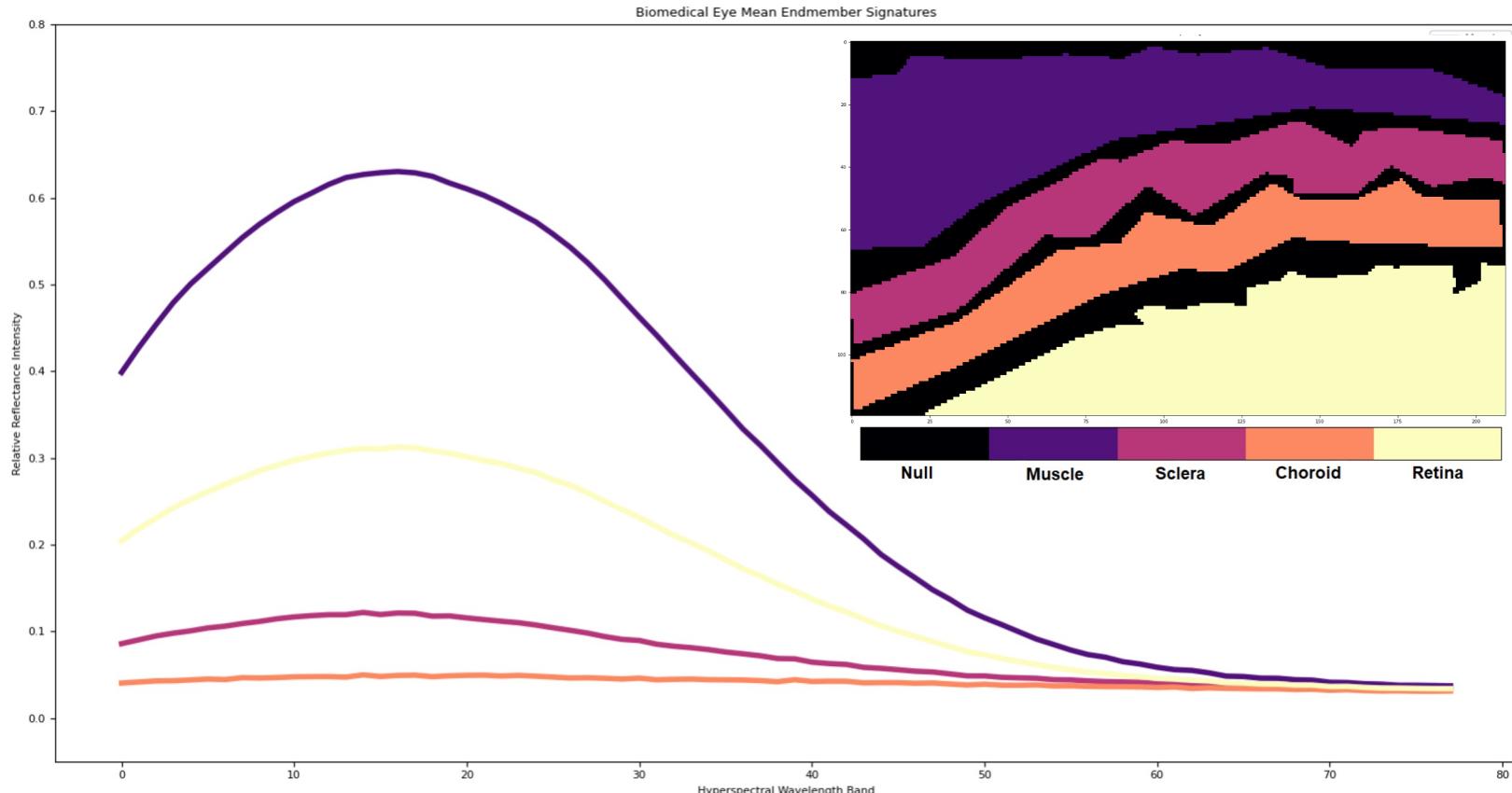
**Endmembers:**



Eye Image Band 1



# Reference Dataset



# What We've Accomplished

We've created a **open-source** Python package for Biomedical Hyperspectral Imaging called **BHSIpy**:

## Linear Unmixing Methods:

Supervised Methods: CVXOPT, Gradient Descent, Active Set

Unsupervised Methods: UFCLSU

## Segmentation Methods:

Supervised: 3D-HyperUNet

Semi-Supervised: SAM, SIDSAM, JMSAM, NS3, K Means

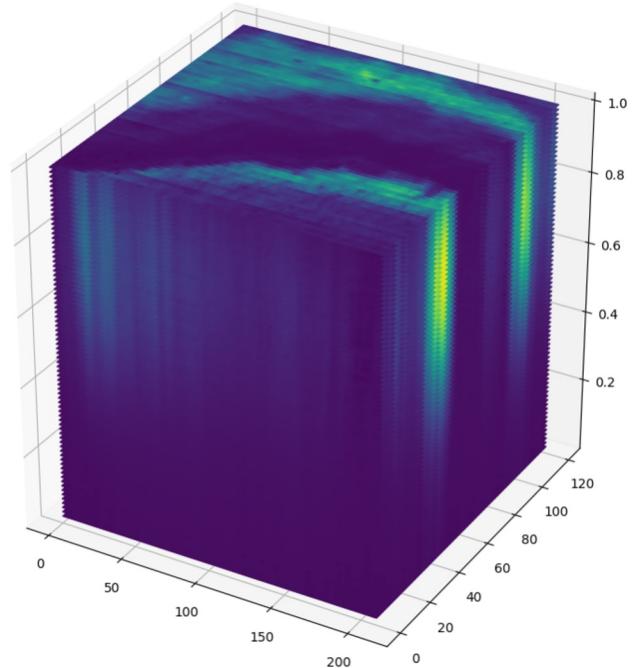
## Dimension Reduction:

Linear Band Selection and Principal Component Analysis

## Visualization Methods:

3D Plotting Method for Hyperspectral Cubes

General Layer Plots for Unmixing and Segmentation

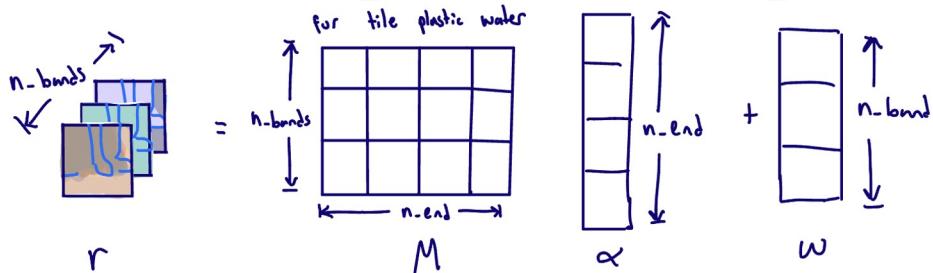


# Hyperspectral Unmixing

- ◆ What is Unmixing?
- ◆ Unmixing Methods
- ◆ Results

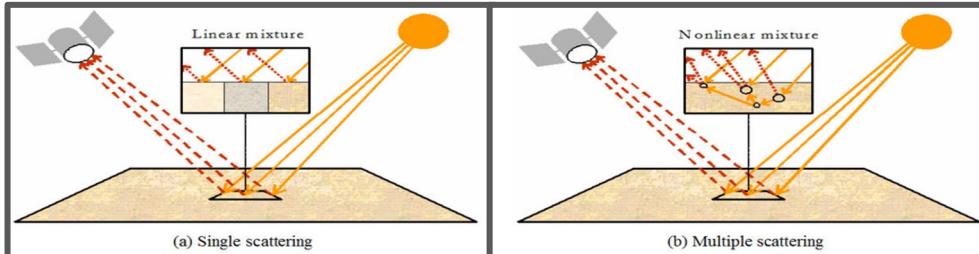
# Linear Hyperspectral Unmixing

Pixels are assumed to be **linear combinations** of the endmember signatures in the image:



**Formally:**

$$r = \begin{bmatrix} | & | & | \\ m_1 & m_2 & \cdots & m_{n_e} \\ | & | & | \end{bmatrix} \begin{bmatrix} a_1 \\ a_2 \\ \vdots \\ a_{n_e} \end{bmatrix} + \begin{bmatrix} w_1 \\ w_2 \\ \vdots \\ w_{n_b} \end{bmatrix} = \sum_{i=1}^{n_e} m_i \alpha_i + w = M\alpha + w$$



Linear Unmixing aims to solve the following **optimization** problem:

$$\min_{\alpha} \|M\alpha - r\|_2^2$$

**Unconstrained:**  $\alpha = (M^T M)^{-1} M^T r$

**Non-negative Constraint:**  $\alpha_i \geq 0$

Solved using **convex optimization** packages or **active-set methods** or **gradient descent**

**Fully Constrained:**  $\alpha_i \geq 0 \quad \sum a_i = 1$

Reformat in Non Negative Constrained Least Squares Problem\*\*:

$$M^* = \begin{bmatrix} \delta M \\ 1^T \end{bmatrix} \quad r^* = \begin{bmatrix} \delta r \\ 1 \end{bmatrix}$$

$$\min_{\alpha} \|M^* \alpha - r^*\|_2^2 \quad \alpha_i \geq 0$$

\*\* D. Heinz and C.-I Chang, "Fully constrained least squares linear spectral mixture analysis method for material quantification in hyperspectral imagery," IEEE Transactions on Geoscience and Remote Sensing, vol. 39, no. 3, pp. 529-545, 2001.

# Linear Unmixing Methods

## Iterative Active Set Methods

At the beginning of the project, we had used CVXOPT, however, we found that it was slow and computationally heavy.

We switch over to a faster **active-set** method\*:

**MATLAB:** *lsqlnneg*

**Python:** *nls*

From what we had seen, nobody really bothered to code the function in Python until we did.

## Gradient Descent Methods

- Applying Gradient Descent to Solve FCI SLI Problems\*

$$\alpha^* = \arg \min_{\alpha} J(\alpha)$$

$$\text{subject to } \alpha_i \geq 0, \quad i = 1, \dots, R$$

$$\sum_{i=1}^R \alpha_i = 1.$$

$$\alpha_j = \frac{w_j}{\sum_{\ell=1}^R w_\ell}$$

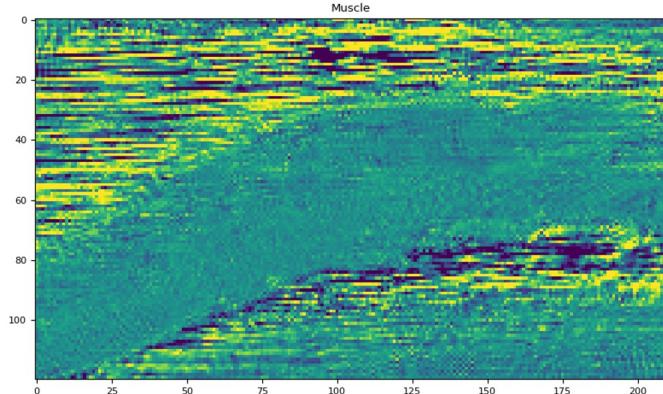
$$0 < \mu_i(k) \leq \frac{1}{[\nabla J(\alpha)]_i}$$

$$\alpha(k+1) = \alpha(k) + \dots$$

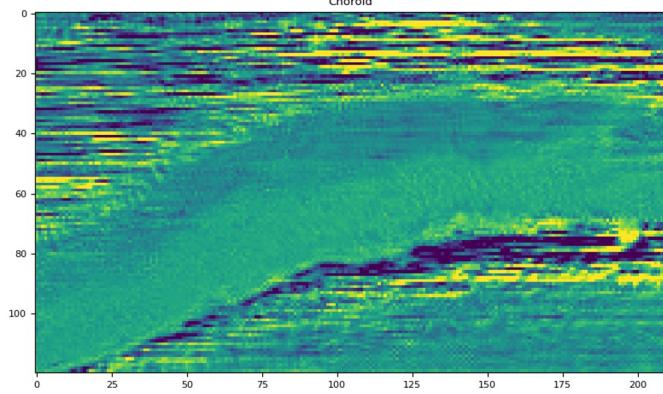
$$+ \mu \operatorname{diag}\{\alpha(k)\} [\nabla_{\alpha} J(\alpha) - \mathbf{1} \nabla_{\alpha} J(\alpha)^{\top} \alpha(k)]$$

# Unmixing Results

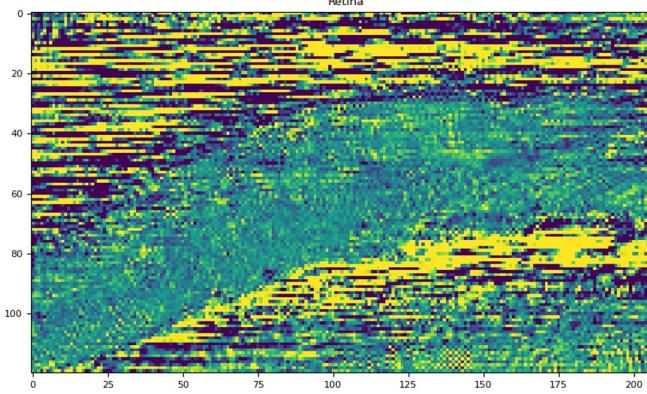
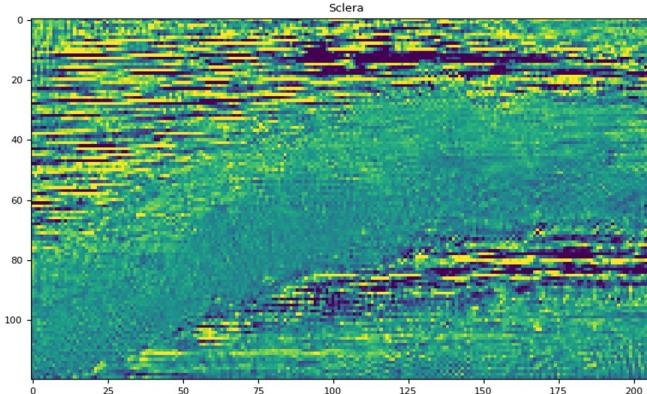
$$\alpha = (M^T M)^{-1} M^T r$$



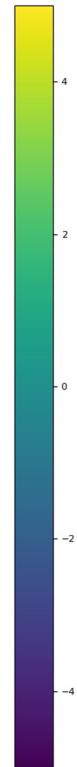
Unconstrained Least Squares Unmixing  
Biomedical Rat Eye Image



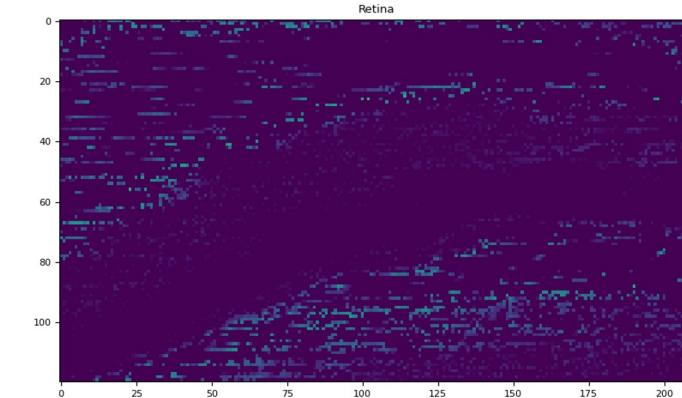
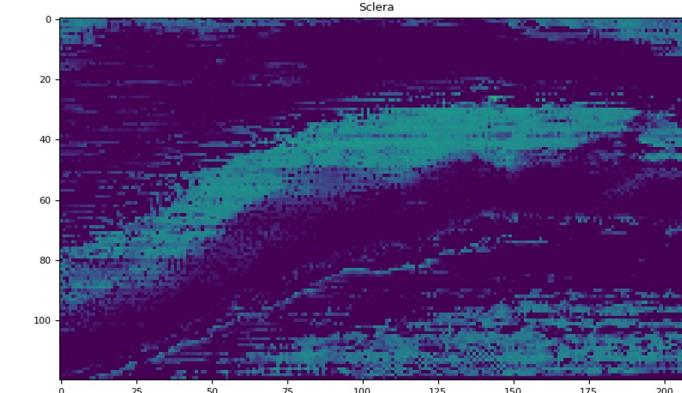
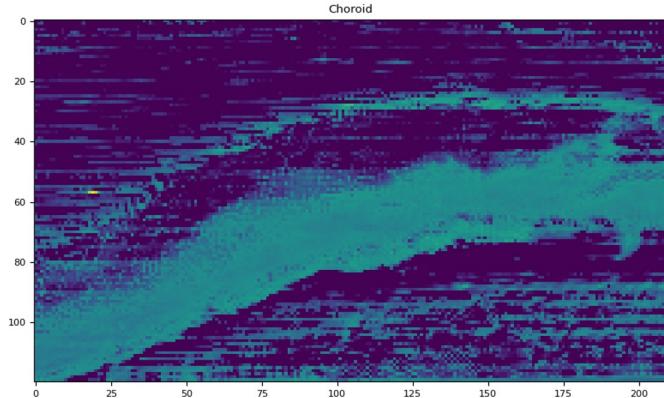
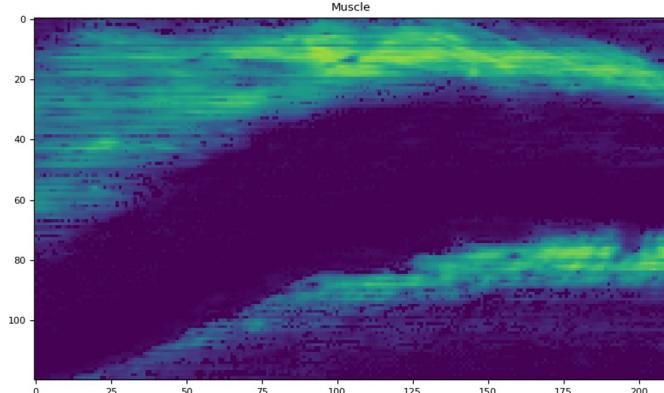
Pixelwise Average LSE: 0.0011



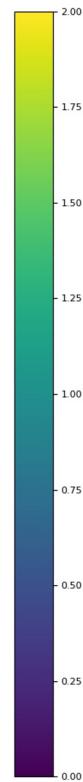
Running Time: 0.074 s



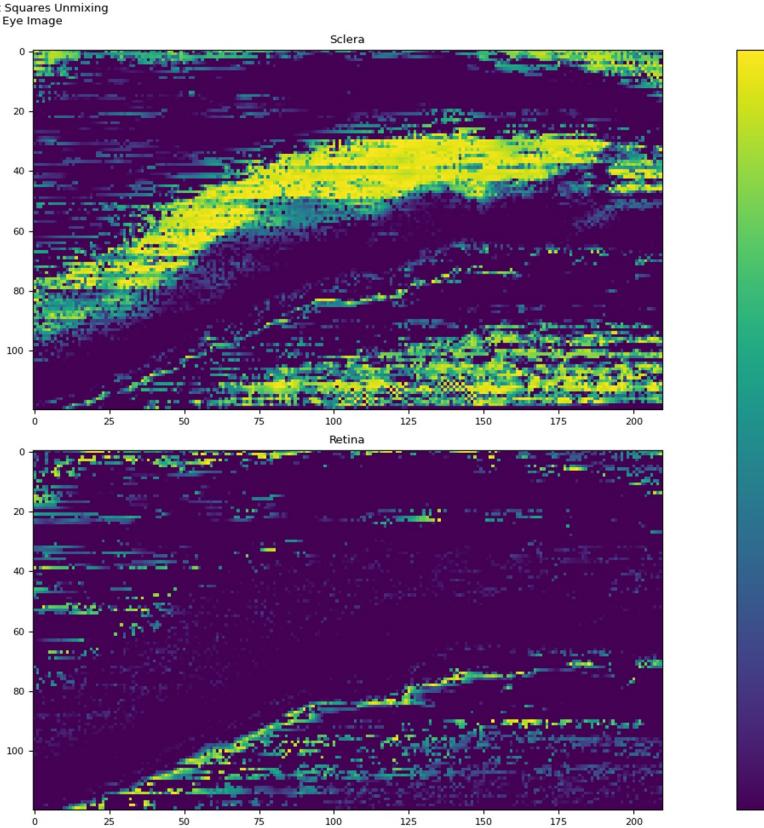
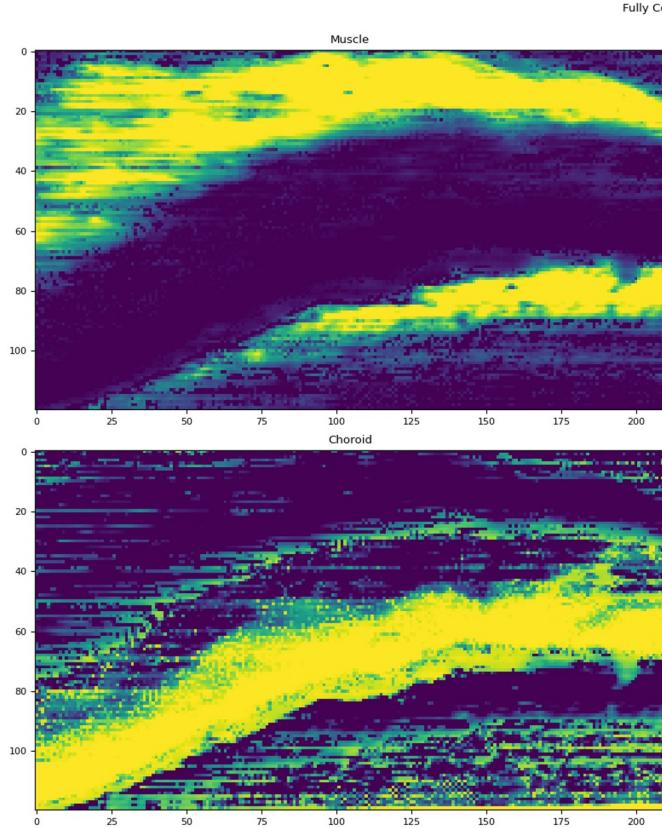
# Unmixing Results



$$\alpha_i \geq 0$$



# Unmixing Results



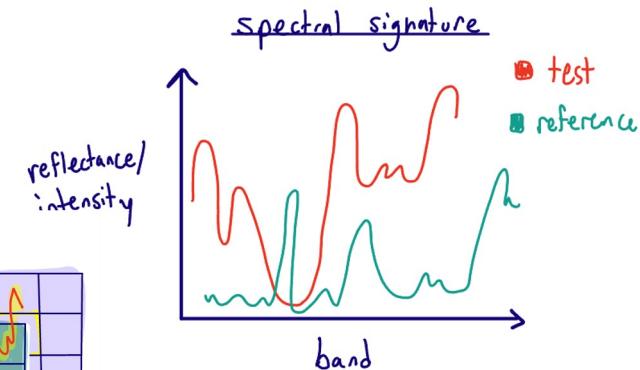
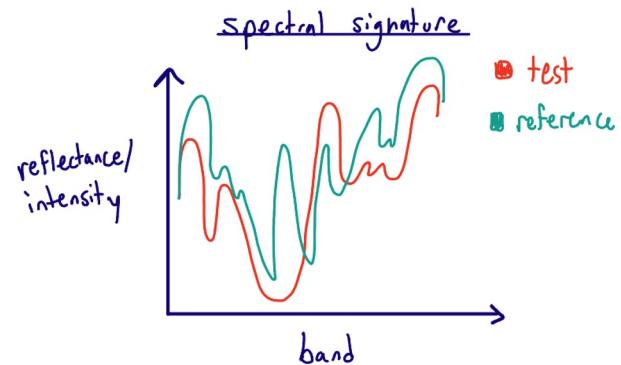
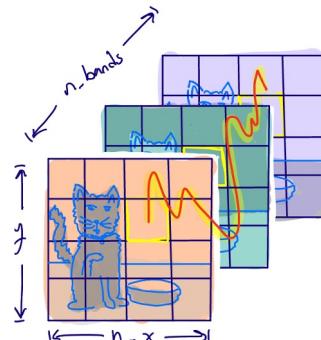
# Hyperspectral Segmentation

- ◆ What is Segmentation?
- ◆ Segmentation Measures Overview
- ◆ Results

# What is Segmentation?

A method to determine endmembers in an image

- Assume each pixel is ***pure***
- Compare a pixel from the image, the ***test pixel***, with the ***reference spectrum*** of an endmember
- Use various measure to determine similarity of the test ***spectral signature*** and reference ***spectral signature***



# Segmentation Measures

We represent both the test and reference signatures as n-dimensional vectors:

$$s = (s_1, s_2, \dots, s_{n_{\text{bands}}})$$

$$t = (t_1, t_2, \dots, t_{n_{\text{bands}}})$$

Techniques for measuring similarity:

$$1. \|s - t\| = \left[ \sum_{i=1}^N [s_i - t_i]^2 \right]^{\frac{1}{2}}$$

$$2. \theta_{t,s} = \arccos \left( \frac{t \cdot s}{\|t\| \|s\|} \right)$$

$$3. SID(\mathbf{r}, \mathbf{r}') = \sum_{n=1}^L p_j \log \left( \frac{p_j}{q_j} \right) + \sum_{n=1}^L q_j \log \left( \frac{q_j}{p_j} \right)$$

## Measures:

- SAM
- SID

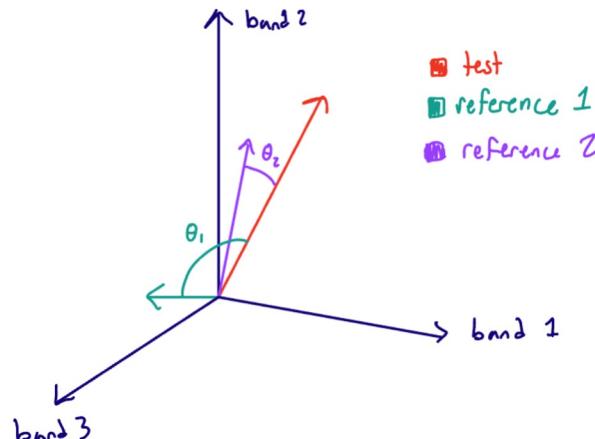
## Mixed

## Measures:

- SID-SAM
- JMSAM
- NS3

Spectral signature array

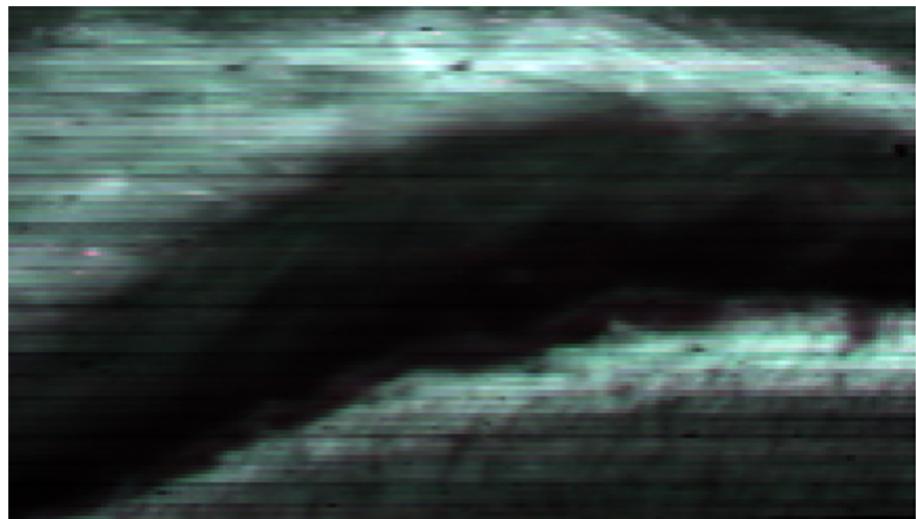
band	1	2	3	4	5	6	7	8
intensity	78	67	46	49	50	80	92	102



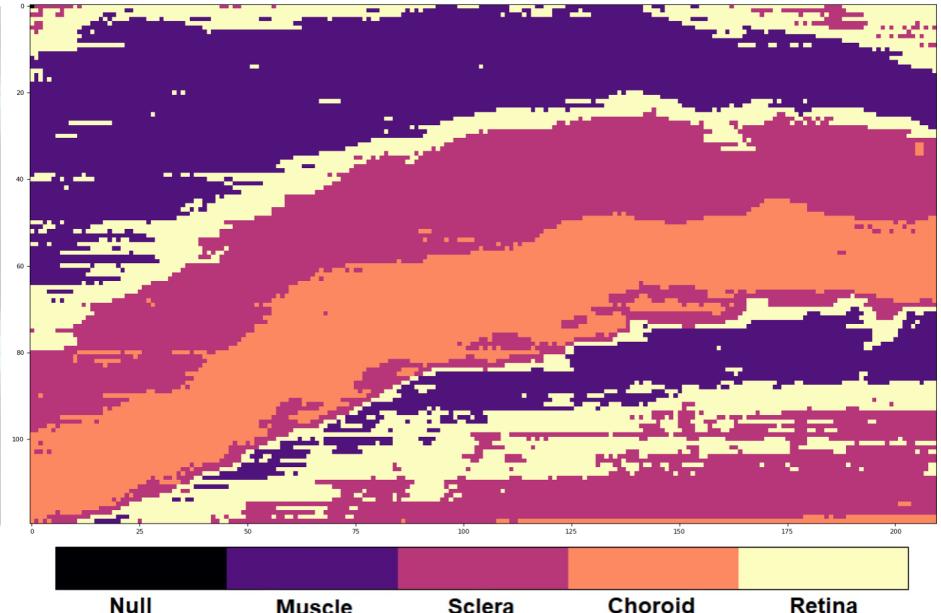
# Spectral Angle Mapper Segmentation

Angle

Original Image



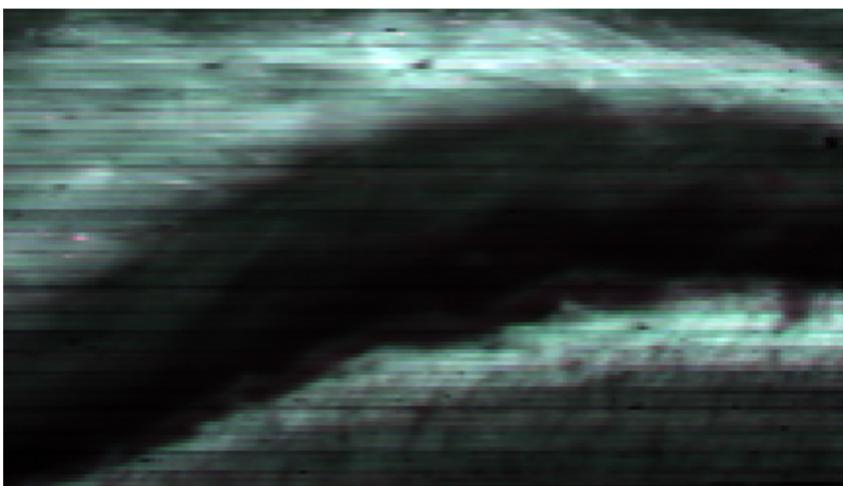
Segmented Image



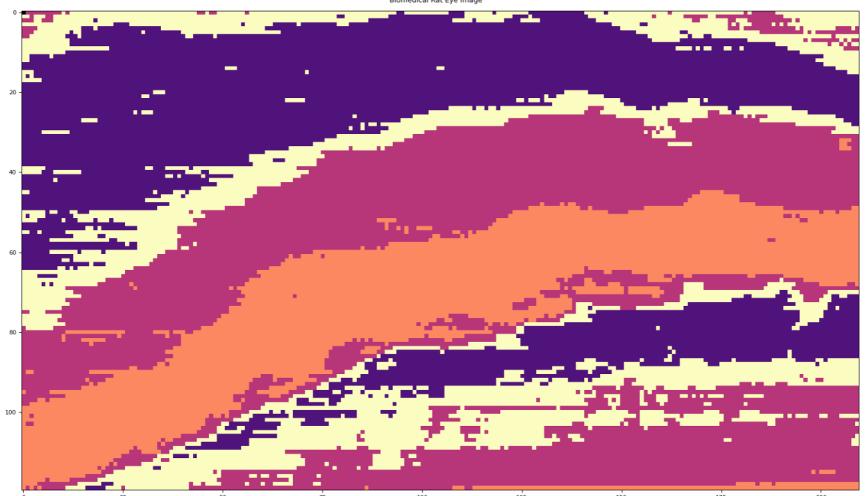
# SIDSAM TAN Segmentation

Angle, Probabilistic

Original Image



Segmented Image

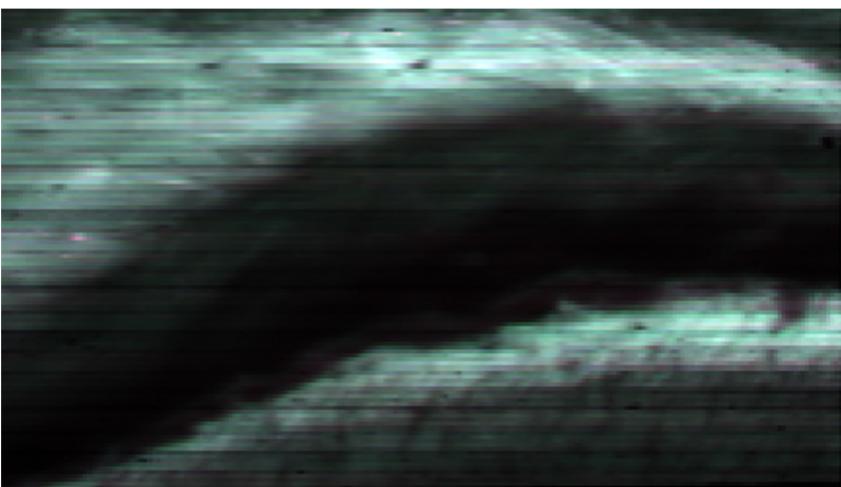


Null      Muscle      Sclera      Choroid      Retina

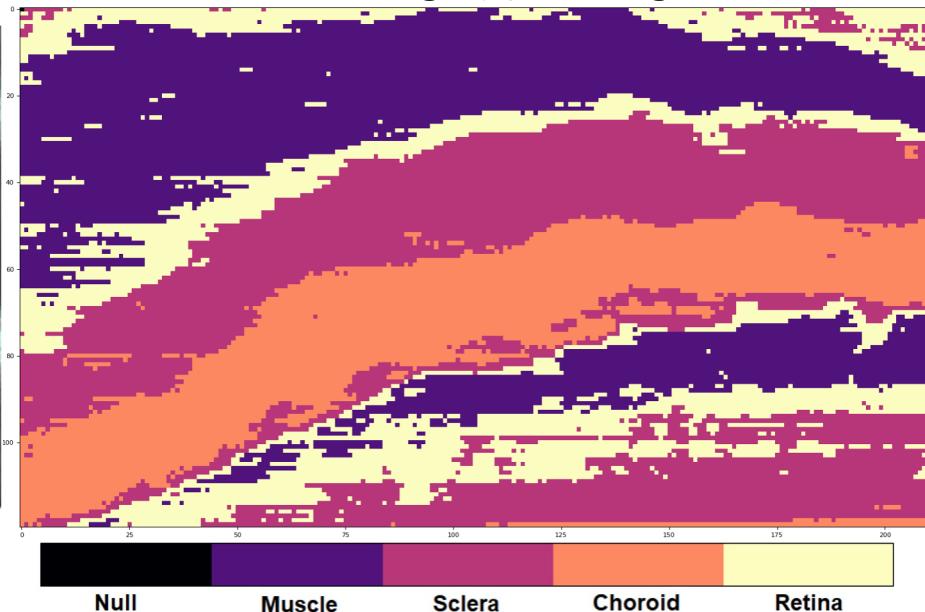
# JMSAM TAN Segmentation

Angle, Distance

Original Image



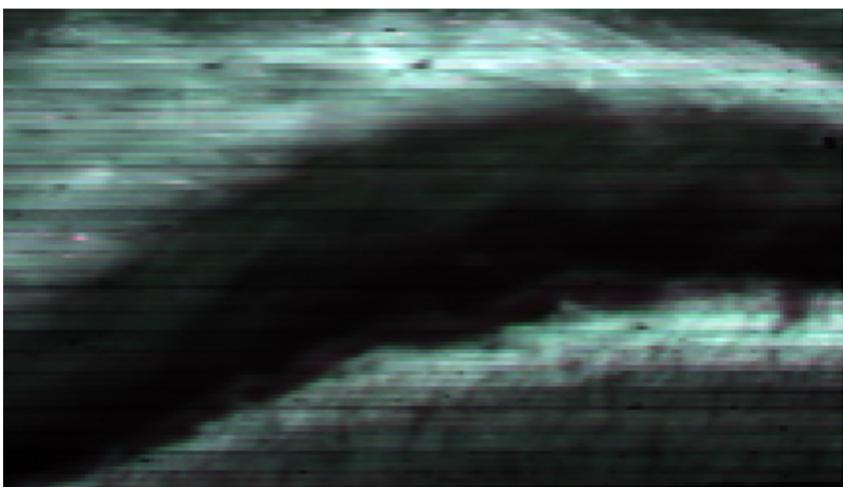
Segmented Image



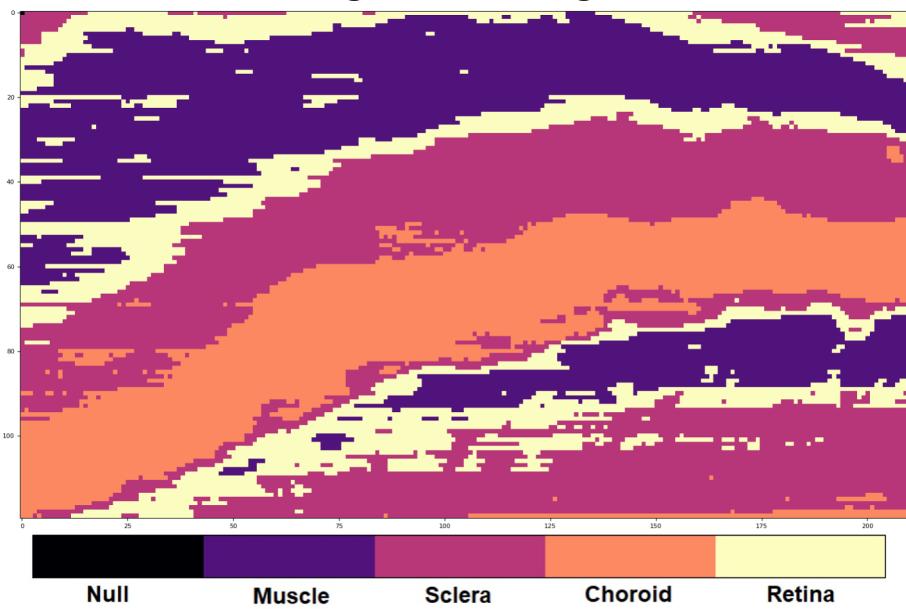
# NS3 Segmentation

Angle, Distance

Original Image



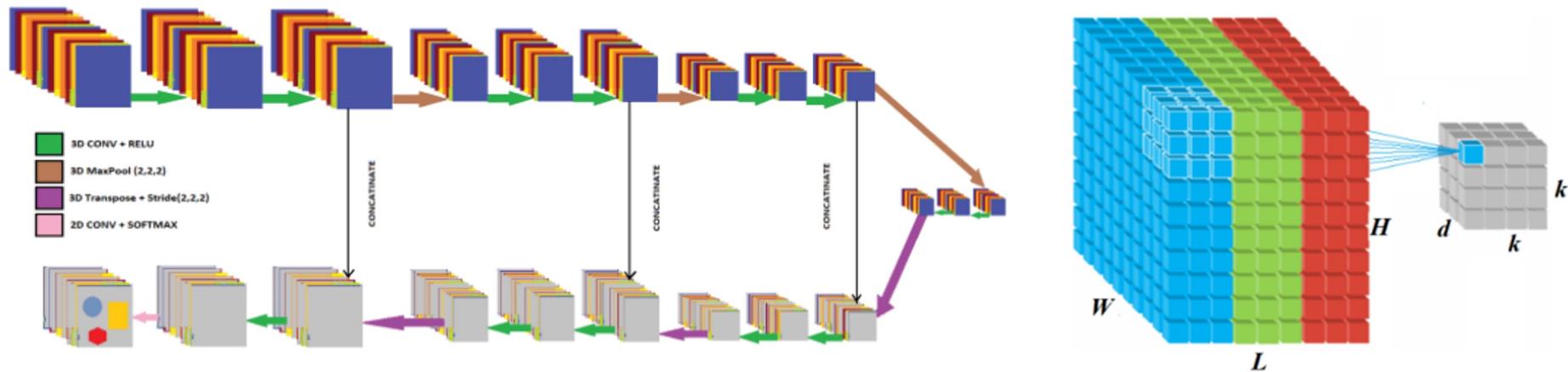
Segmented Image



# Deep Learning Segmentation

We adapt the **3D Hyper-UNet\*** for geospatial hyperspectral and multispectral imaging proposed by *Nischal et al.* to the realm of biomedical hyperspectral imaging

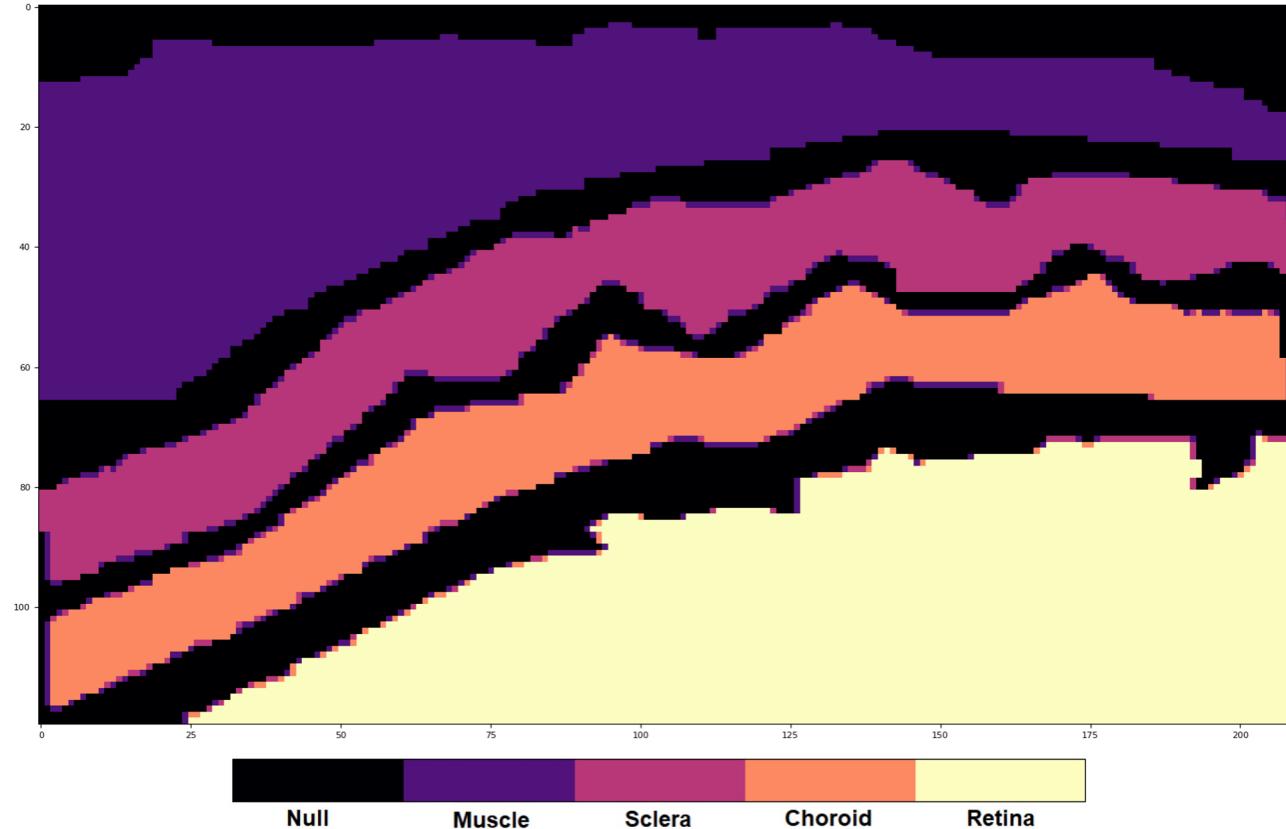
Combines both **spatial** and **spectral** information using 3D kernel convolutions



# Deep Learning Segmentation Results

```
unet = U_Net(dataset = 'biomedical_image', num_epochs=1000, reduced_size_x_y=144, n_features=5)
```

Trainable params: 5,028,101  
Non-trainable params: 1,344



# Dimension Reduction

- ◆ Linear Predictor Band Selection

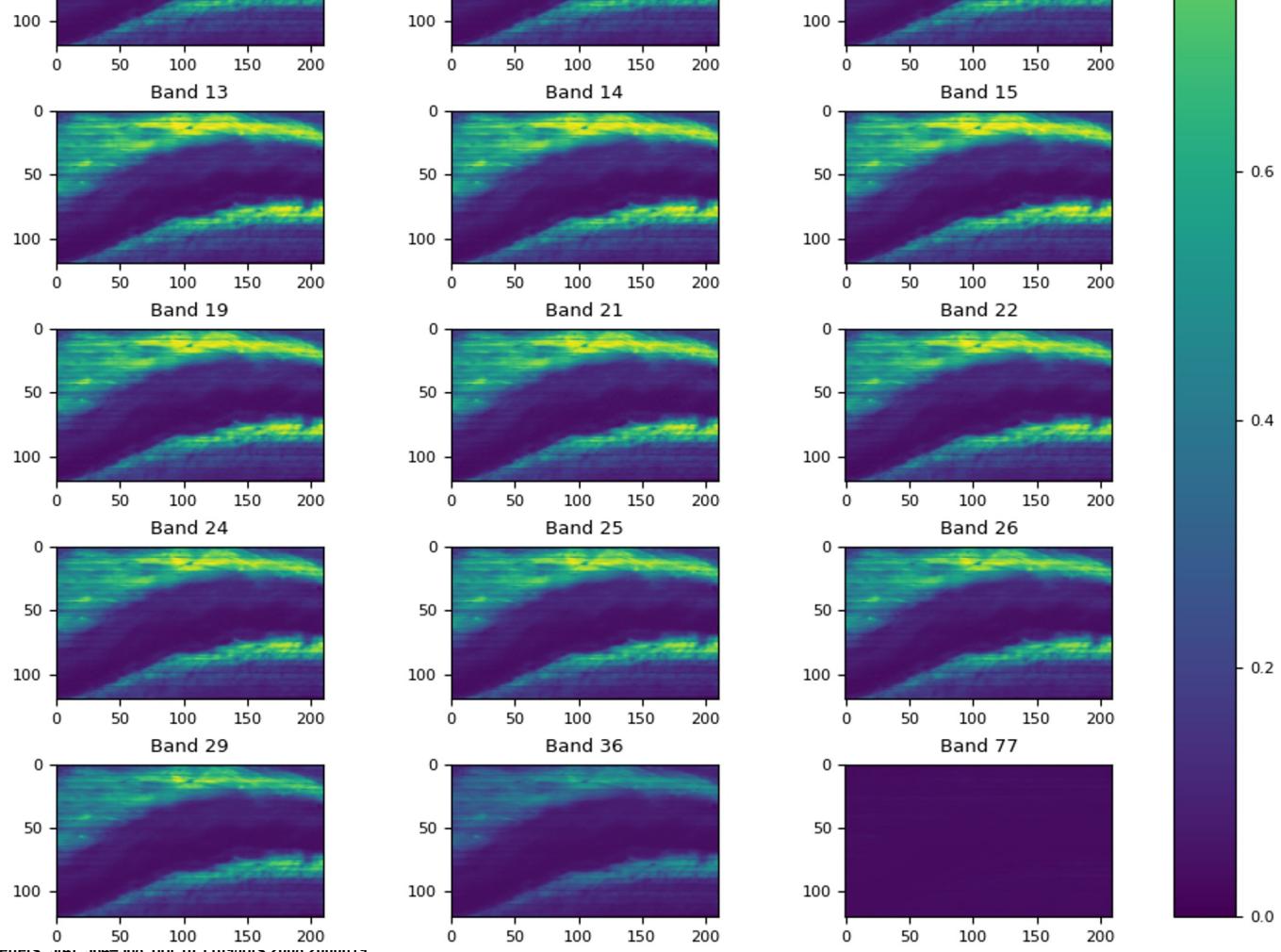
# Linear Band

## Motivation:

As the band width is so large  
the hyperspectral image  
We do not gain that much  
keeping adjacent bands  
choose only the most relevant  
our unmixing and segmentation

## Algorithm\*:

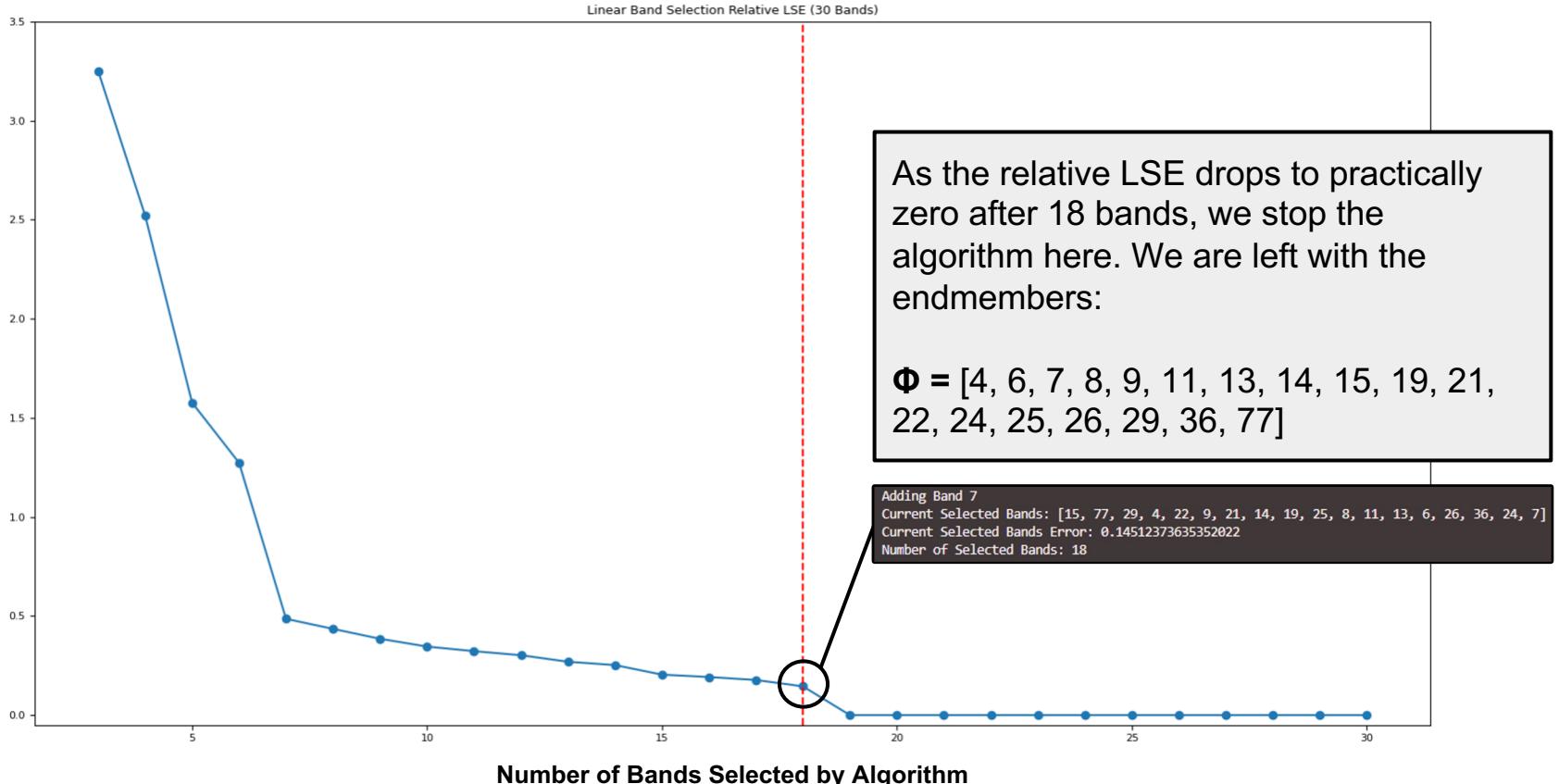
Start with an initial set of bands  
linear predictor model using the first band  
next band with the greatest correlation  
the model for  $\Phi$ , Repeat  
bands are found or error is less than  $\epsilon$



[\*]Du, Q., & Yang, H. (2008). Similarity-Based Unmixing for Hyperspectral Image Analysis. *IEEE Geoscience and Remote Sensing Letters*, 5(4), 504–508. doi:10.1109/LGRS.2008.2000019

# Linear Band Selection Results

LSE with respect to previous selected bands



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# Acknowledgments

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**We want to thank the following:**

**Fields Institute** for supporting us and this project

**Dr. Na Yu and Dr. You Liang** at TMU for their support and guidance in this project

**Dr. Yeni Yucel** at St Michaels for the motivation behind this project, the biomedical data we could show you today, and valuable suggestions from biomedical point of view

**Janak Bhanushali** at TMU for his help at the beginning of the project

# What We've Accomplished

We've created a **open-source** Python package for Biomedical Hyperspectral Imaging called **BHSIpy**:



## Linear Unmixing Methods:

Supervised Methods: CVXOPT, Gradient Descent, Active Set

Unsupervised Methods: UFCLSU

## Segmentation Methods:

Supervised: 3D-HyperUNet

Semi-Supervised: SAM, SIDSAM, JMSAM, NS3, K Means

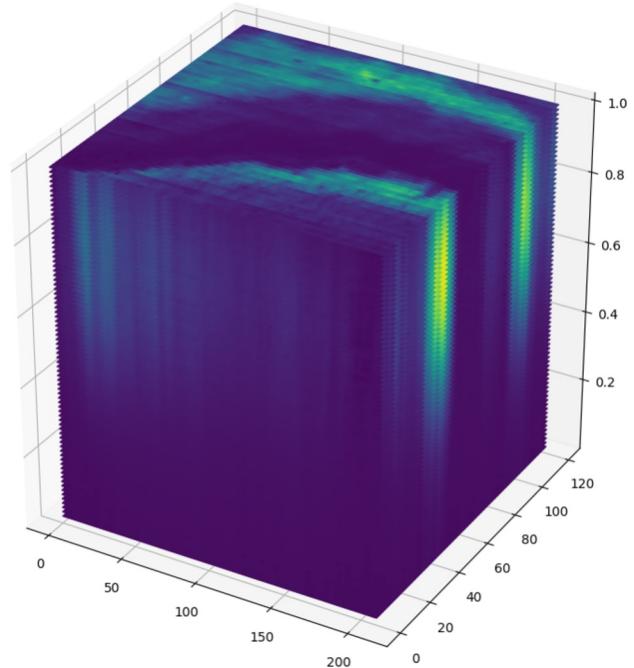
## Dimension Reduction:

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## Visualization Methods:

3D Plotting Method for Hyperspectral Cubes

General Layer Plots for Unmixing and Segmentation



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# Questions?

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