

# Spectral Unmixing and Segmentation of Biomedical Hyperspectral Images

Fields Undergraduate Summer Research Program Project 9



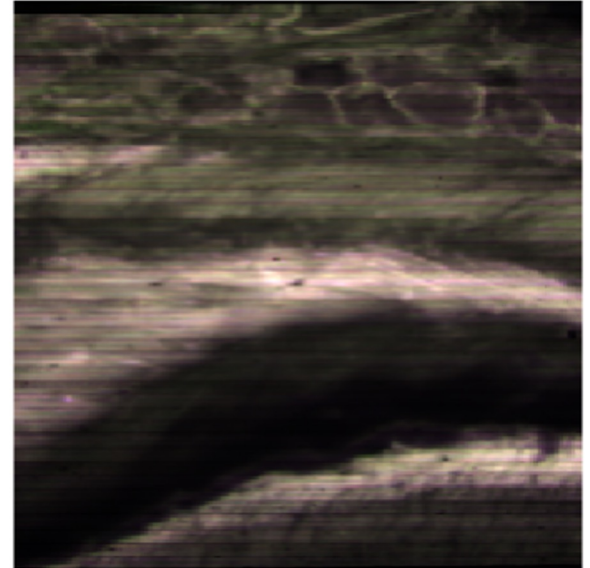
Xuanze (Charlie) Li, Aleksandar Popovic, Hannah Johnson

Supervised by Dr. Na Yu & Dr. You Liang



# Outline

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

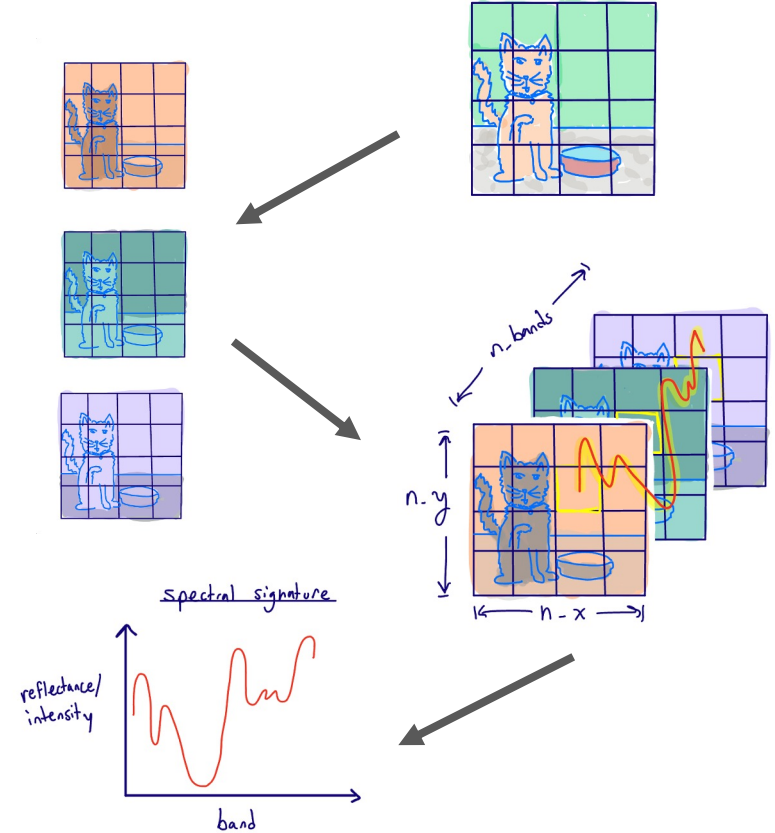


# Project Introduction

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

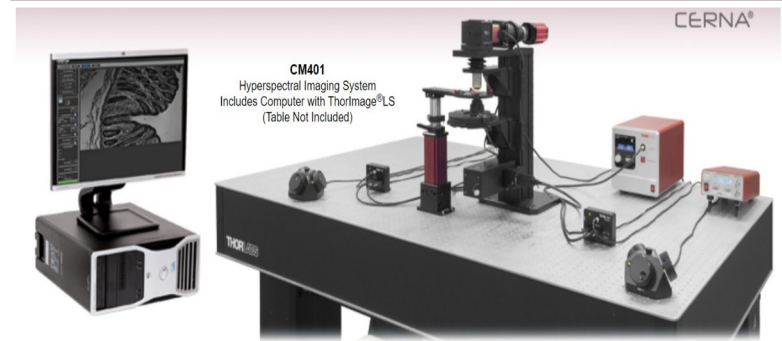
# What is Hyperspectral Imaging?

- Imaging camera measures light reflectance off of objects
- The camera separates the image by wavelength
- Different **endmembers**, or materials, in the 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



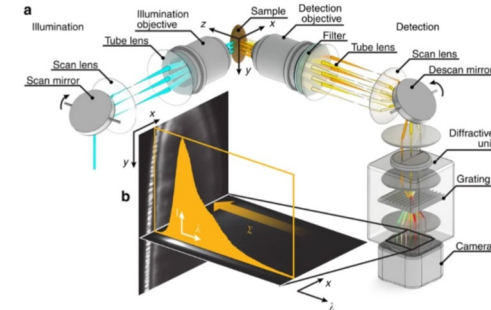
# Biomedical Hyperspectral Imaging

- 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/newgrouppage9.cfm?objectgroup\\_id=11095](https://www.thorlabs.com/newgrouppage9.cfm?objectgroup_id=11095)

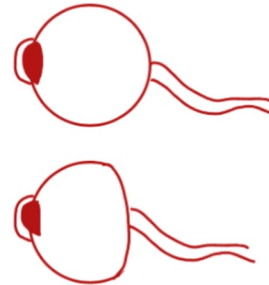
Figure 1: Hyperspectral SPIM setup and image formation.



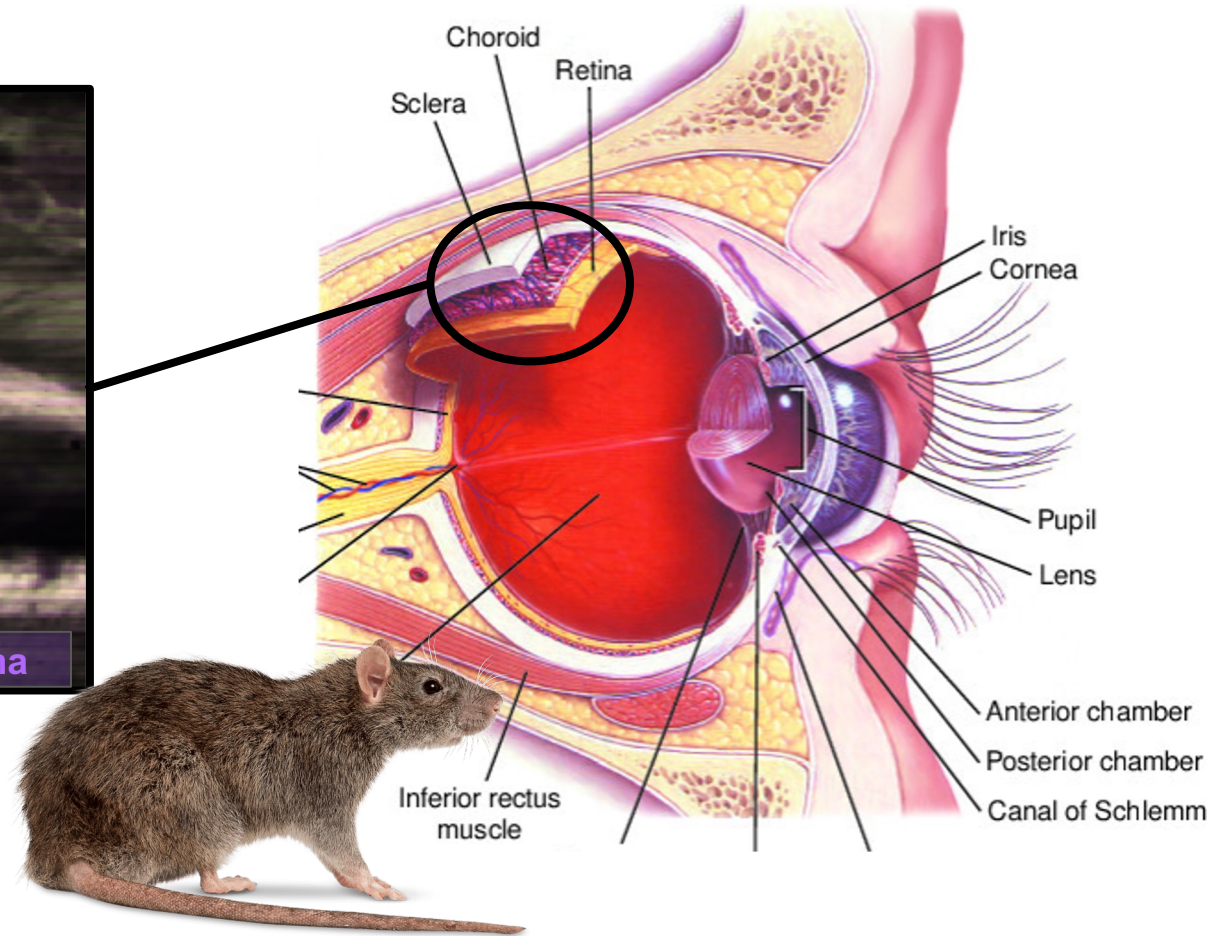
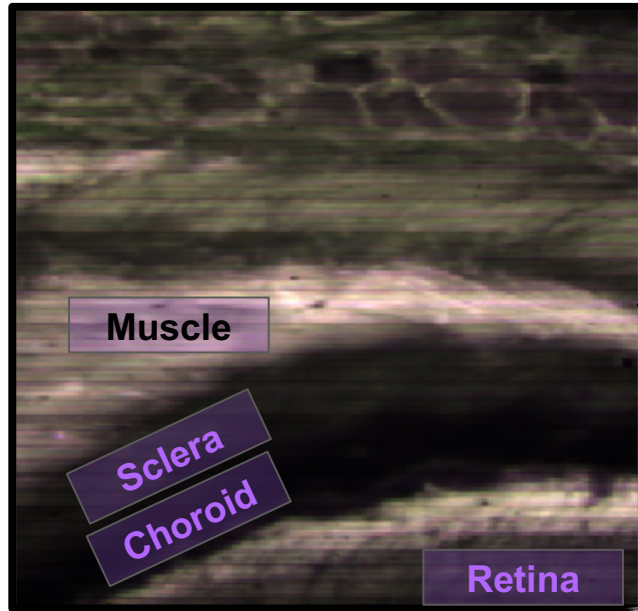
<https://www.nature.com/articles/ncomms8990>

# 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**

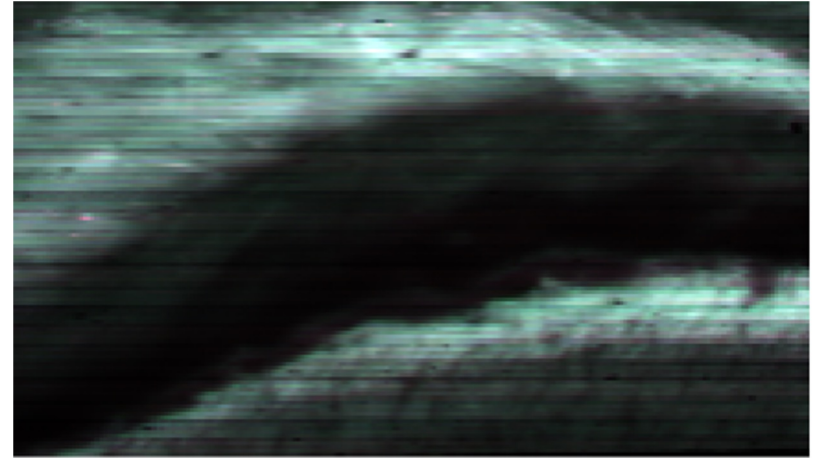
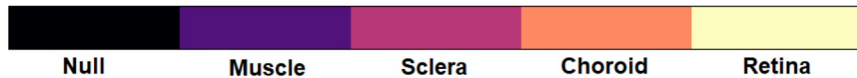
**Spectral Range:** 528 to 836 nm (NUV to NIR)

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

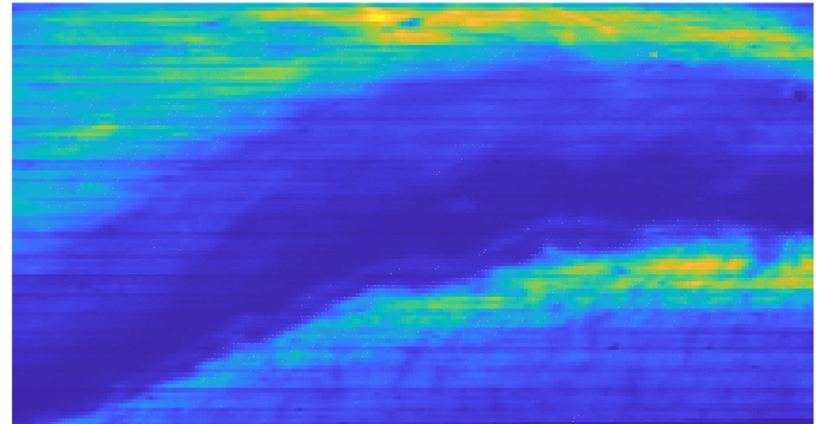
**Dataset Information:**

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

**Endmembers:**

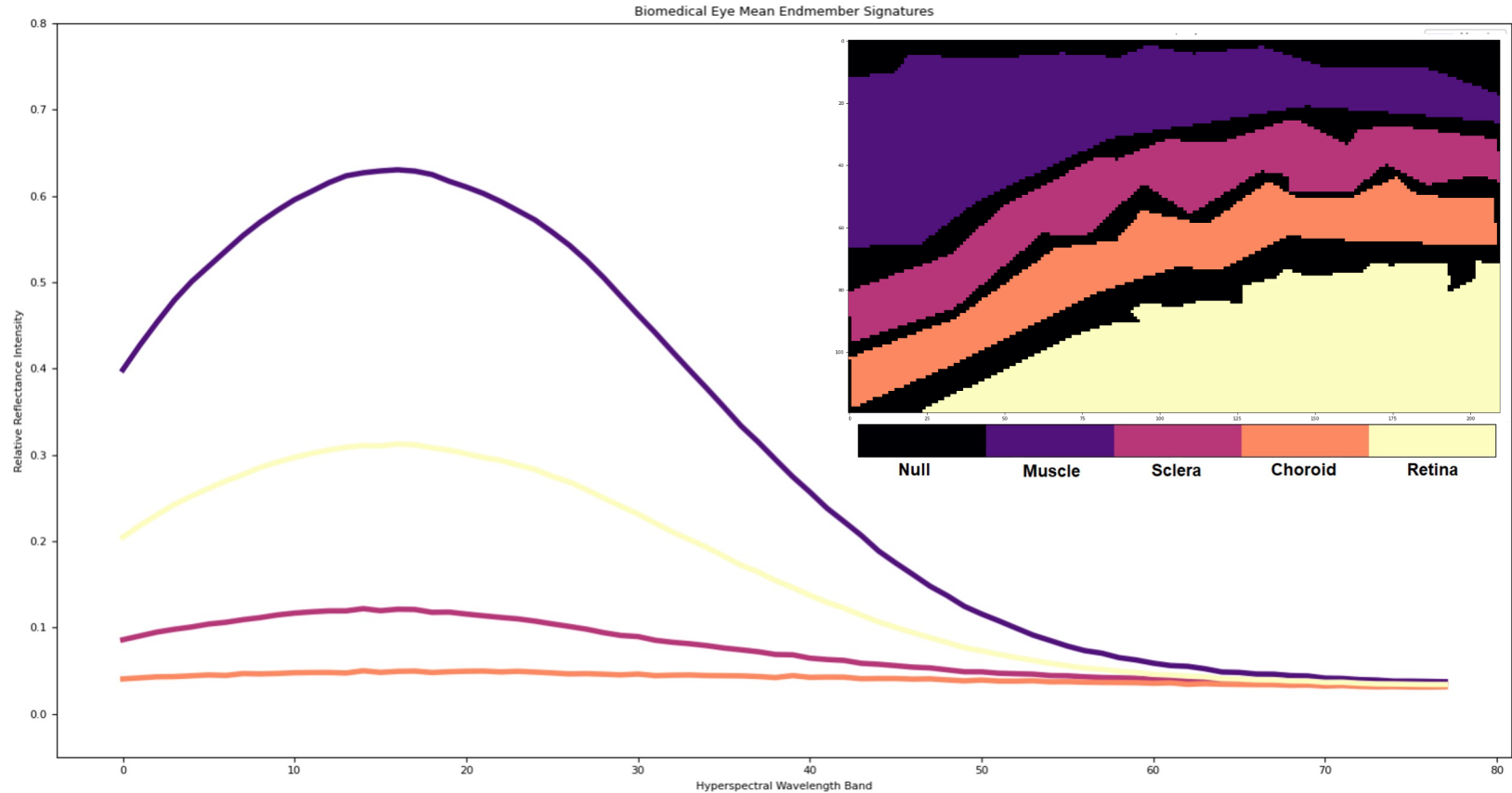


Eye Image Band 1





# 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

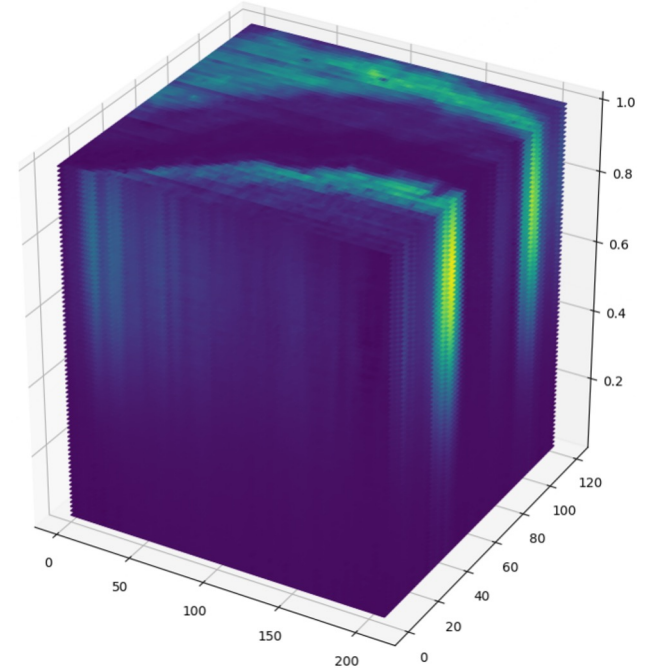
## Dimension Reduction:

Linear Band Selection and Principal Component Analysis

## Visualization Methods:

3D Plotting Method for Hyperspectral Cubes

General Layer Plots for Unmixing and Segmentation

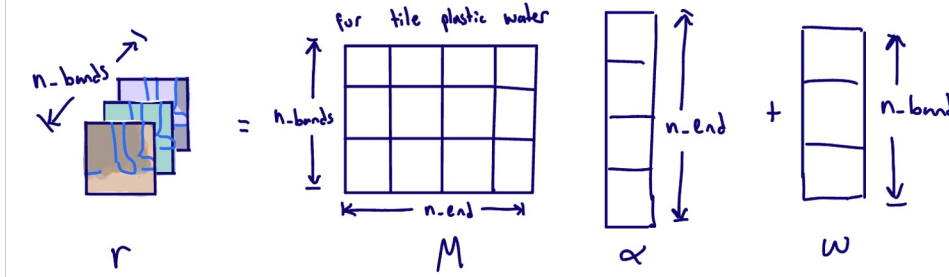


# Hyperspectral Unmixing

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

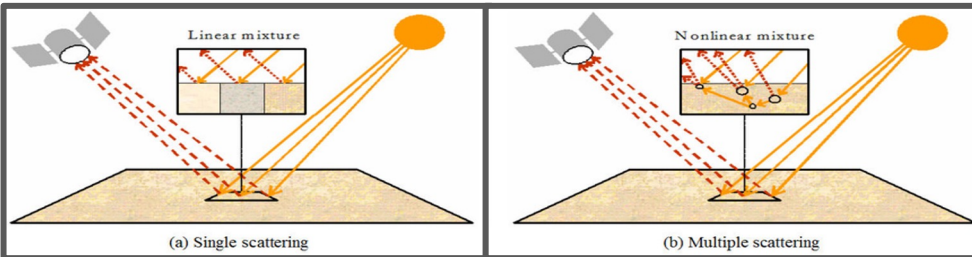
# Linear Hyperspectral Unmixing

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



**Formally:**

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



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\*\*:

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

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

# 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:** *lsqnonneg*

**Python:** *nnls*

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



[\*] Lawson, C. L. and R. J. Hanson. *Solving Least-Squares Problems*. Upper Saddle River, NJ: Prentice Hall. 1974. Chapter 23, p. 161.

## Gradient Descent Methods

- Applying Gradient Descent to Solve FCI SII Problems\*

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

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

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

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

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

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

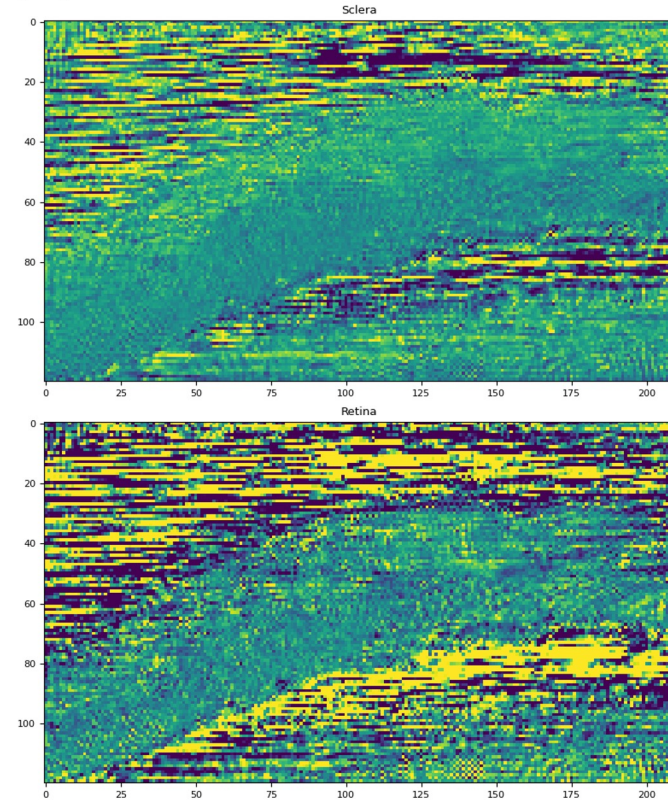
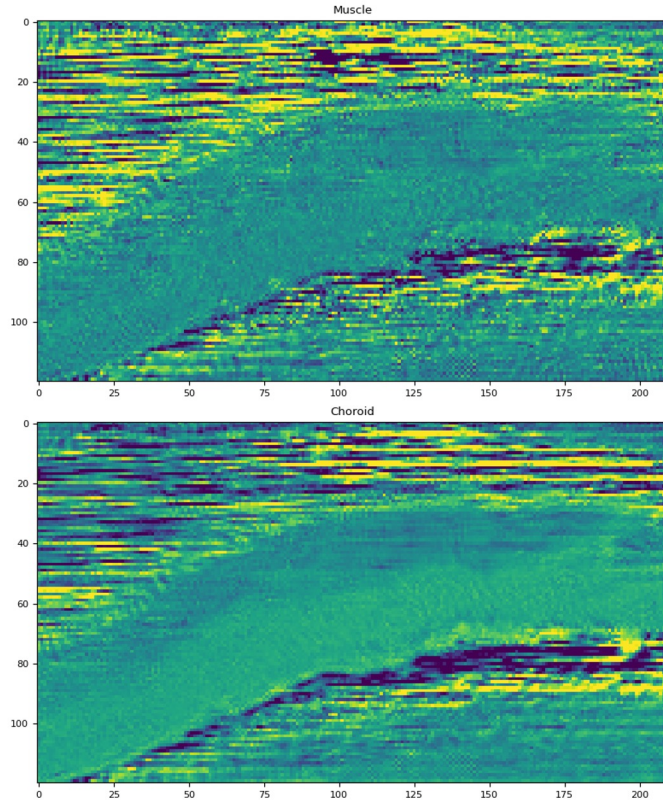
$$+ \mu \text{diag}\{\alpha(k)\} [\nabla_{\alpha} J(\alpha) - \mathbf{1} \nabla_{\alpha} J(\alpha)^{\top} \alpha(k)]$$

[\*] Jie Chen, Cédric Richard, Henri Lantéri, Céline Theys, Paul Honeine. A Gradient Based Method for Fully Constrained Least-Squares Unmixing of Hyperspectral Images. Proc. IEEE workshop on Statistical Signal Processing (SSP). 2011. Nice, France. pp.301-304.

# Unmixing Results

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

Unconstrained Least Squares Unmixing  
Biomedical Rat Eye Image



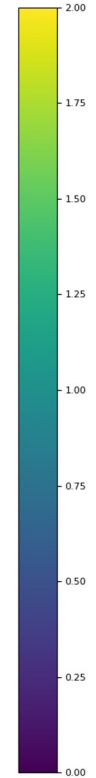
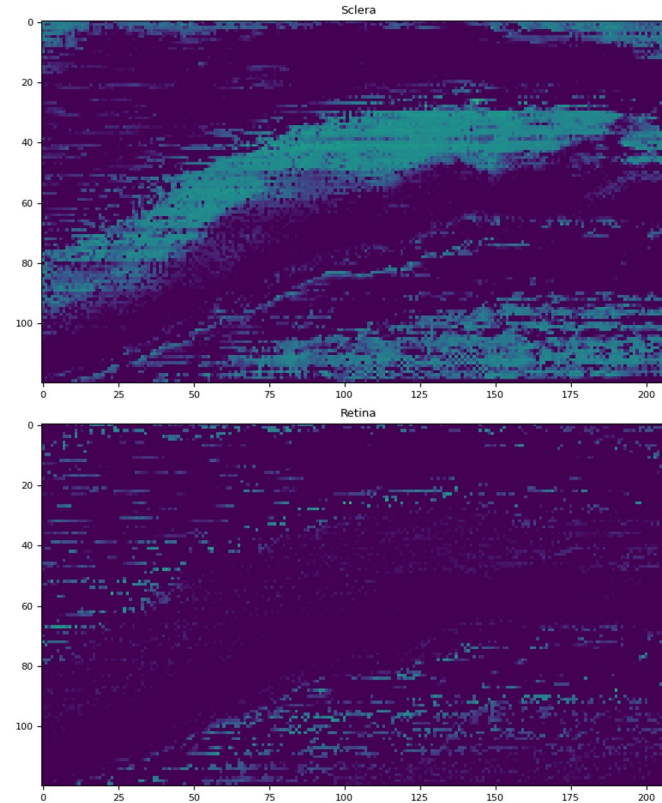
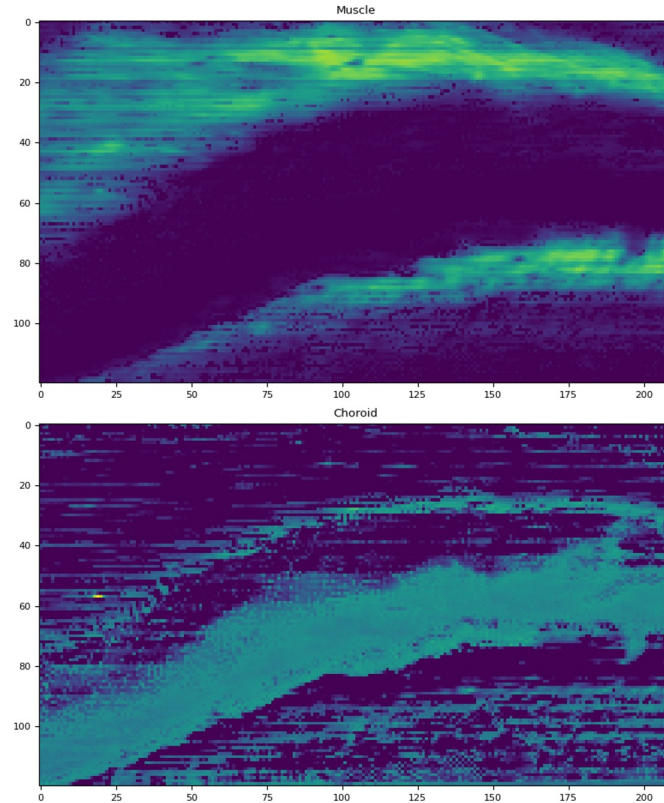
Pixelwise Average LSE: 0.0011

Running Time: 0.074 s

# Unmixing Results

$$\alpha_i \geq 0$$

Constrained Least Squares Unmixing  
Biomedical Rat Eye Image



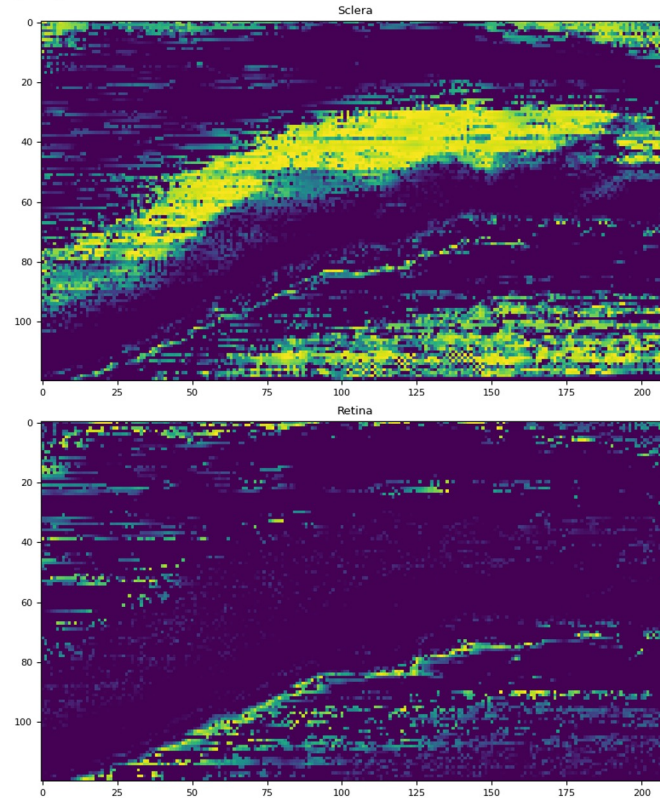
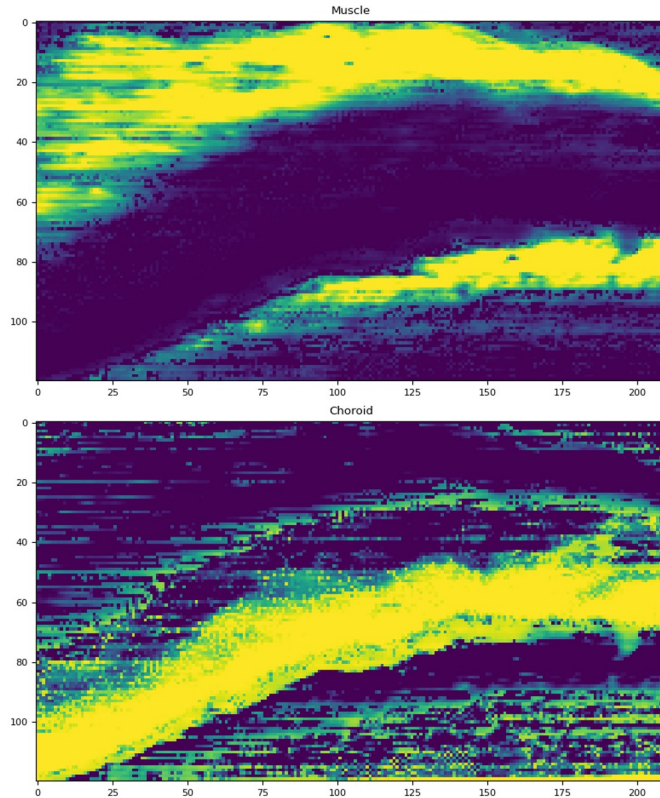
Pixelwise Average LSE: 0.0062

Running Time: 16.583 s

# Unmixing Results

$$\alpha_i \geq 0 \quad \sum a_i = 1$$

Fully Constrained Least Squares Unmixing  
Biomedical Rat Eye Image



Pixelwise Average LSE: 0.1532

Running Time: 17.808 s



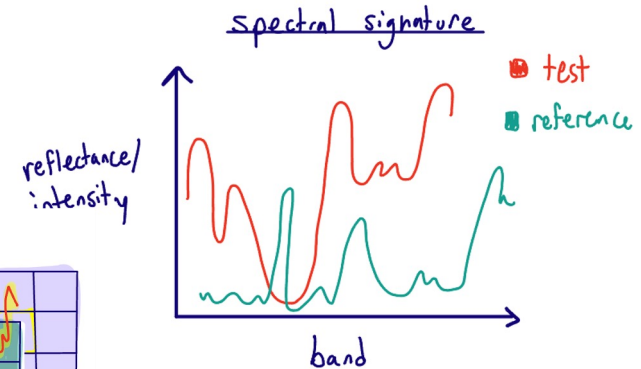
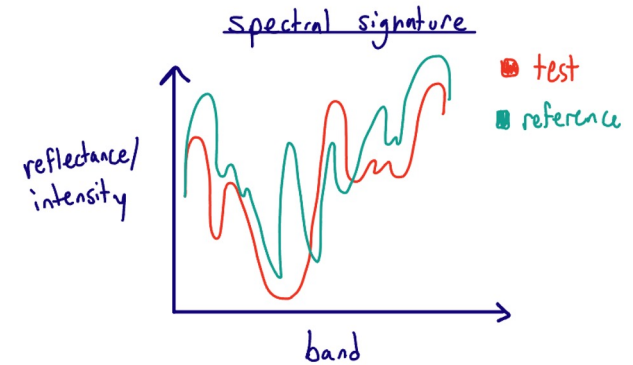
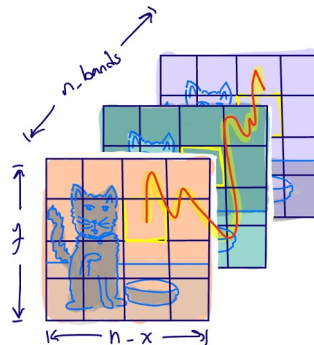
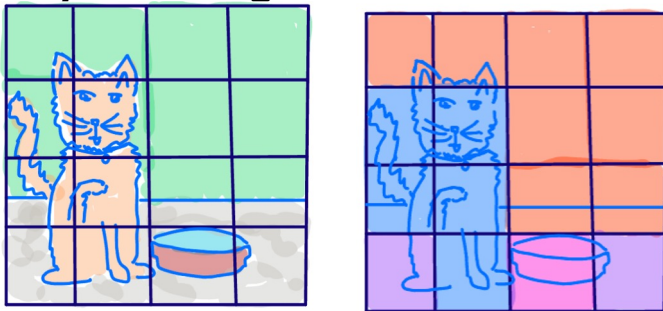
# Hyperspectral Segmentation

- ◆ 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

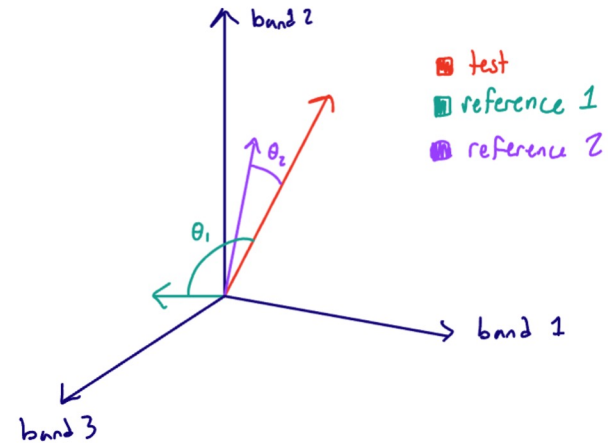
Mixed

Measures:

- SID-SAM
- JMSAM
- NS3

Spectral signature array

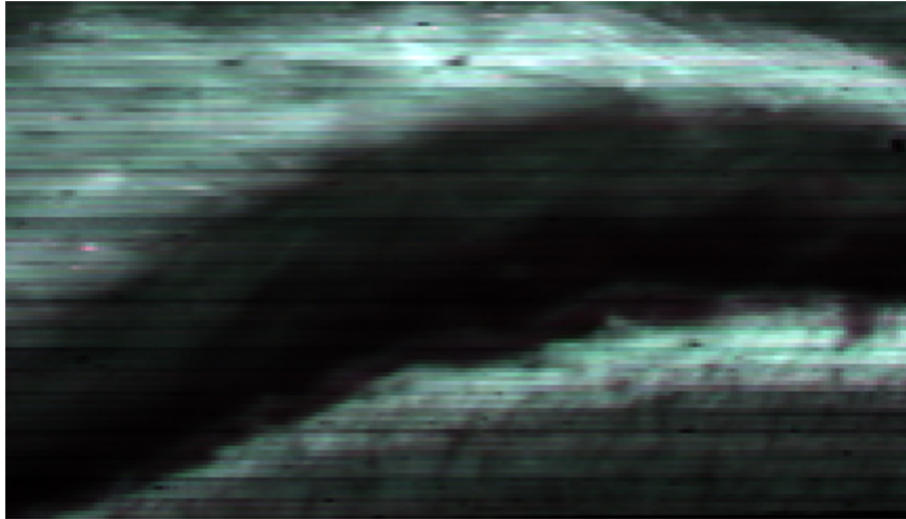
band	1	2	3	4	5	6	7	8
intensity	78	67	46	49	50	80	92	102



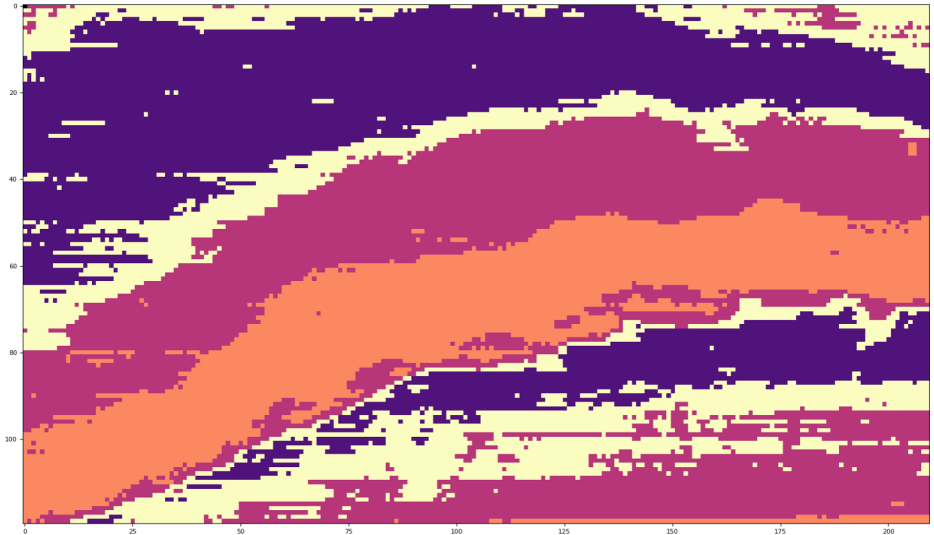
# Spectral Angle Mapper Segmentation

Angle

Original Image



Segmented Image



Null

Muscle

Sclera

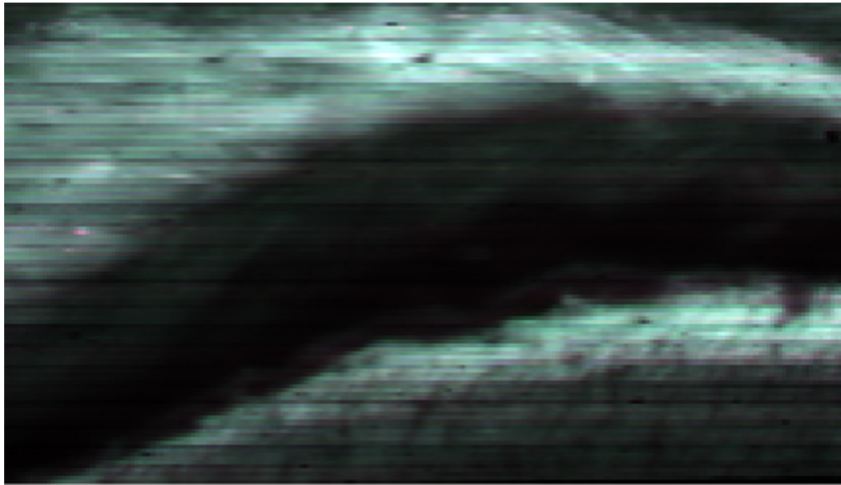
Choroid

Retina

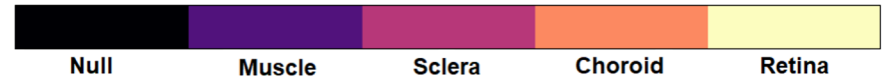
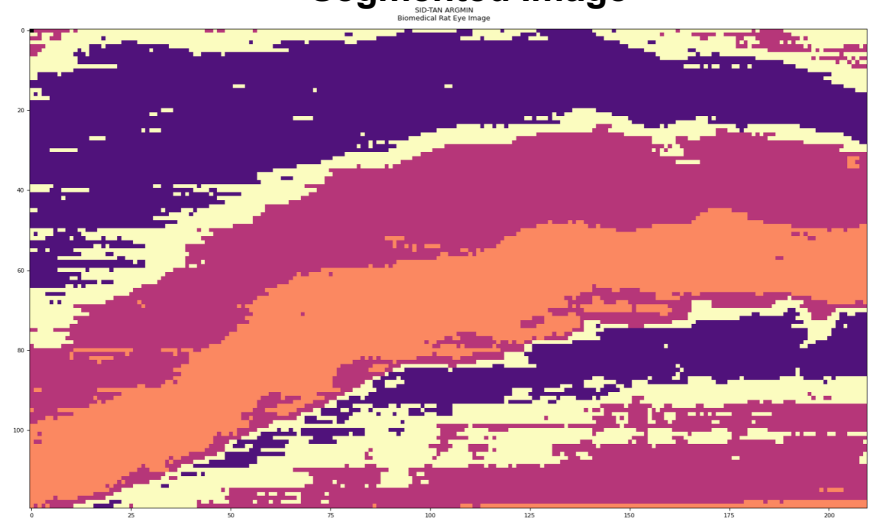
# SIDSAM TAN Segmentation

Angle, Probabilistic

Original Image



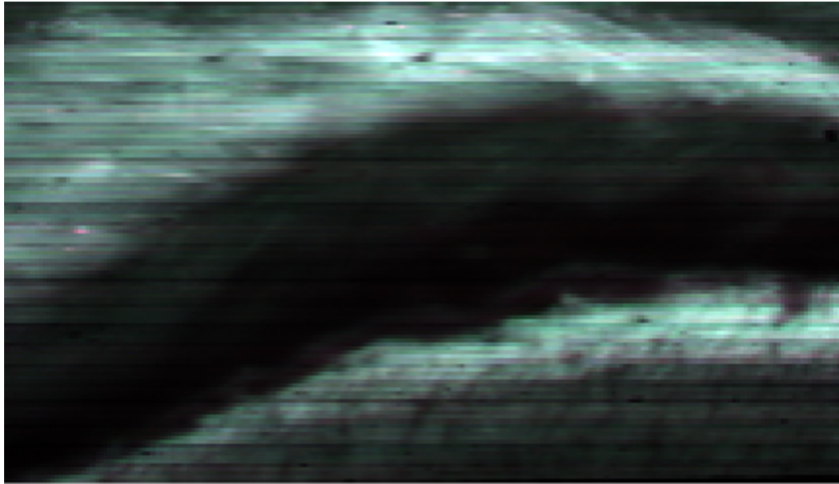
Segmented Image



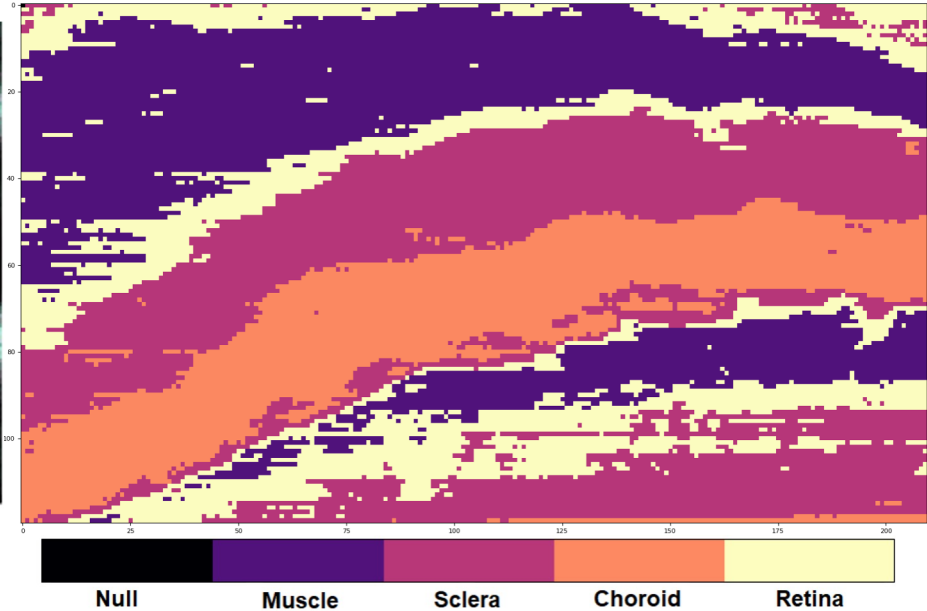
# JMSAM TAN Segmentation

Angle, Distance

Original Image



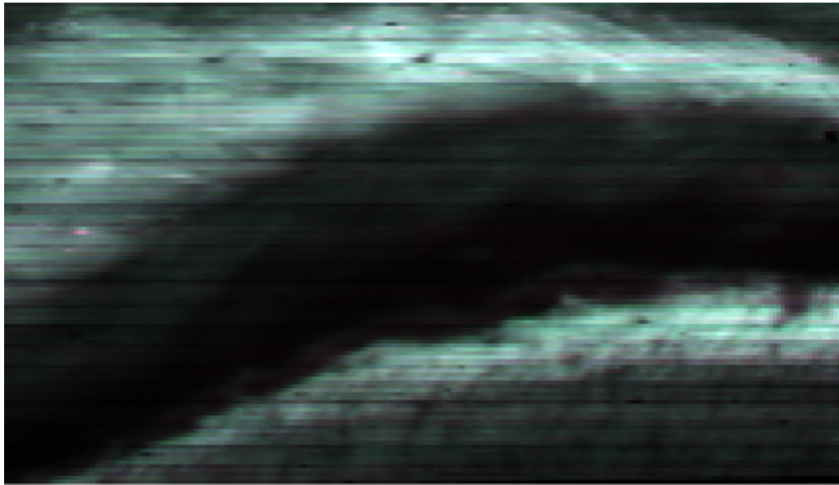
Segmented Image



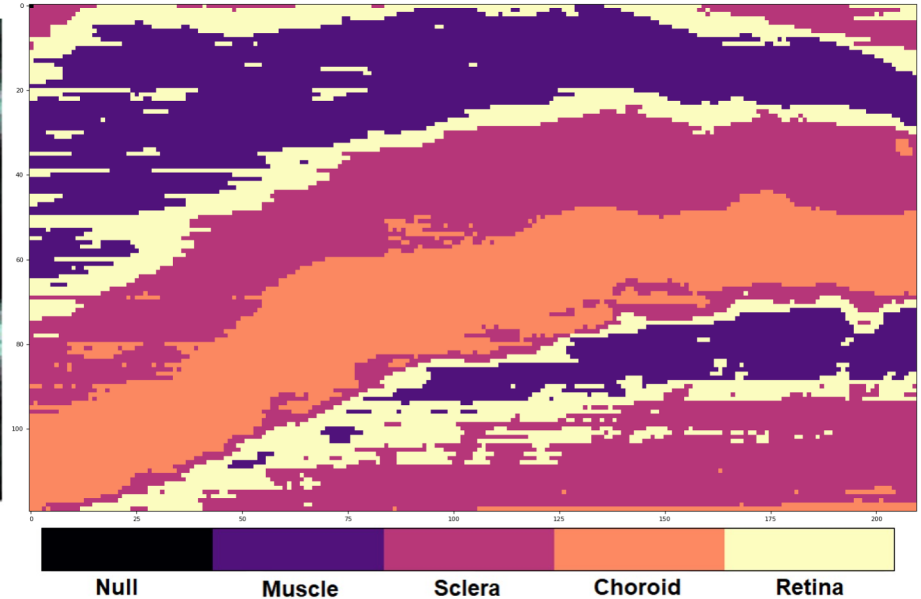
# NS3 Segmentation

Angle, Distance

Original Image



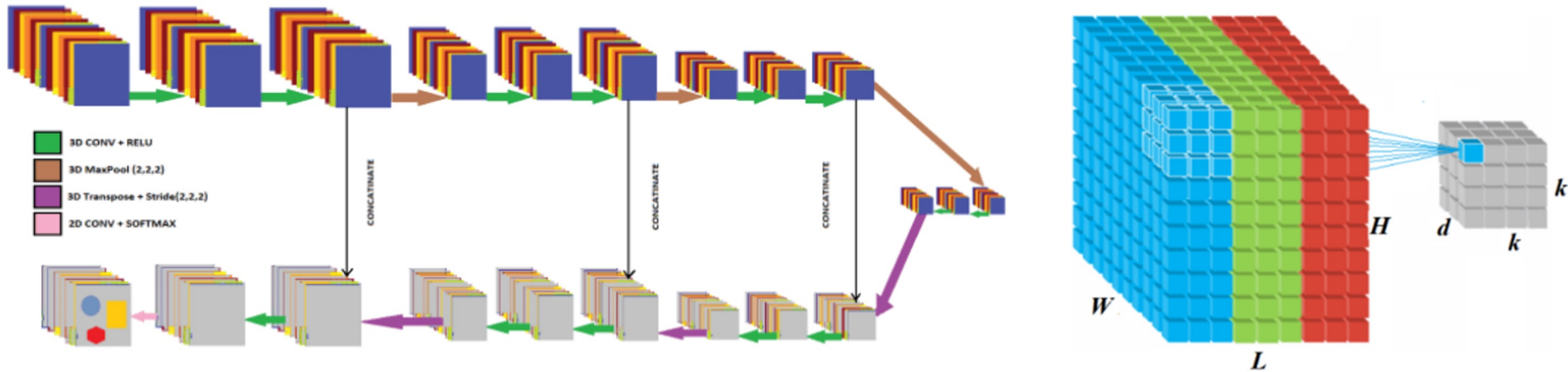
Segmented Image



# 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

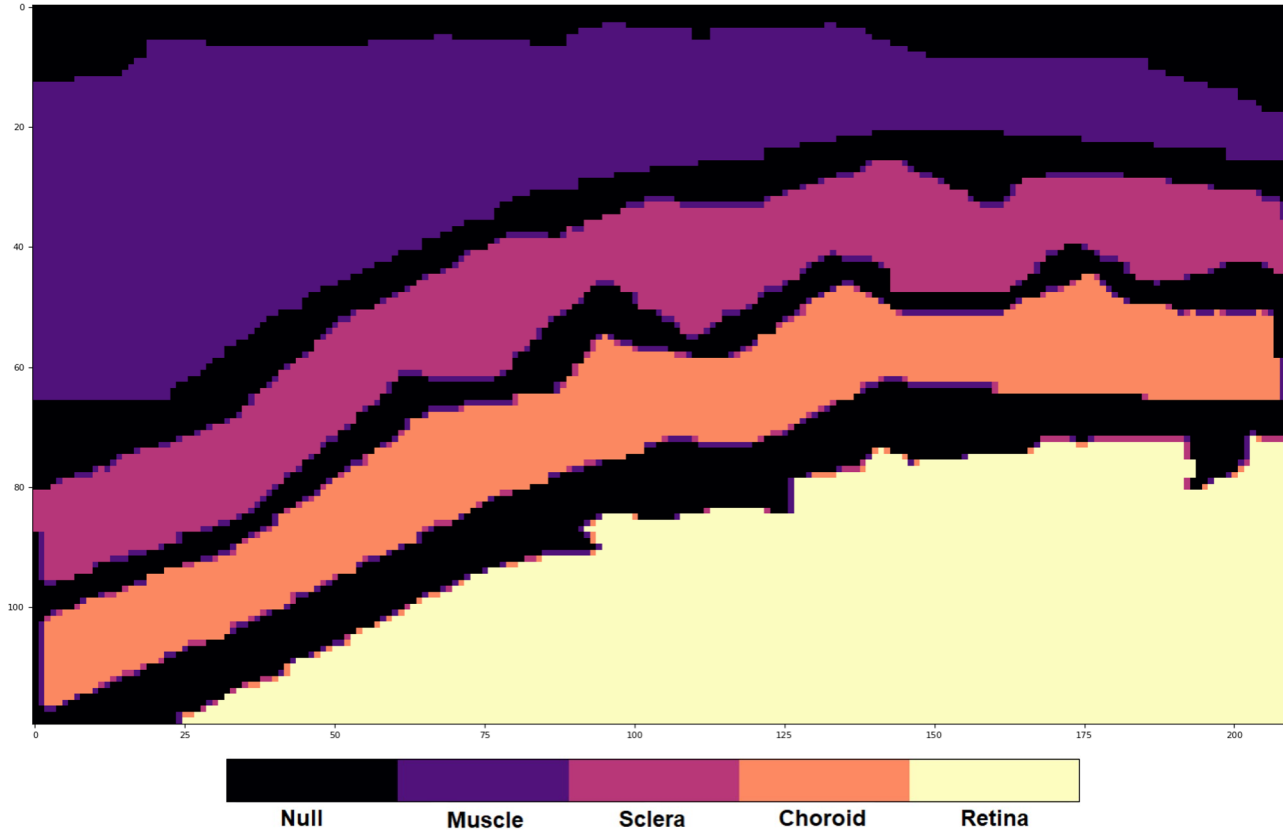




# Deep Learning Segmentation Results

```
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



IoU: 0.8054  
Loss: 0.0386

# Dimension Reduction

- ◆ Linear Predictor Band Selection

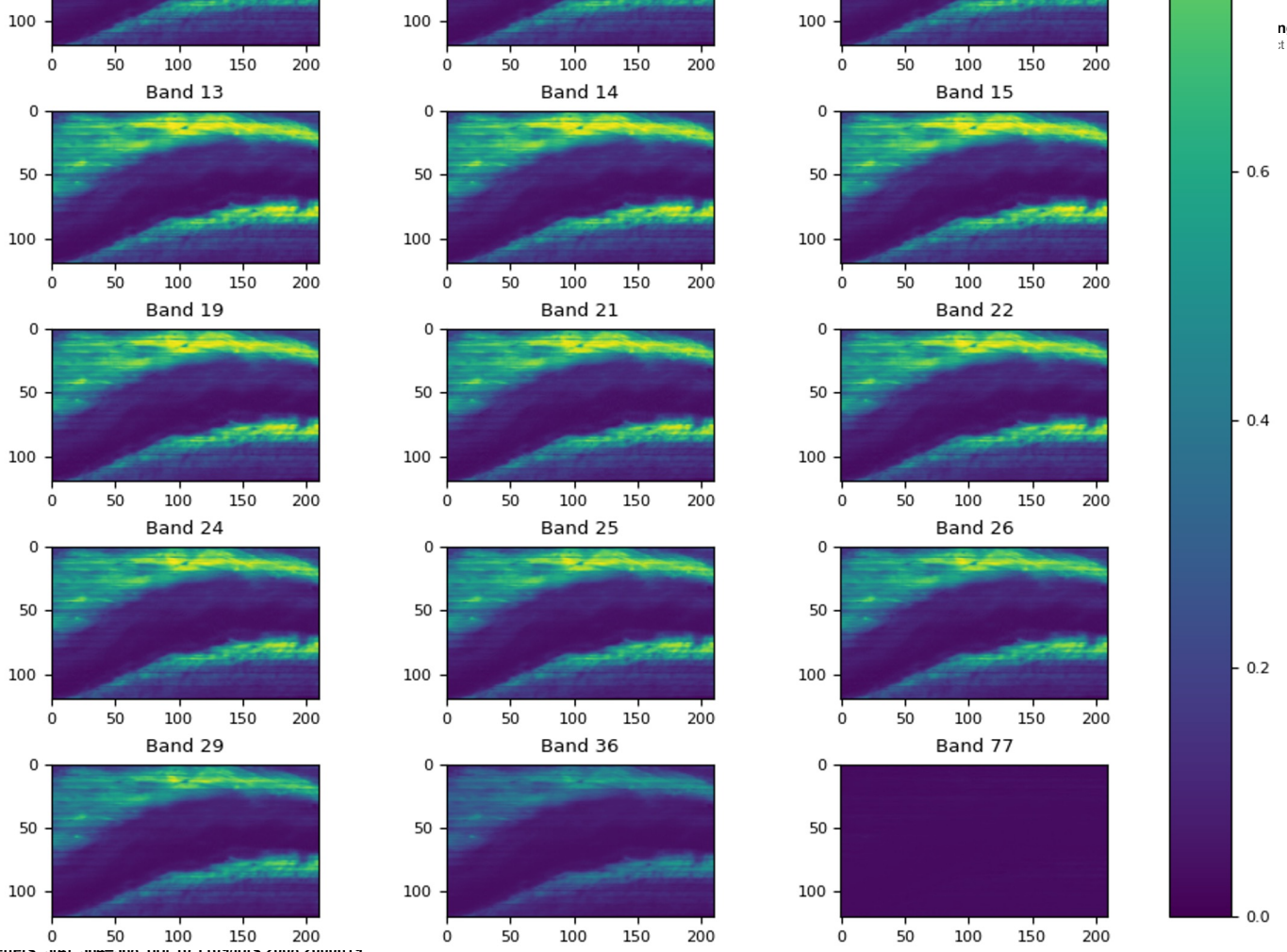
# Linear Bands

## Motivation:

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

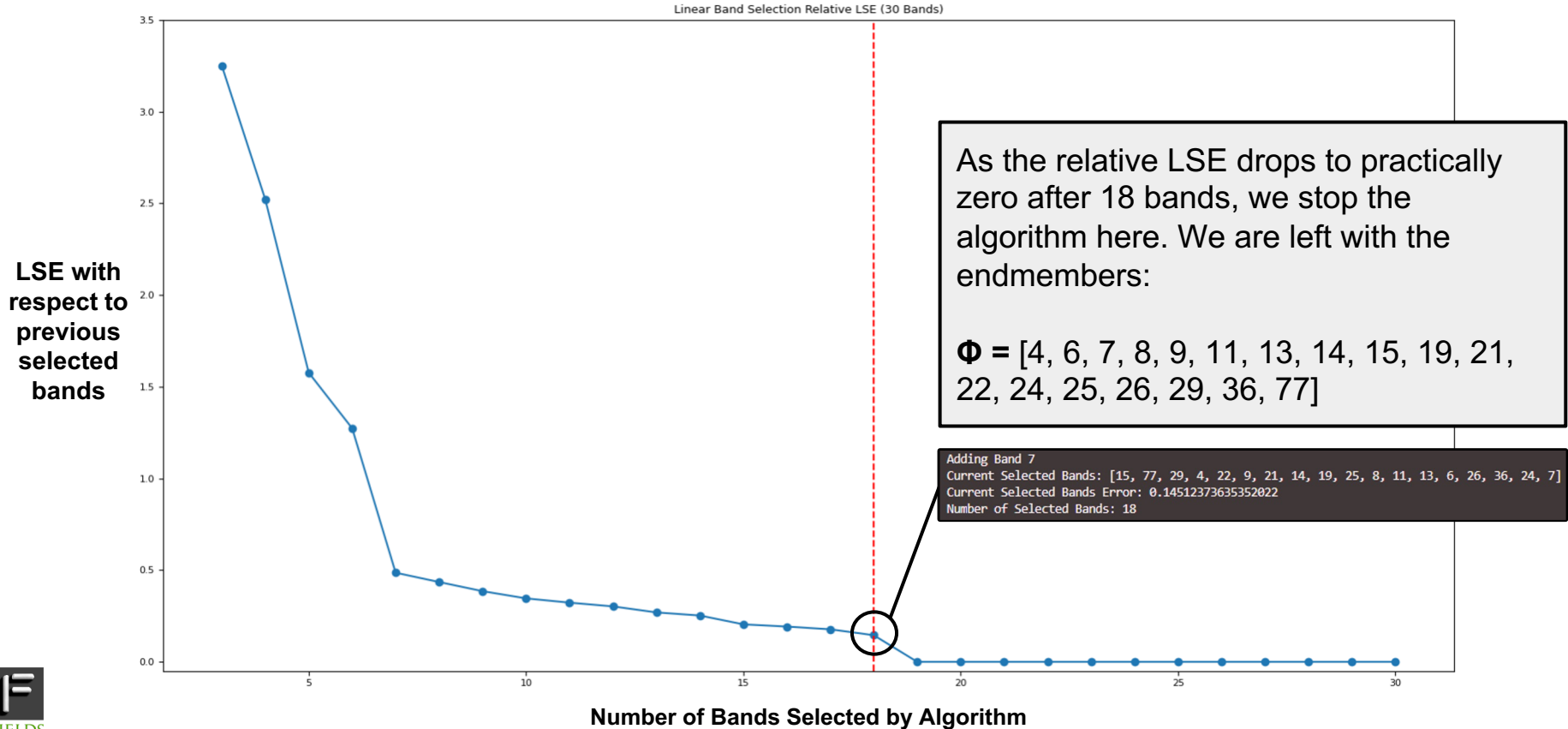
## Algorithm\*:

Start with an initial set of linear predictor models. Use the next band with the greatest error to update the model for  $\Phi$ . Repeat until no more bands are found or error is small.



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

# Linear Band Selection Results



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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

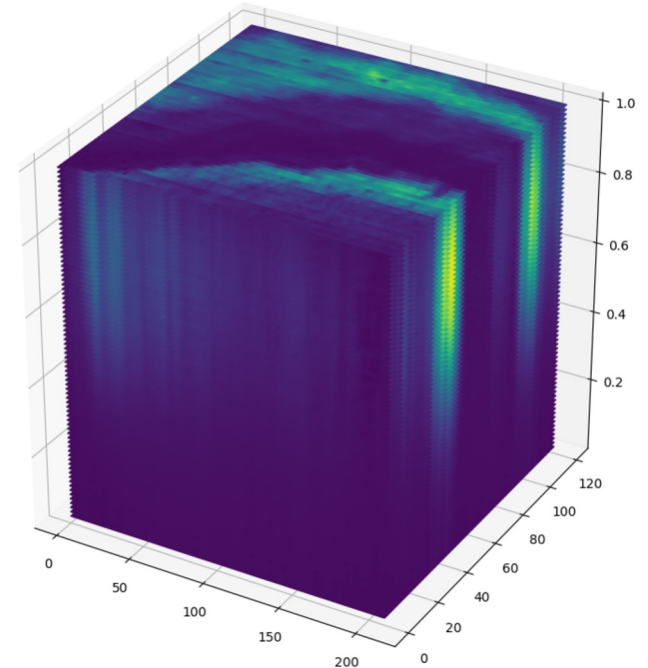
## Dimension Reduction:

Linear Band Selection and Principal Component Analysis

## Visualization Methods:

3D Plotting Method for Hyperspectral Cubes

General Layer Plots for Unmixing and Segmentation



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# Questions?

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