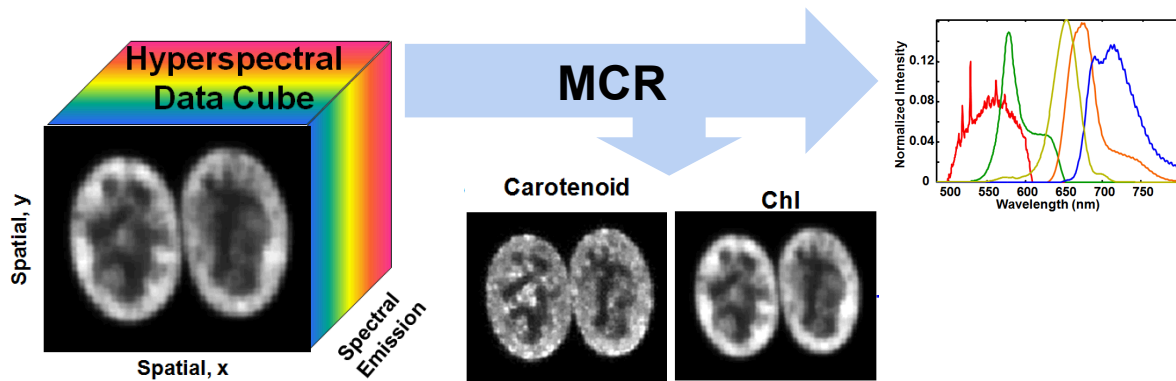


Adapted from Howard Vindin - Own work, CC BY-SA 4.0,  
<https://commons.wikimedia.org/w/index.php?curid=40722030>

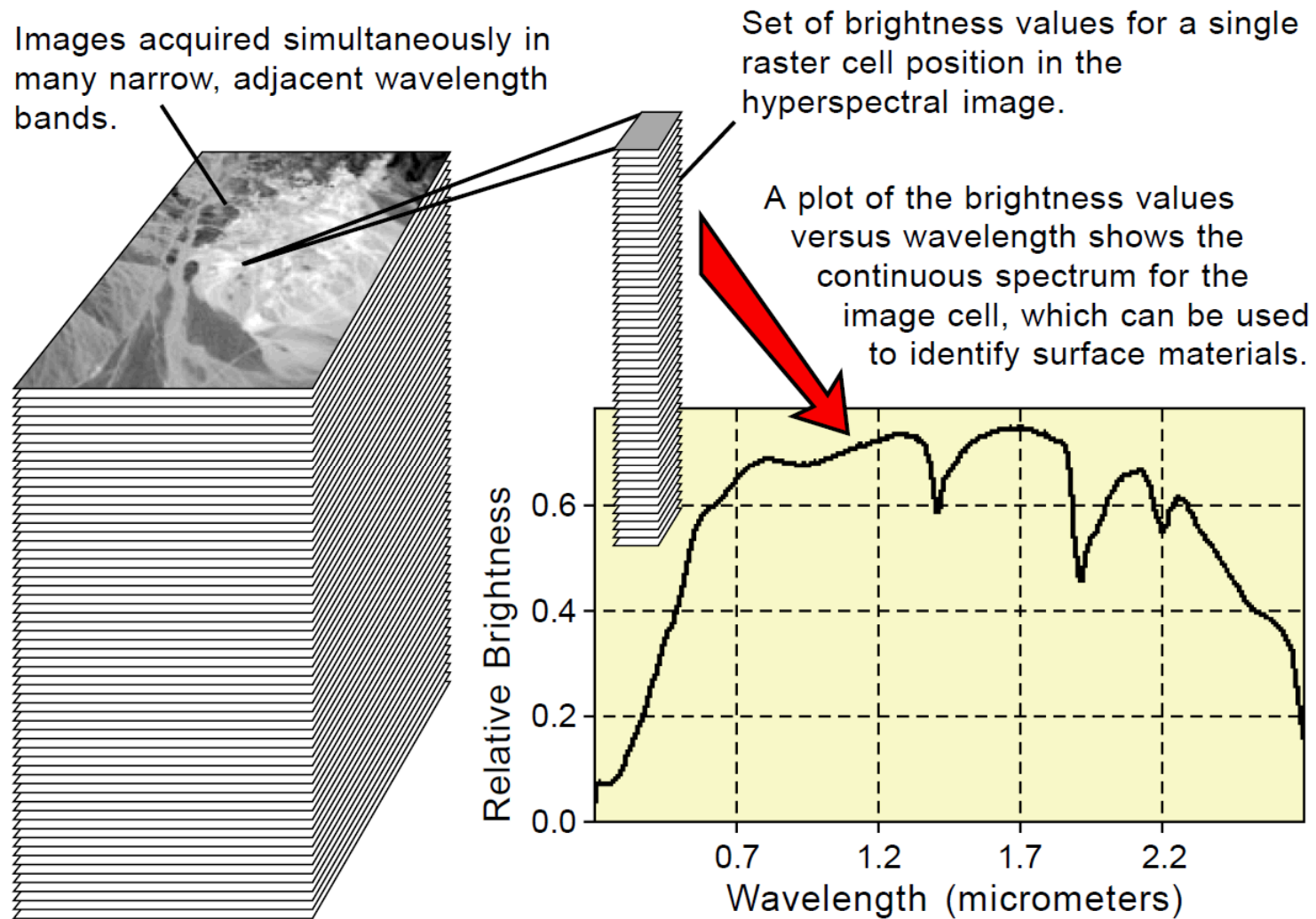


# Hyperspectral Super-Resolution Imaging and Data Analysis

Stephen M. Anthony

- **Why hyperspectral?** – What is hyperspectral imaging and what are its benefits?
- **Sandia's hyperspectral microscopes** – What are the systems I typically work with?
- **Analyzing hyperspectral data** – What can multivariate curve resolution (MCR) do for you?
- **Improving MCR** – Ongoing work to improve its capabilities.
- **Trilinear Data** – How to leverage additional information.

# Introduction to Hyperspectral Imaging



Smith, R. B. (2012) Introduction to Hyperspectral Imaging. [Microimages](http://www.microimages.com/documentation/Tutorials/hyprspect.pdf)  
<http://www.microimages.com/documentation/Tutorials/hyprspect.pdf>

# Why Use Hyperspectral Imaging?

Conventional Fluorescence Image

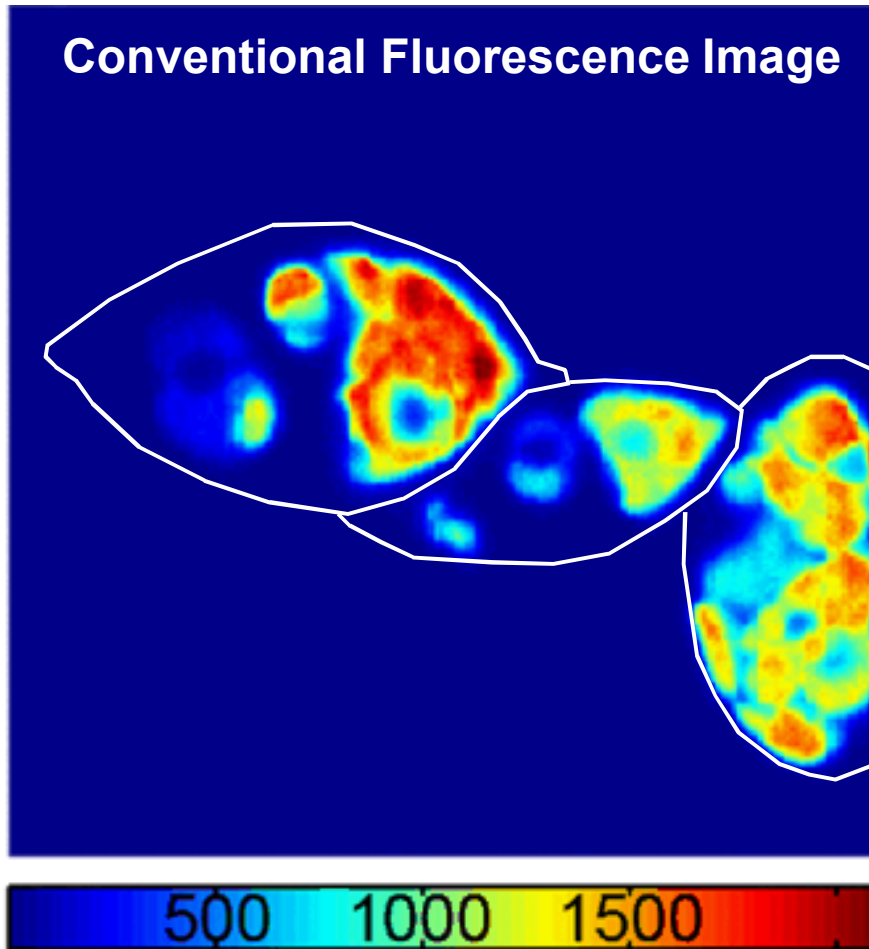
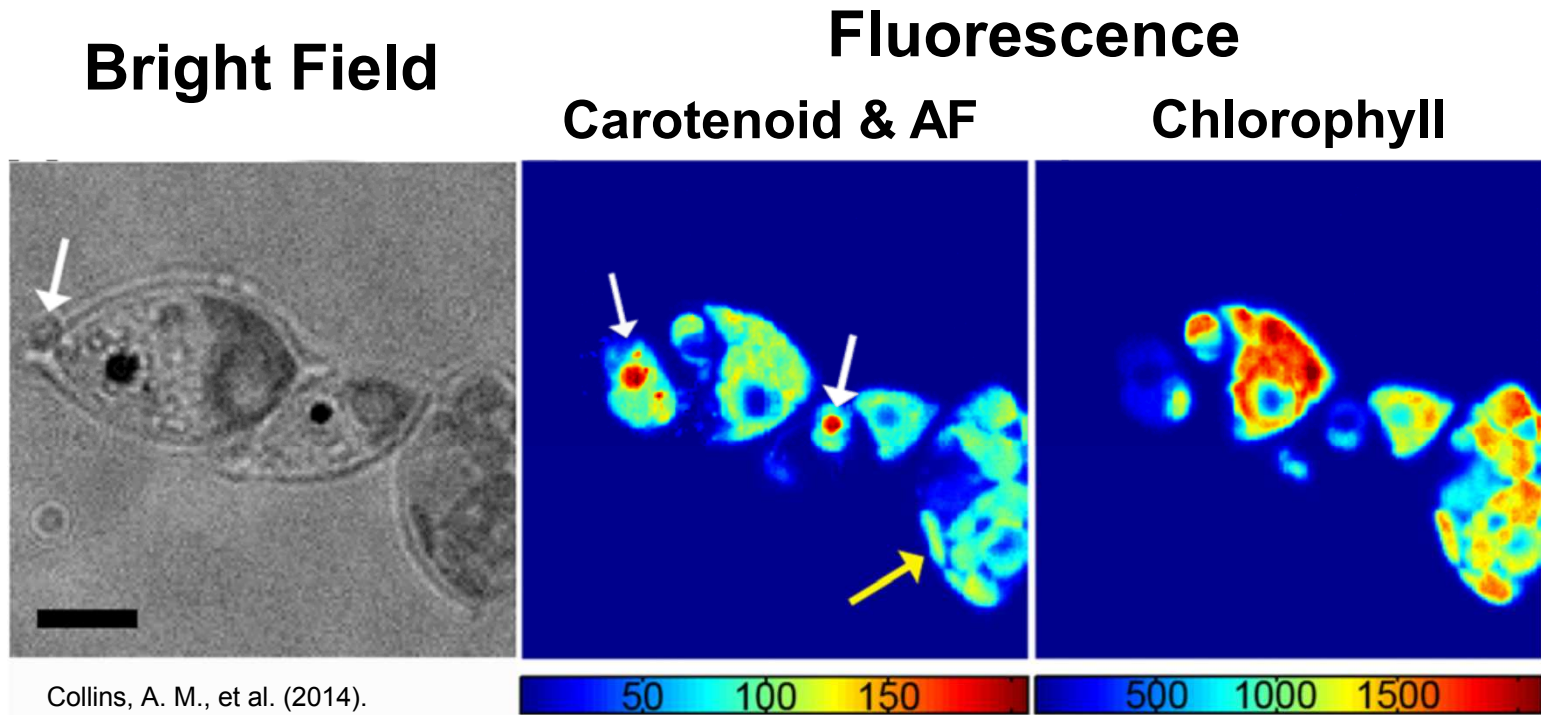


Image of the endogenous fluorescence from *S. dimorphus* (algae) undergoing parasitic infection by *A. protococcarum*.

- Approximate cell borders are hand-drawn in white.
- Two of the cells contain parasitic vacuoles.
- **Can you spot the parasitic vacuoles?**

Adapted from Collins, A. M., et al. (2014). "Host Cell Pigmentation in *Scenedesmus dimorphus* as a Beacon for Nascent Parasite Infection." *Biotechnology and Bioengineering* **111**(9): 1748-1757.

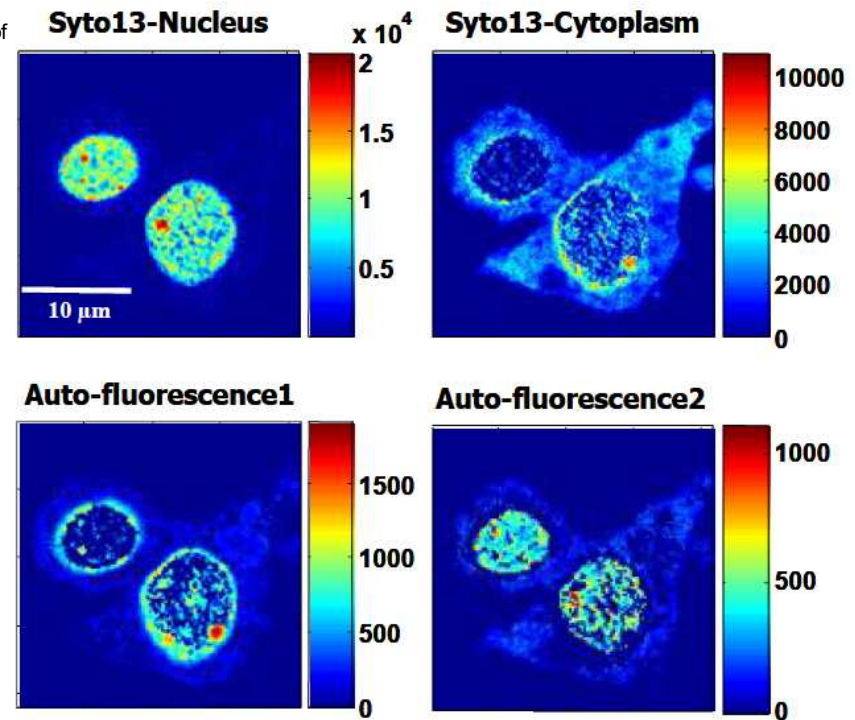
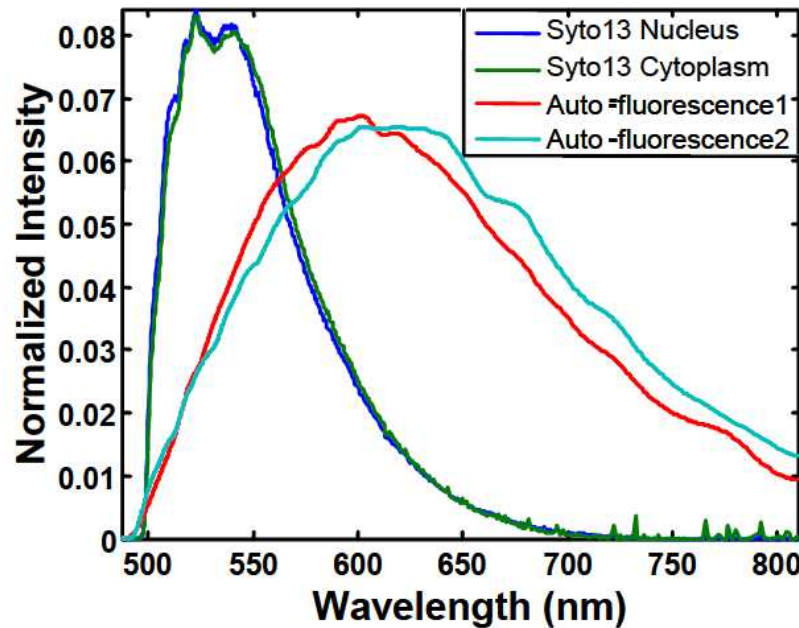
# Why Use Hyperspectral Imaging?



- Parasitic vacuoles (white arrows) are easily spotted using the combined carotenoid and autofluorescence signal.
- Spotting them is nearly impossible when examining all the fluorescence together as the chlorophyll signal dominates.
- **Hyperspectral imaging reveals otherwise hidden features.**

# Why multispectral is not enough

Haaland, D. M., et al. (2007). "Hyperspectral confocal fluorescence imaging of cells" *Next-Generation Spectroscopic Technologies* 6765: 76509-76509.



Left) Fluorescence spectra for two Syto 13 and two autofluorescence emission components. Right) Relative concentration of the components' spatial distributions in mouse macrophage cells (Raw 264.7).

- Multispectral imaging (e.g. filter-based microscopes) would only distinguish Syto 13 from autofluorescence – two components.
- **Hyperspectral imaging can distinguish nearly identical spectra.**

# Hyperspectral Imaging Applications

- **Fluorescence or Raman microscopy – cell signaling**
- **Agriculture (satellite or drone-based) – monitoring crop locations (poppy fields), crop health, disease outbreak**
- **Chemical detection – Detect airborne chemical hazards at ppm levels up to 5 km away**
- **Mineralogy – disturbed ground indicative of improvised explosive devices**

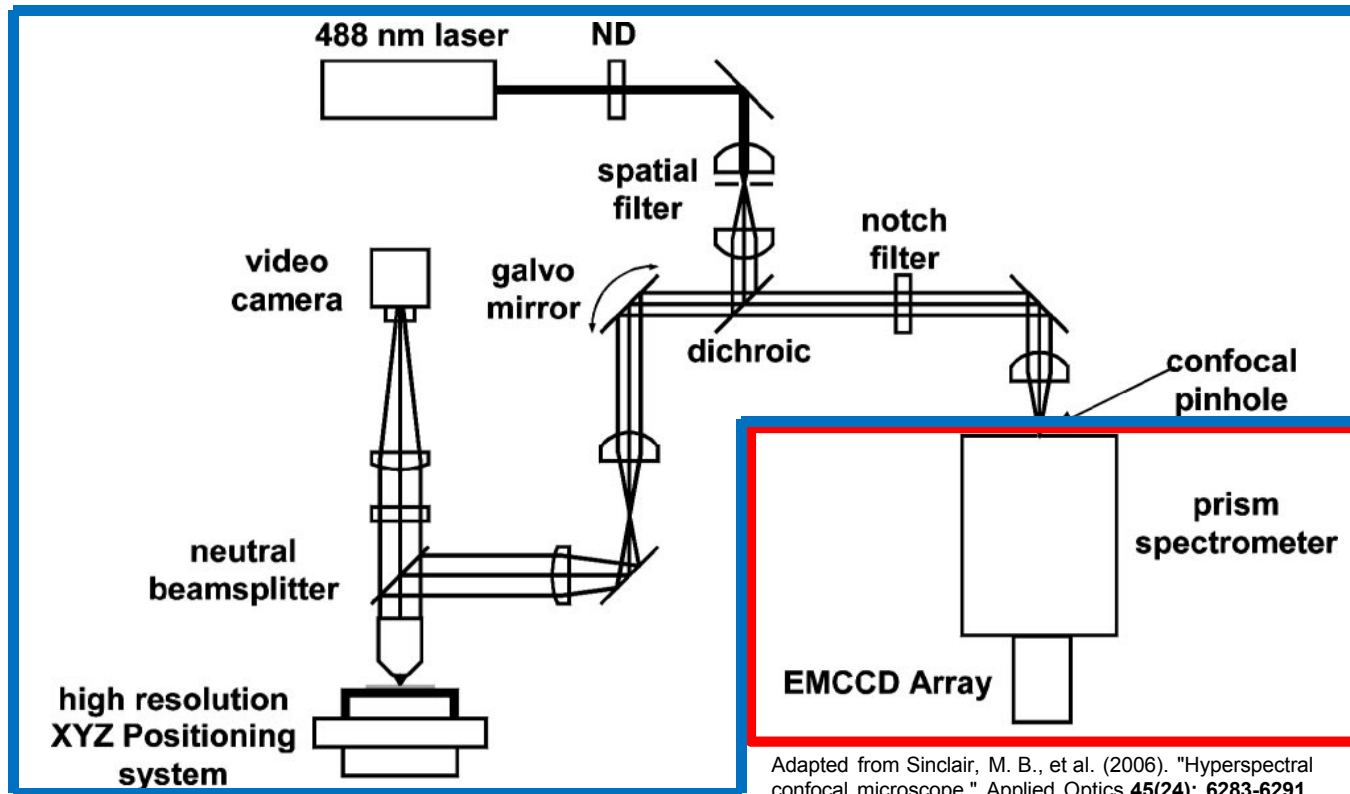


# Outline

- **Why hyperspectral?** – What is hyperspectral imaging and what are its benefits?
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# How to Build a Hyperspectral Microscope

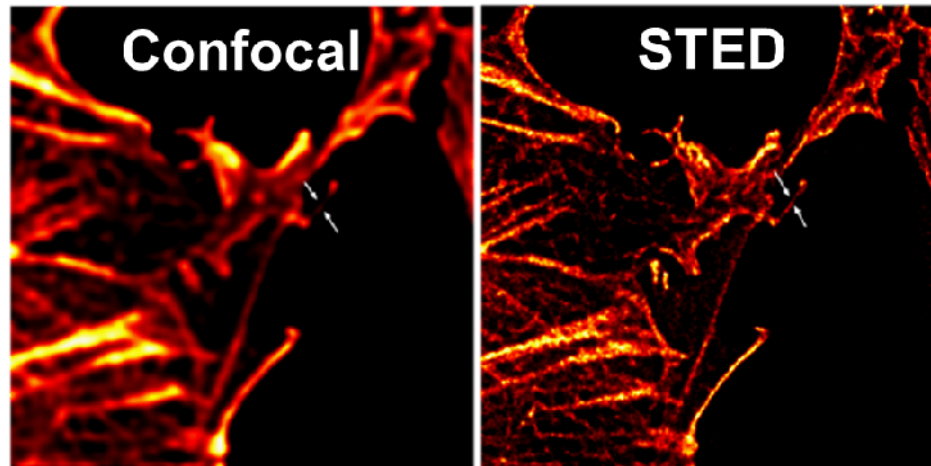


Schematic diagram of Sandia's hyperspectral confocal microscope

**Hyperspectral Confocal Microscope =**  
**Confocal Microscope + Spectrometer**

# Hyperspectral STED Microscope

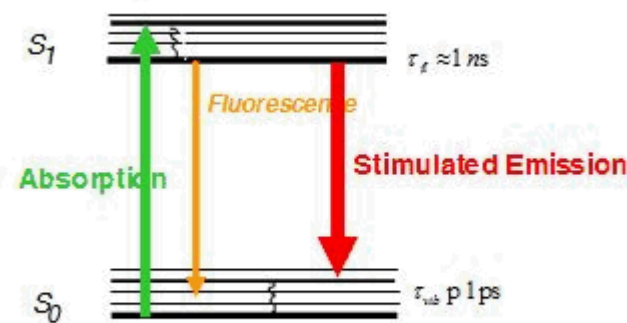
- Stimulated Emission Depletion (STED) is a super-resolution microscopy technique
  - Super-resolution microscopy won the 2014 Nobel prize in chemistry
- STED dramatically improves the spatial resolution ( $\sim 30$  nm typical,  $< 3$  nm reported)



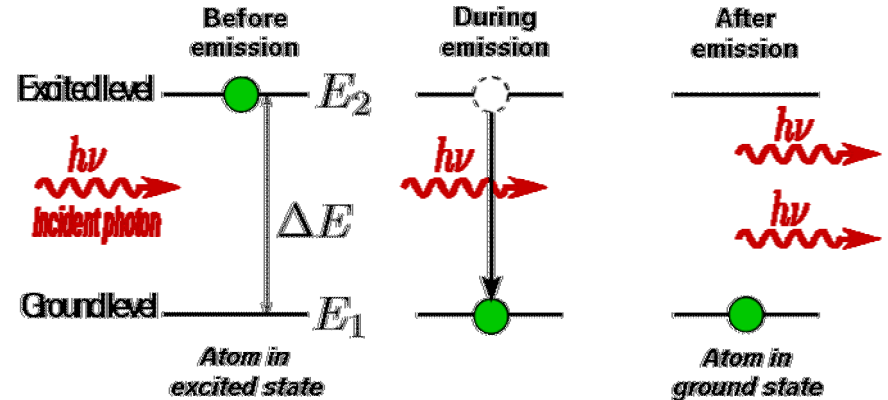
- Building world's first hyperspectral STED microscope
  - Patented by Jeri Timlin (8631) and Jesse Aaron

# Stimulated Emission Details

## Basic Principle of Stimulated Emission



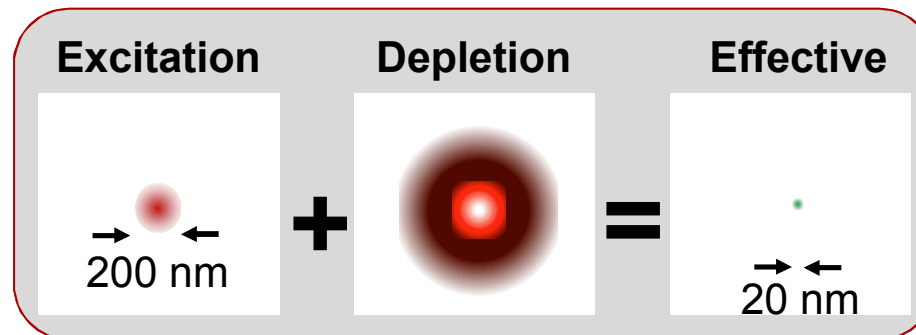
[https://en.wikipedia.org/wiki/File:STED\\_Jablonski.jpg](https://en.wikipedia.org/wiki/File:STED_Jablonski.jpg)



$$E_2 - E_1 = \Delta E = h\nu$$

By V1adis1av - Own work, GFDL, <https://commons.wikimedia.org/w/index.php?curid=3983414>

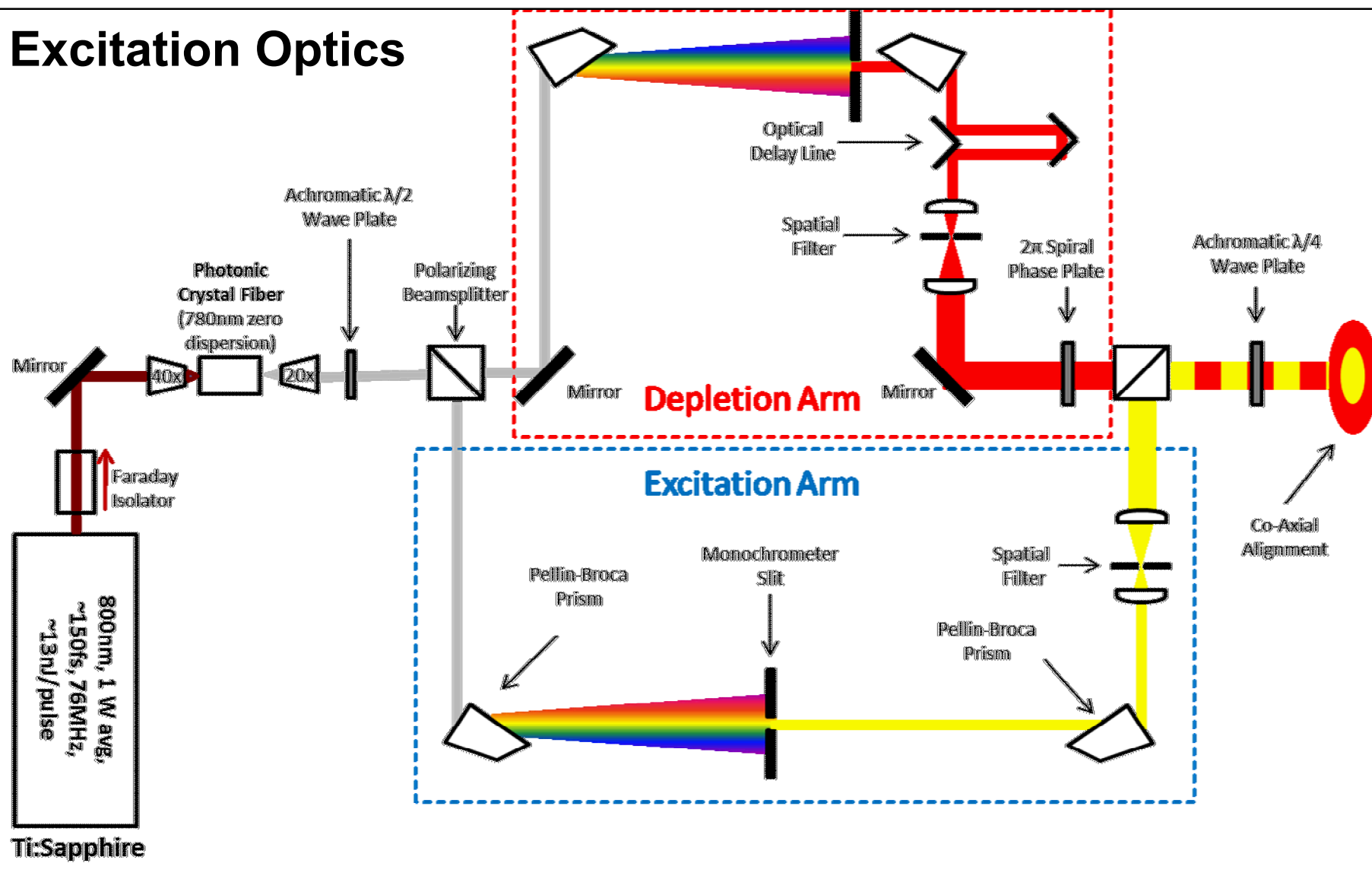
## Generating the STED Point Spread Function (PSF)



Neither beam PSF can exceed the diffraction limit, but the effective PSF can!

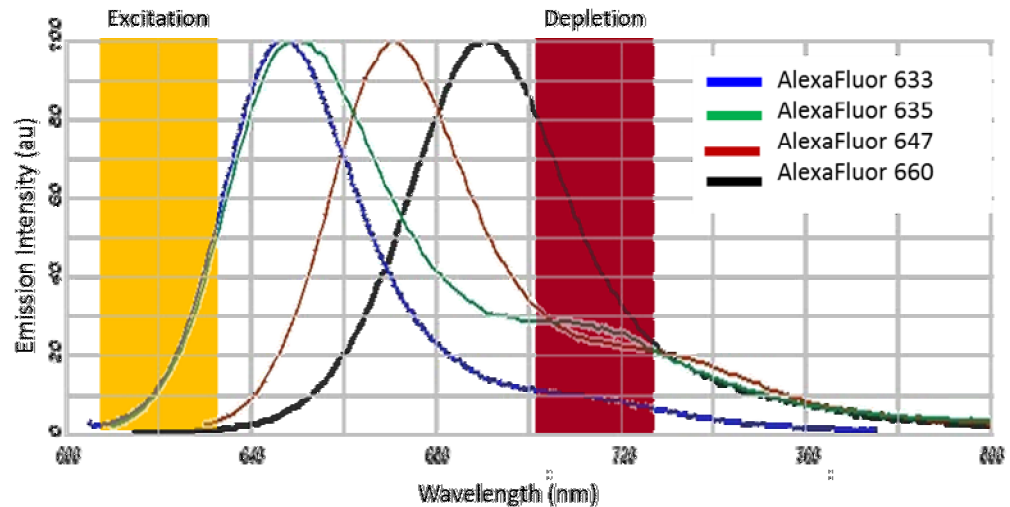
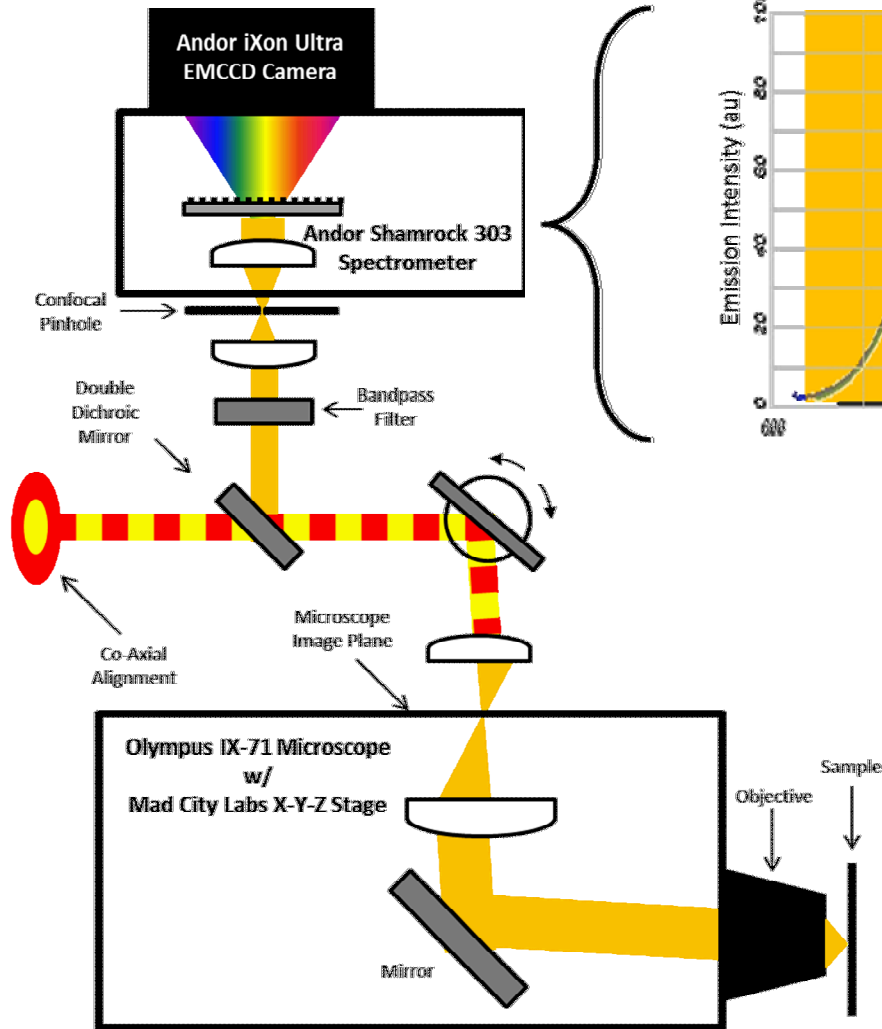
# Building a Hyperspectral STED

## Excitation Optics



# Building a Hyperspectral STED

## Detection Optics



## Design Considerations

- Tunable wavelength for both excitation and depletion beams
- Can be optimized for any STED fluorophore with exchange of a single optic (the dichroic)

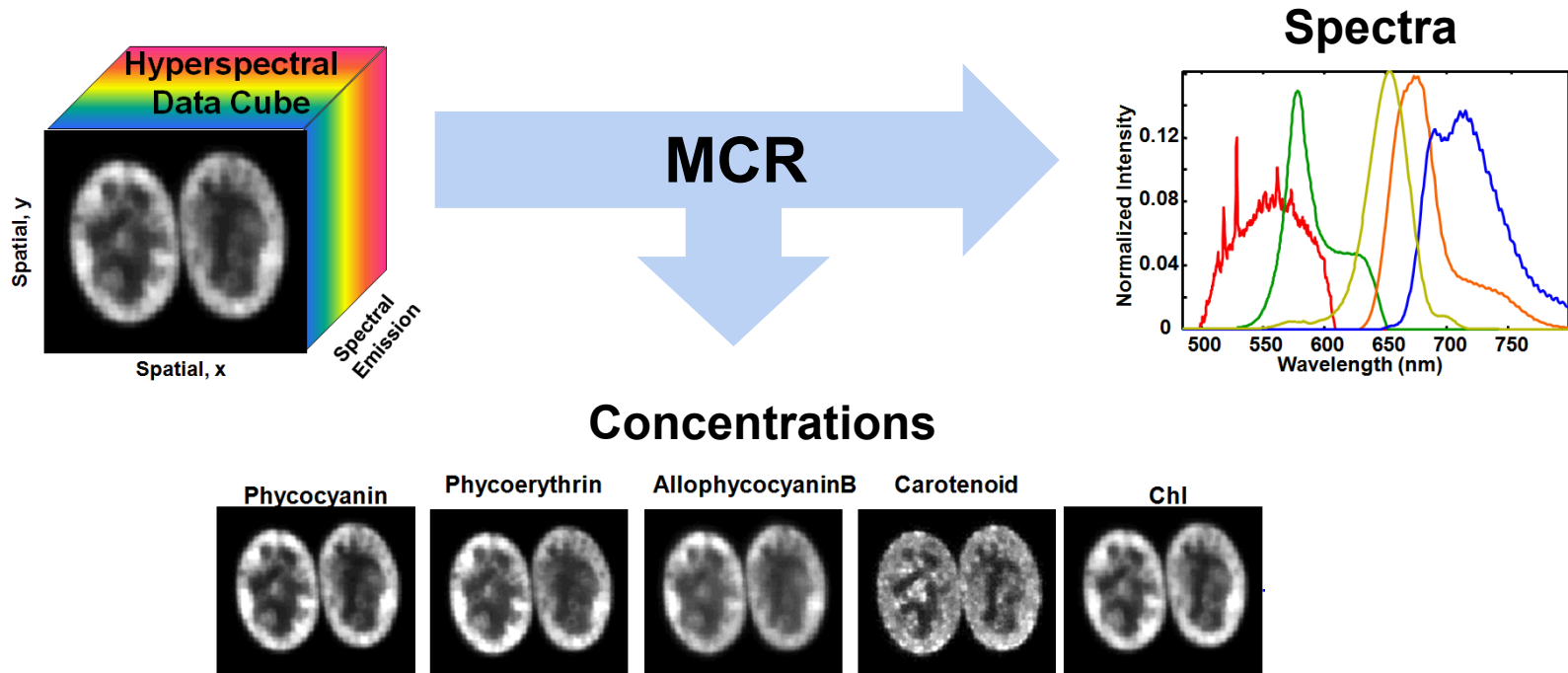
# Outline

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# Multivariate Curve Resolution (MCR)

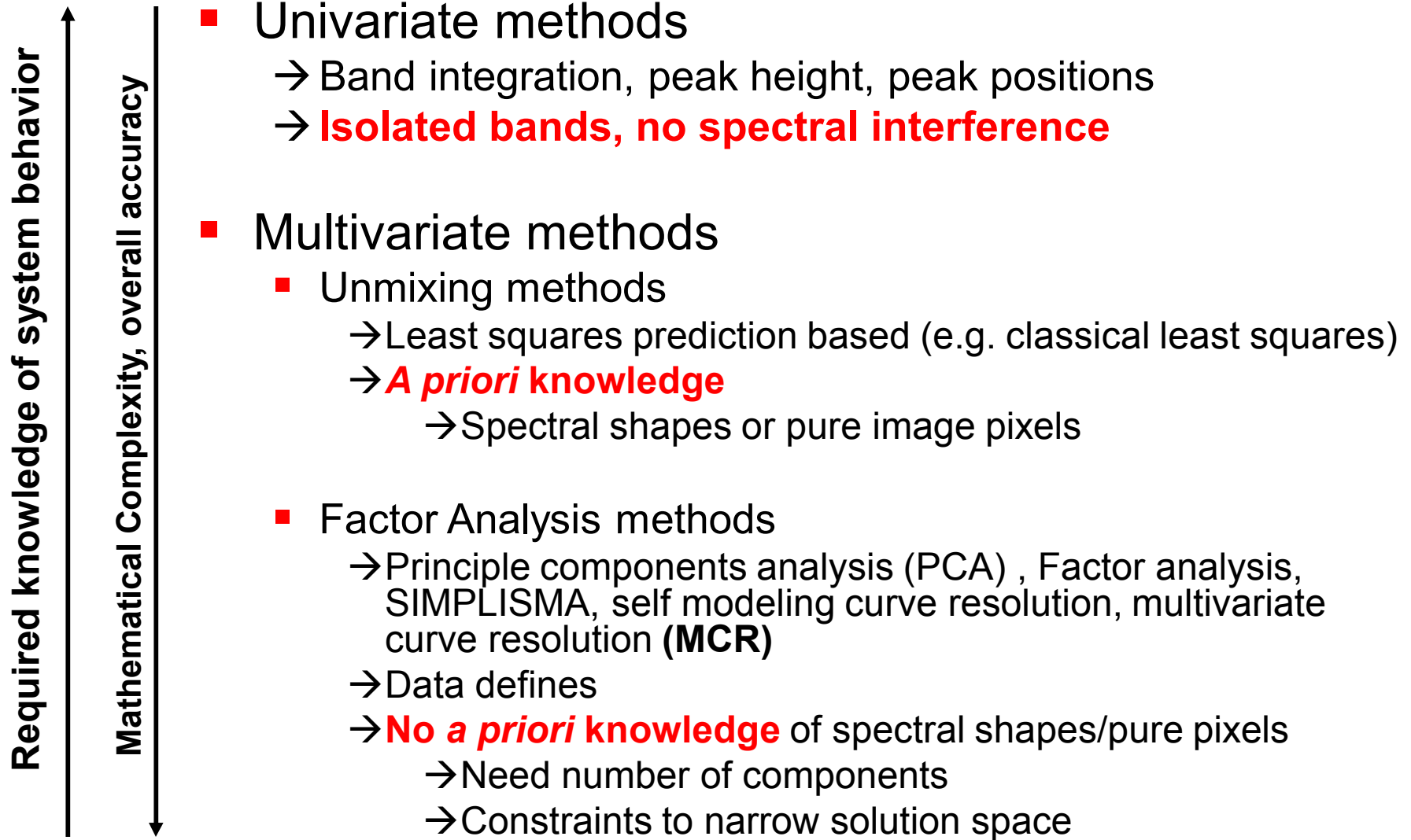
Chemometric factor analysis (such as MCR) extracts:

- 1) the spectra of the pure compounds &
- 2) their relative concentrations at each position.



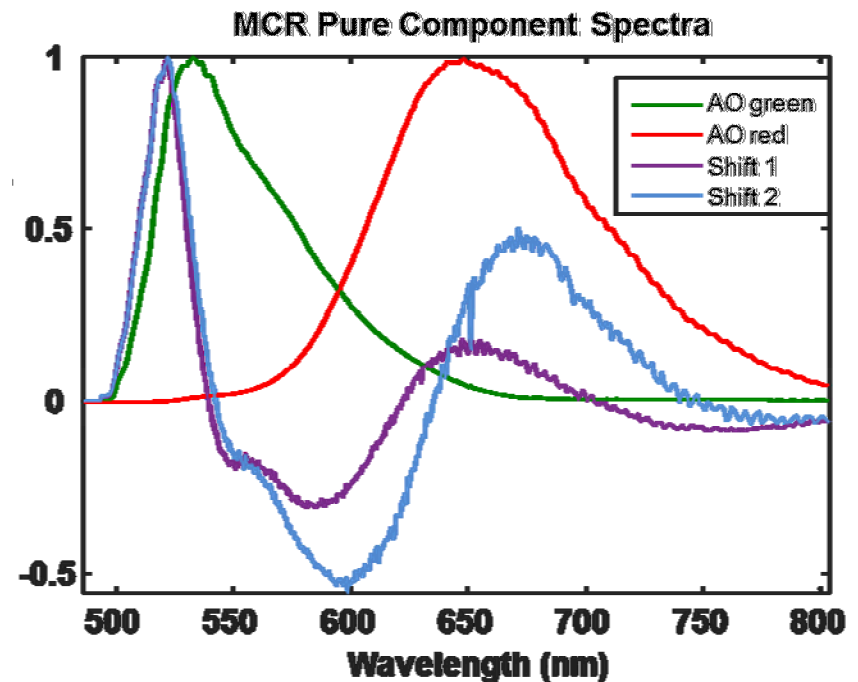
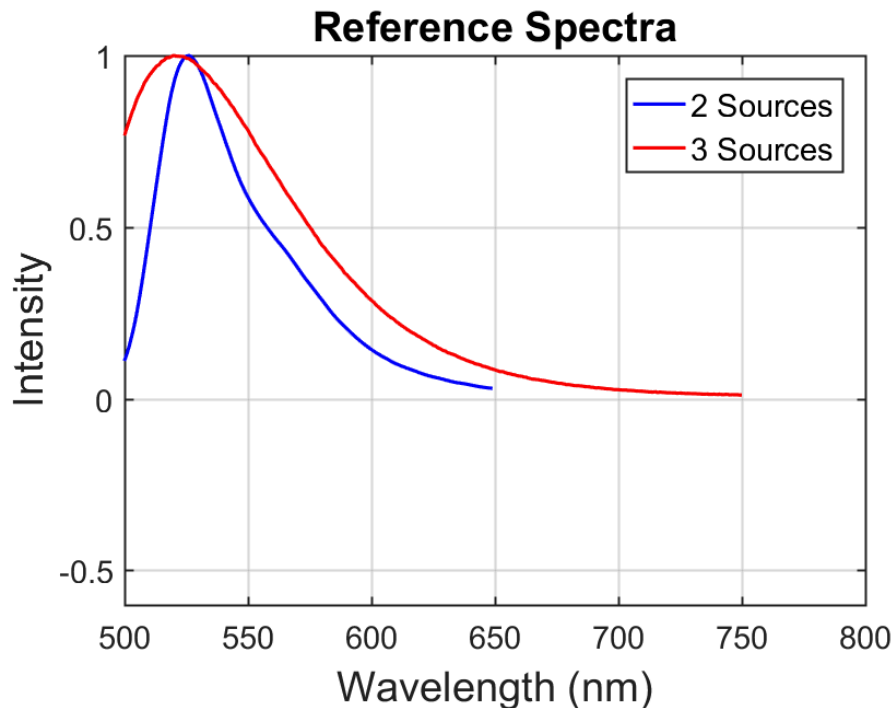


# Common Spectral Image Analysis Methods



# Why Aren't the Spectra Known?

“Acridine Orange is a cell-permeant nucleic acid binding dye that emits **green fluorescence** when bound to dsDNA and **red fluorescence** when bound to ssDNA or RNA.” - ThermoFisher Scientific



**Reference spectra are not always available, and when available do not always capture the complete spectral properties.**

# Broadly Applicable

**Works for any data satisfying the linear additive model:**

$$\mathbf{D} = \mathbf{CS}^T + \mathbf{E}$$

- **Fluorescence spectroscopy**
- **Raman spectroscopy**
- **Mass spectroscopy**
- **Infrared satellite imagery**

# MCR Assumptions

## 1. Assumes a linear additive model:

$$\mathbf{D} = \mathbf{C}\mathbf{S}^T + \mathbf{E}$$

$\mathbf{D}$  = Data matrix                      nPoints X nWavelengths

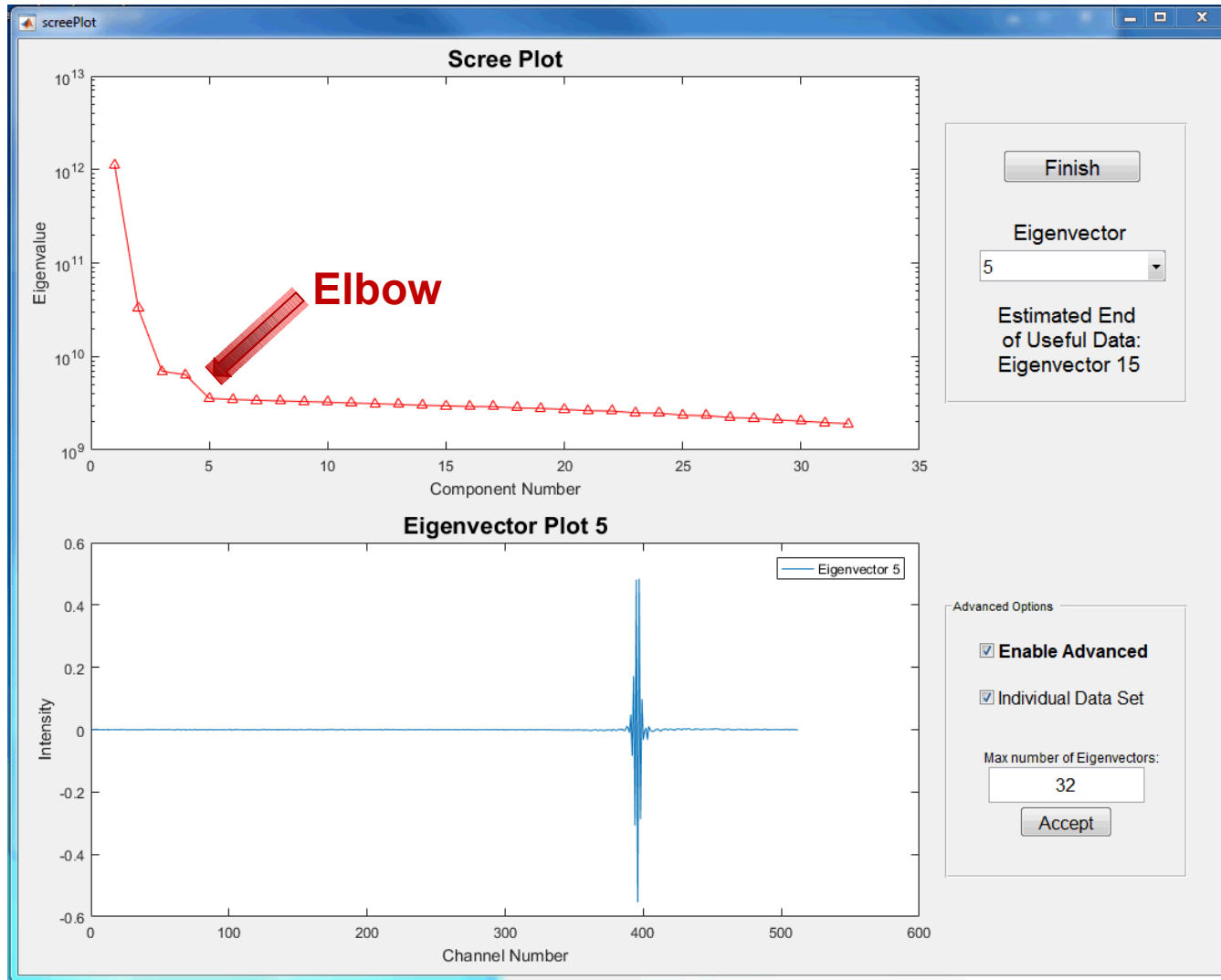
$\mathbf{C}$  = Concentrations matrix              nPoints X nComponents

$\mathbf{S}^T$  = (Spectra matrix)<sup>Transpose</sup>      nComponents X nWavelengths

$\mathbf{E}$  = Noise (error) matrix              nPoints X nWavelengths

## 2. The # of spectral components is known or can be estimated

# Determining the # of Spectra



Scree plots allow rough estimation of the number of spectra.

Elbow (transition to flat) in scree and eigenvectors that look like noise indicate no more components.

# Basic Operation

$$\mathbf{D} = \mathbf{CS}^T + \mathbf{E}$$

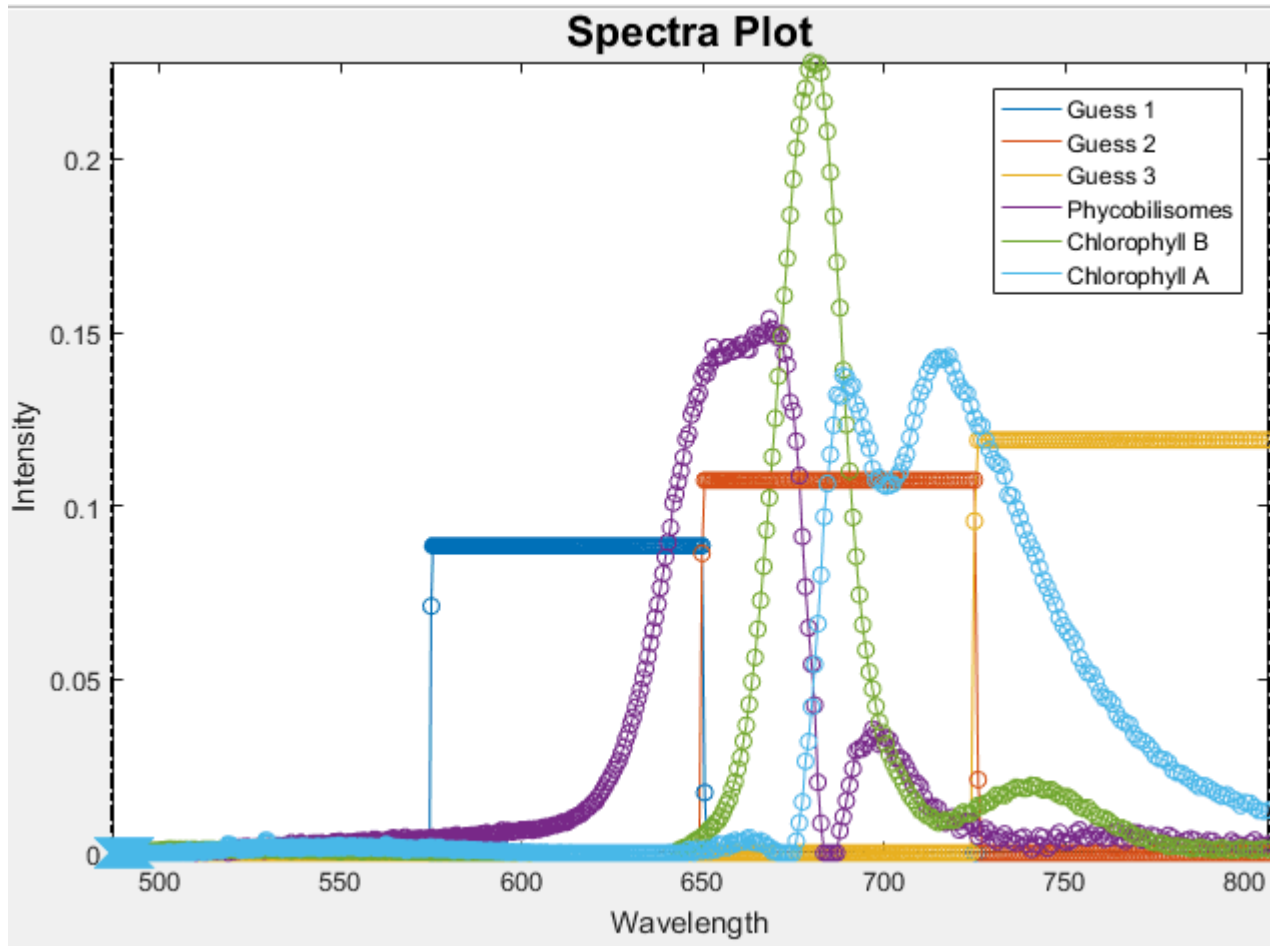
- D is known
  - If C were known, could solve for S
  - If S were known, could solve for C
- Constrained Alternating Least Squares
  1. Provide an initial guess for S (or C)
  2. Solve for C based upon current S guess, enforcing constraints
  3. Solve for S based upon current C guess, enforcing constraints
  4. Repeat steps 2 & 3
  - 5. Iterations converge on solution**

# Advantages of MCR

- Extracts underlying relationships from complex data sets
- No *a priori* knowledge needed
- Signals below the noise level can be detected!
- Physically meaningful constraints
  - Negative concentrations not allowed
  - Negative intensities not allowed
- Efficient algorithms developed at Sandia
  - Keenan, M. R. and P. G. Kotula (2003). Apparatus and system for multivariate spectral analysis, Google Patents.



# No *A Priori* Spectra Required



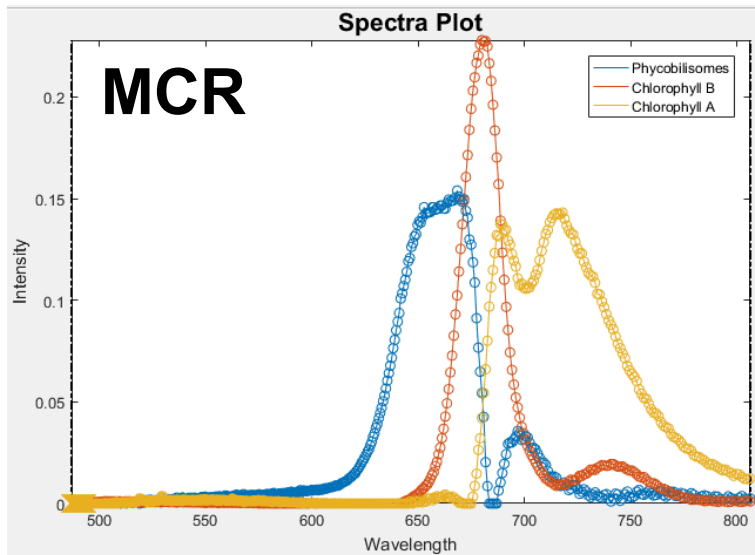
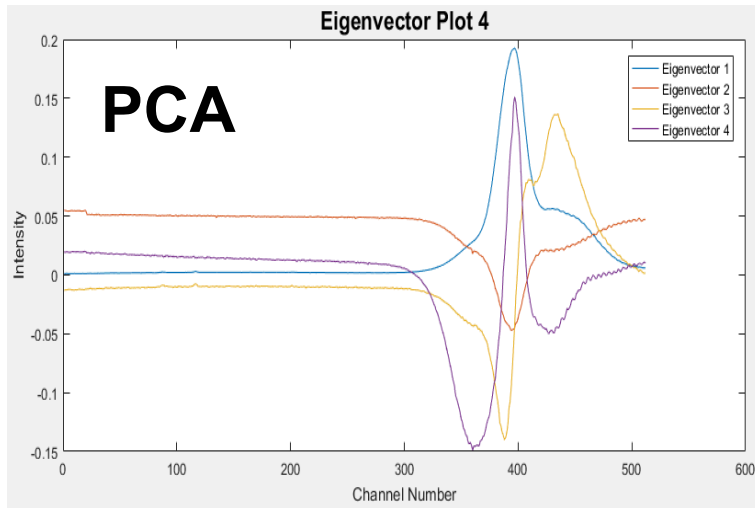
Hyperspectral Confocal  
Fluorescence Microscopy  
of R61 *Acaryochloris*

Even when  
initialized with  
naïve guesses,  
MCR determines  
the spectrum of  
chlorophyll B and  
localizes where it is  
present.

# Why MCR vs. PCA?

- Three related factor analysis methods
  - Multivariate Curve Resolution (MCR)
  - Principal Component Analysis (PCA)
  - Independent Component Analysis (ICA)
- All resolve the data into pure spectral components and concentrations without a priori information
- **Different Constraints**
  - MCR – Physical and Chemical Constraints (e.g. no negative concentrations, no negative intensities)
  - PCA – Linearly uncorrelated
  - ICA – Statistically Independent

# Why MCR vs. PCA?



## ■ PCA

- Recovers 4 spectral components when only 3 are present
- Eigenvectors do not look like normal spectra – major negative portions

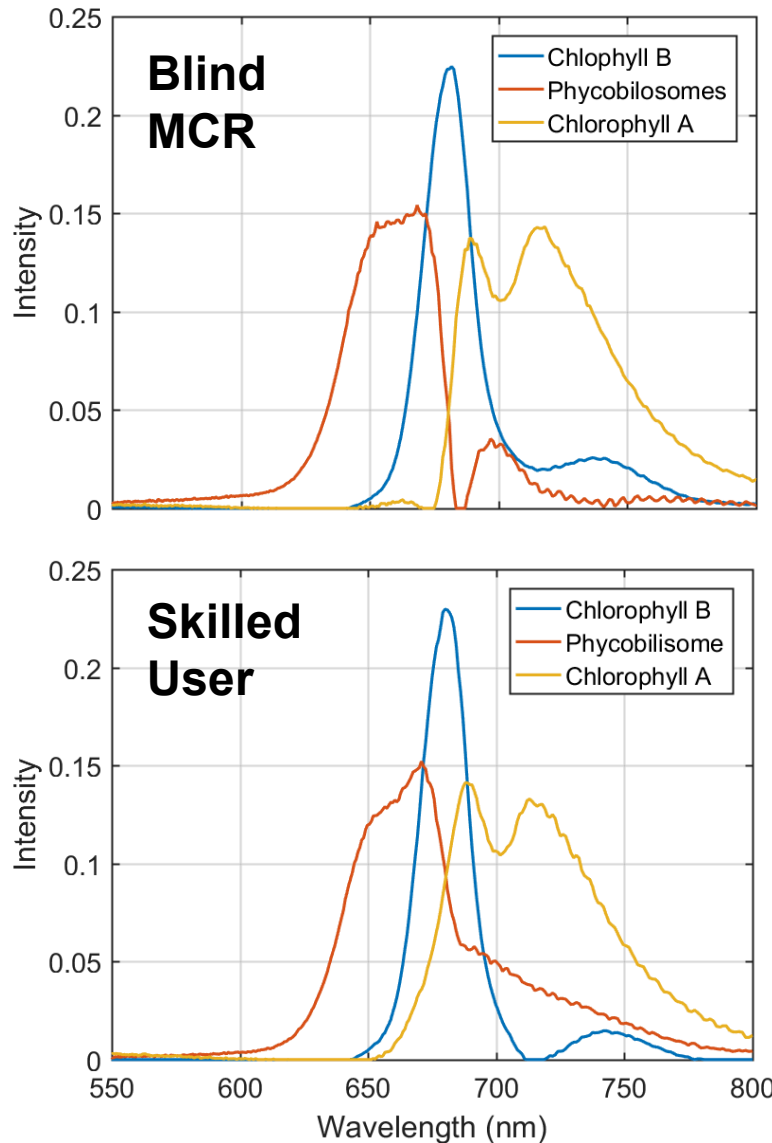
## ■ MCR

- Recovers the correct number of components
- Components generally correspond to actual spectra

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# The Art of MCR



While MCR can provide excellent results, it is currently more of an art than a science.

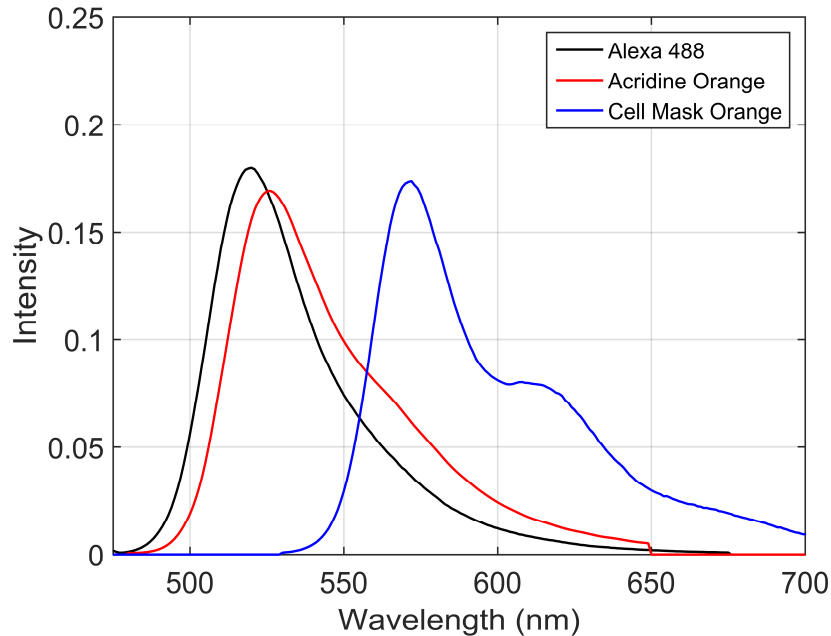
A skilled user will obtain better spectra from MCR, particularly for weaker spectral components.

Relative amounts:

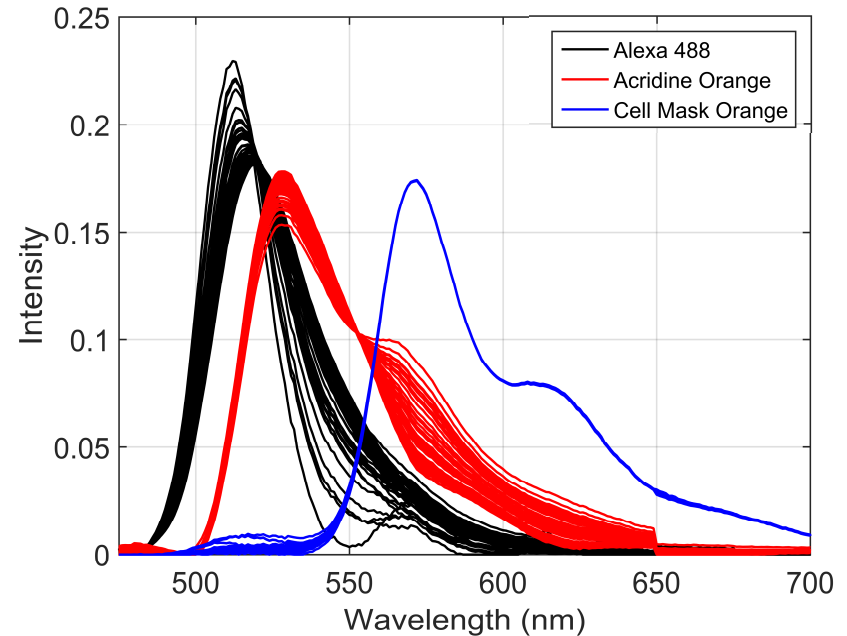
Chlorophyll B:	3.7
Chlorophyll A:	2.2
Phycobilisomes:	1.0

# Multitude of Results

## Pure Spectra



## MCR Spectra (100 different runs)



MCR results for 100 runs on a simulated data set for a 100 x 100 pixel hyperspectral image averaging ~55000 counts for each spectrum initialized with random spectra.

## Why multiple results? Two possibilities:

- Failure to converge – trapped in a local minimum
- Rotational ambiguity – results mathematically equally good

## Also known as the rotation problem:

- Construct an invertible transformation matrix  $M_i$  to operate on the matrices  $C$  and  $S$ .

$$D = CS^T = (CM_i^{-1}) (M_i S^T)$$

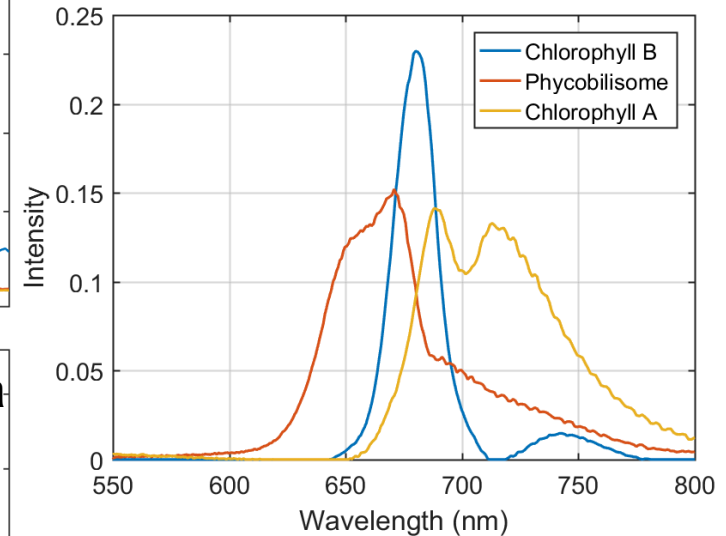
- The resulting matrices  $(CM_i^{-1})$  and  $(M_i S^T)$  are an alternate solution with identical residuals

**Constraints may reduce or eliminate rotational ambiguity.**



# Data Size

## Desired Results 653k spectra



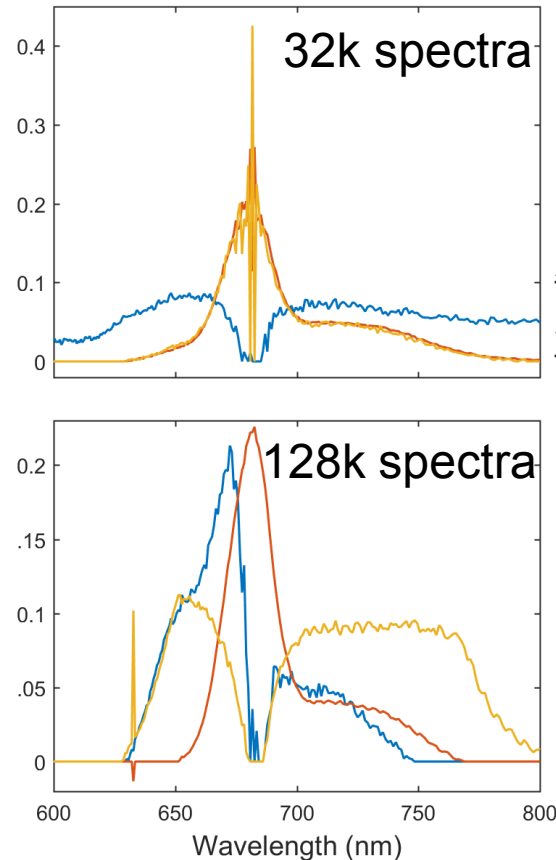
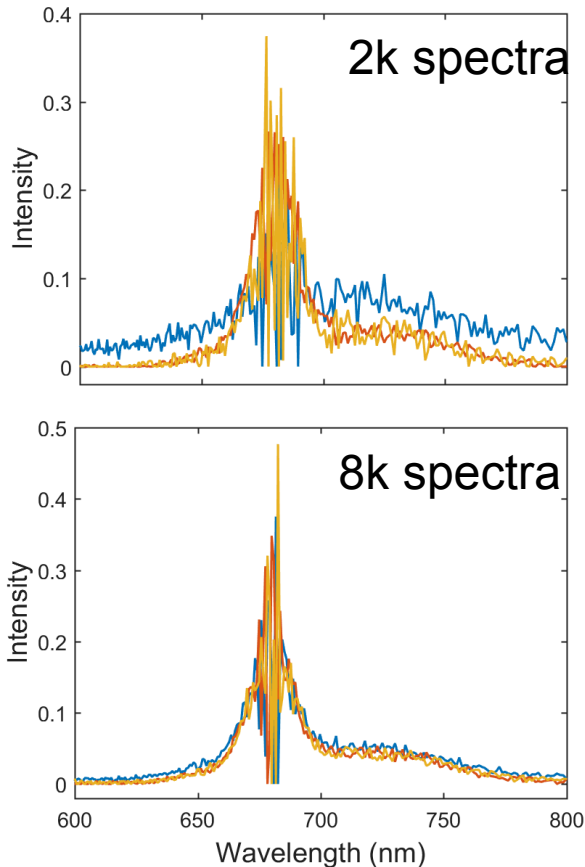
Relative amounts:

Chlorophyll B: 3.7

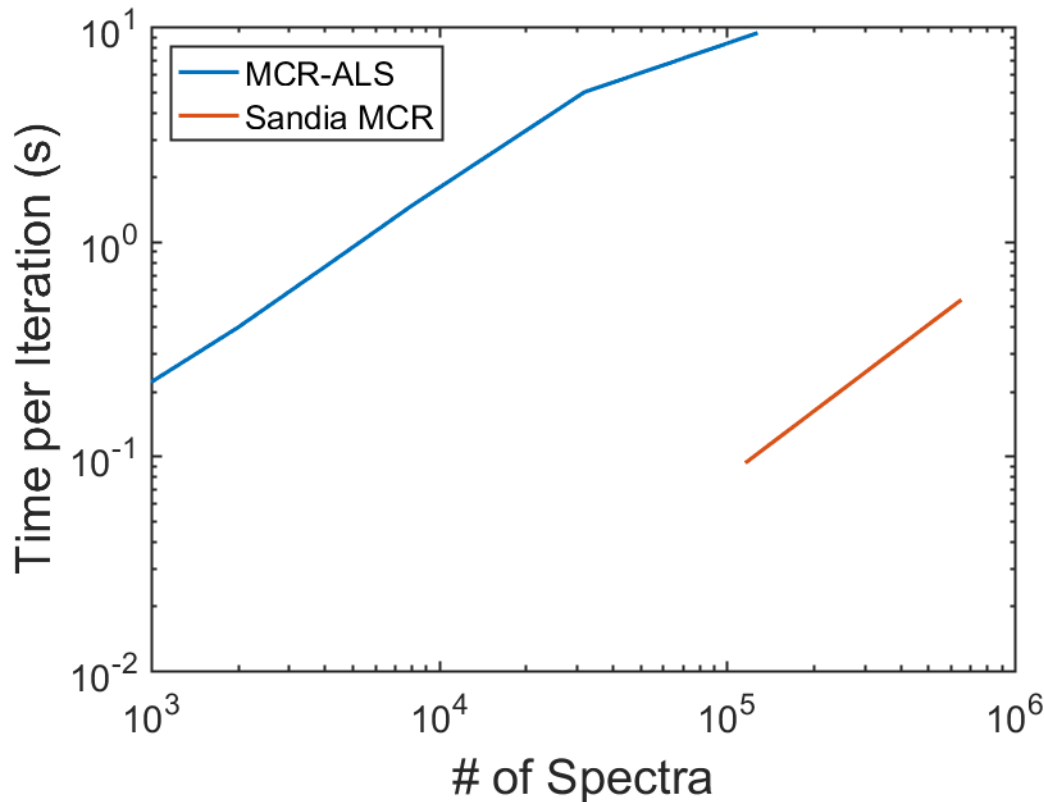
Chlorophyll A: 2.2

Phycobilisomes: 1.0

# Working with more data provides better results



# Computational Requirements



Sandia's MCR algorithm runs ~50 times faster than competing code<sup>1</sup> released in 2015.

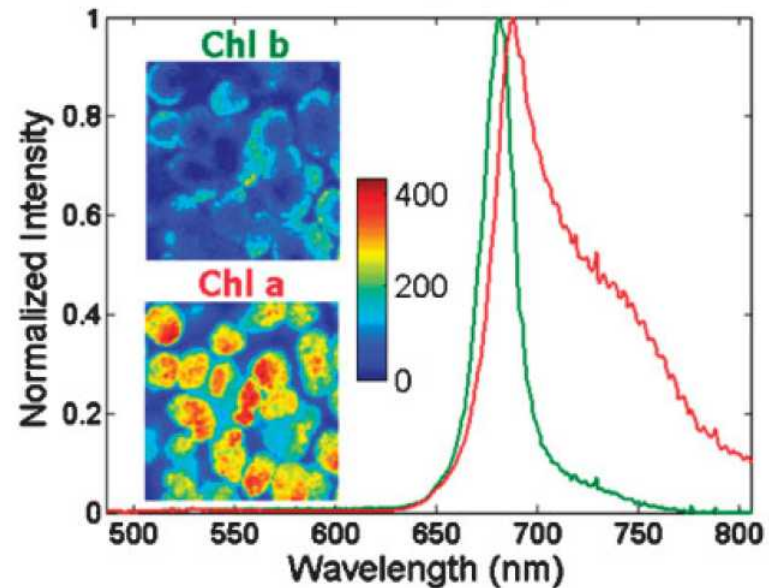
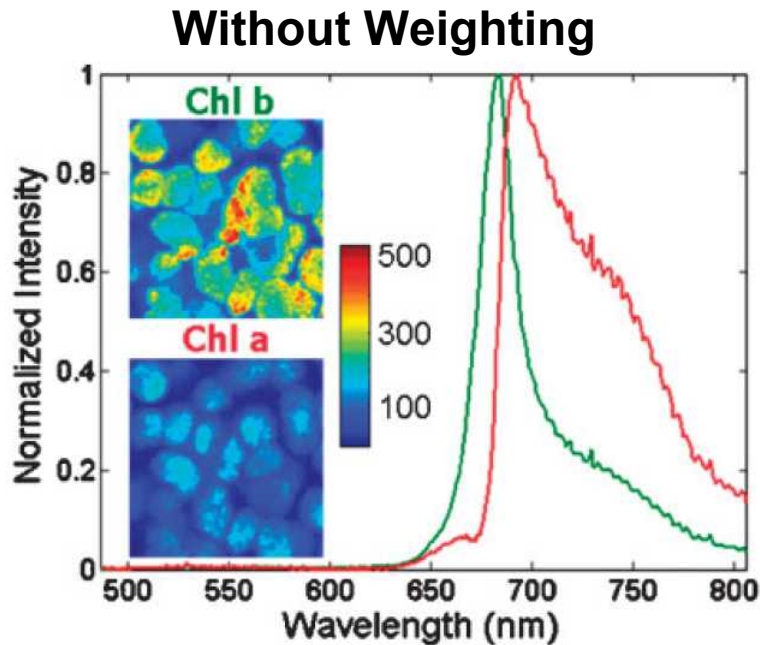
Competing code<sup>1</sup> is widely used – cited 46 times in 2 years.

1) Jaumot, J. et al., "MCR-ALS GUI 2.0: new features and applications." *Chemometrics and Intelligent Laboratory Systems* 140 (2015): 1-12.

Computational time is on a high-end PC

# Ongoing Work – Improved Weighting

Weighted for  
Poisson & Gaussian Noise



Jones, H. D. T., et al. (2008). "Weighting hyperspectral image data for improved multivariate curve resolution results." *Journal of Chemometrics* **22**(9-10): 482-490.

## Proper weighting makes a major difference!

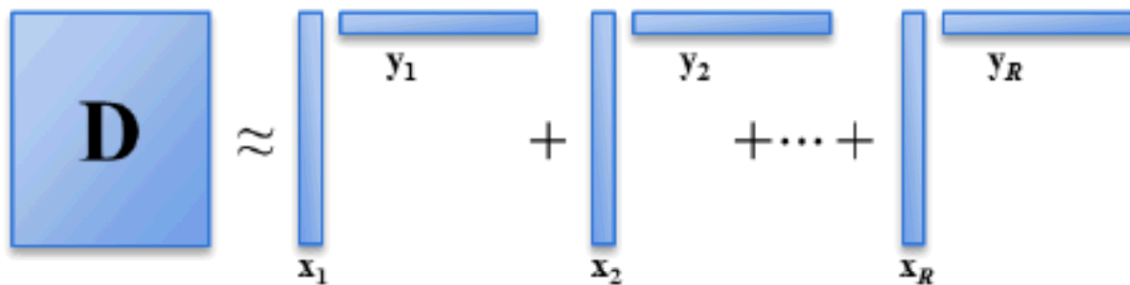
Working on improving the weighting to correctly account for all sources of noise, including the pre-processing steps.

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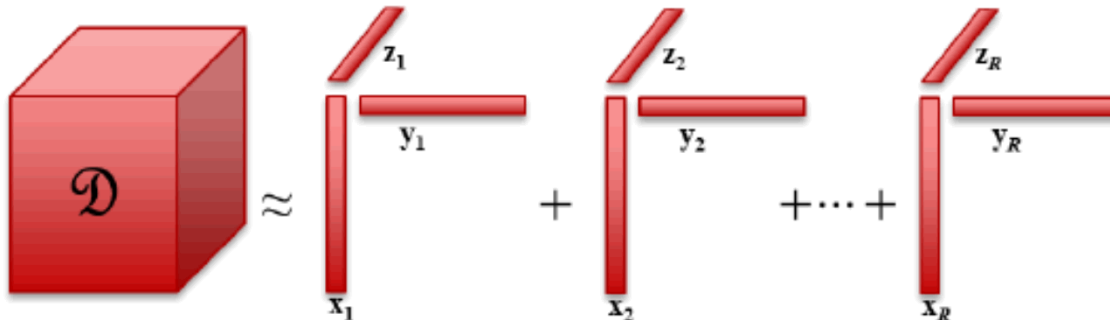
# Bilinear vs. Trilinear Data

$$\mathbf{D} \approx \sum_{i=1}^R \mathbf{x}_i \mathbf{y}_i^T$$



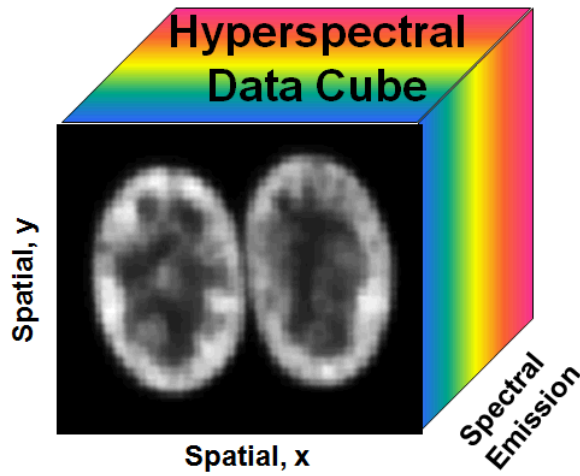
**Bilinear Data** – Each component in the data matrix can be expressed as the product of two vectors.

$$\mathcal{D} \approx \sum_{i=1}^R \mathbf{x}_i \circ \mathbf{y}_i \circ \mathbf{z}_i$$

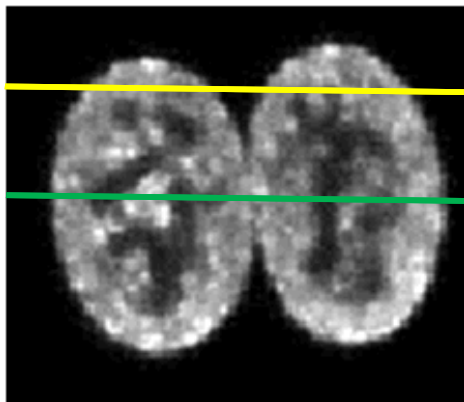


**Trilinear Data** – Each component in the data tensor can be expressed as the product of three vectors.

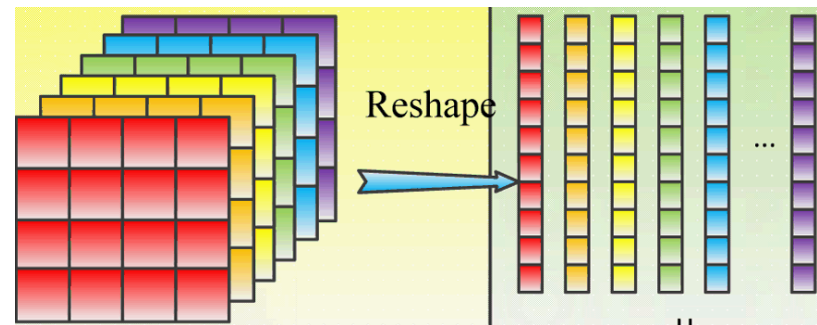
# Hyperspectral Data Cube = Bilinear



**Carotenoid**



While hyperspectral data cubes are 3-dimensional, typically the data is only bilinear, not trilinear. Prior to processing with MCR, the data must be reshaped.



Adapted from Wu, Zhaojun, et al. *Journal of Electronic Imaging* 25.1 (2016): 013037-013037.

In order to be trilinear, for an individual component, the cross sections at different y positions (yellow and green lines) would need to be identical.

# Examples of Trilinear Data

- Hyperspectral fluorescence lifetime
  - Spatial position
  - Wavelength
  - Lifetime
- Gas Chromatography
  - Elution Time
  - Mass spectrum
  - Multiple possibilities (sample number, run temperature)

# Analyzing Trilinear Data

- MCR can be applied to trilinear data, but better methods exist
  - Trilinear data can always be reshaped to generate a bilinear data set.
  - Doing so forfeits one of the great advantages of trilinear data. Trilinear analysis eliminates the rotational ambiguity problem common to bilinear data.
- Trilinear methods exist
  - Parafac, Tucker3 algorithms are examples
  - For trilinear data analysis, talk to Mark van Benthem (1814)



# Summary

- Hyperspectral microscopy and MCR are valuable tools
- Hyperspectral STED will provide super-resolution capability
- Improvements to MCR will:
  - Make MCR less of an art
  - Improve estimation of the weaker components

## Acknowledgements

Collaborators or prior research:

Microscopy: Jeri A. Timlin, Michael B. Sinclair, Jesse S. Aaron

MCR: Mark Van Benthem, Jeri A. Timlin, Michael R. Keenan, Paul G. Kotula

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