

# unmixR: Hyperspectral Unmixing in R

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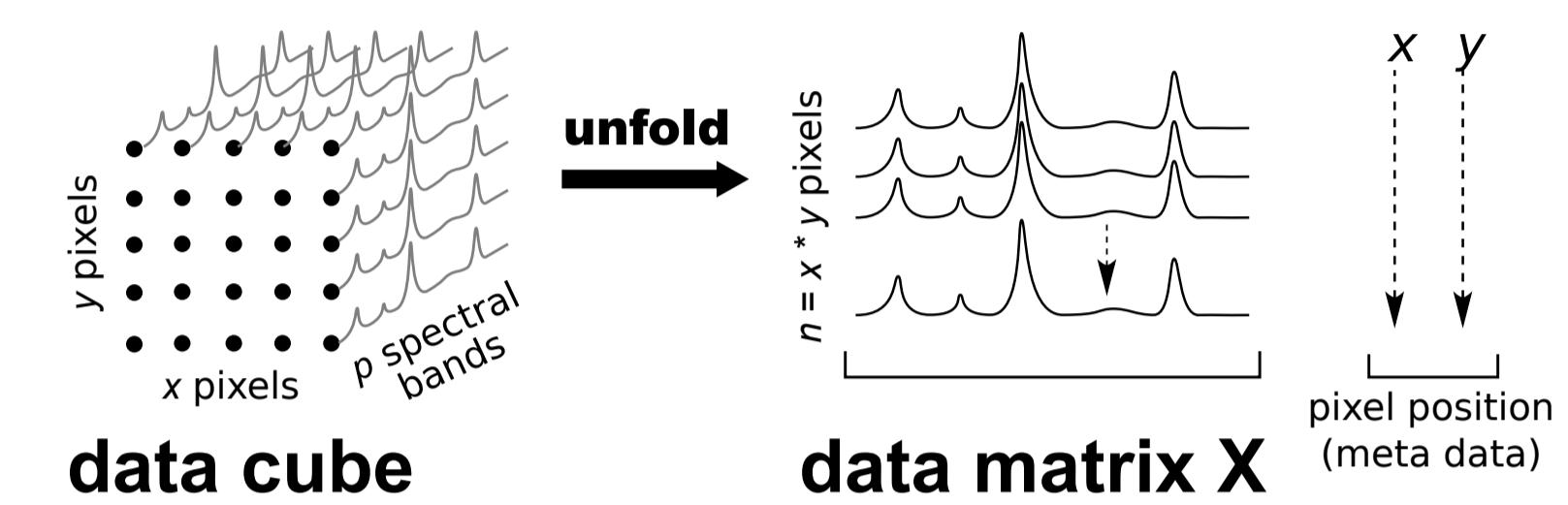
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## Hyperspectral Imaging

**Hyperspectral images** are 3D data sets of spectra collected over an  $x, y$  grid.



**Applications:** remote sensing/ airborne or satellite land imaging, biomedical microspectroscopy and art history investigations

**Spectra:** e.g. visible, near-infrared, mid-infrared, or Raman spectra.

## Spectral Unmixing

Identify  $m$  pure component spectra in data, then derive respective concentrations.

**Bilinear statistical model:**

$$\mathbf{X}^{(n \times p)} = \mathbf{A}^{(n \times m)} \times \mathbf{E}^{(m \times p)} + \boldsymbol{\varepsilon}$$

mixture spectra      abundances      endmember / noise

molar fractions      pure component spectra

Mixture diagram for  $m$  components:  $(m - 1)$ -simplex in  $m - 1$  dimensions ( $\mathbf{A}$ ).

2 components	1-simplex	line
3 components	2-simplex	triangle
4 components	3-simplex	tetrahedron

Vertices are pure component spectra.

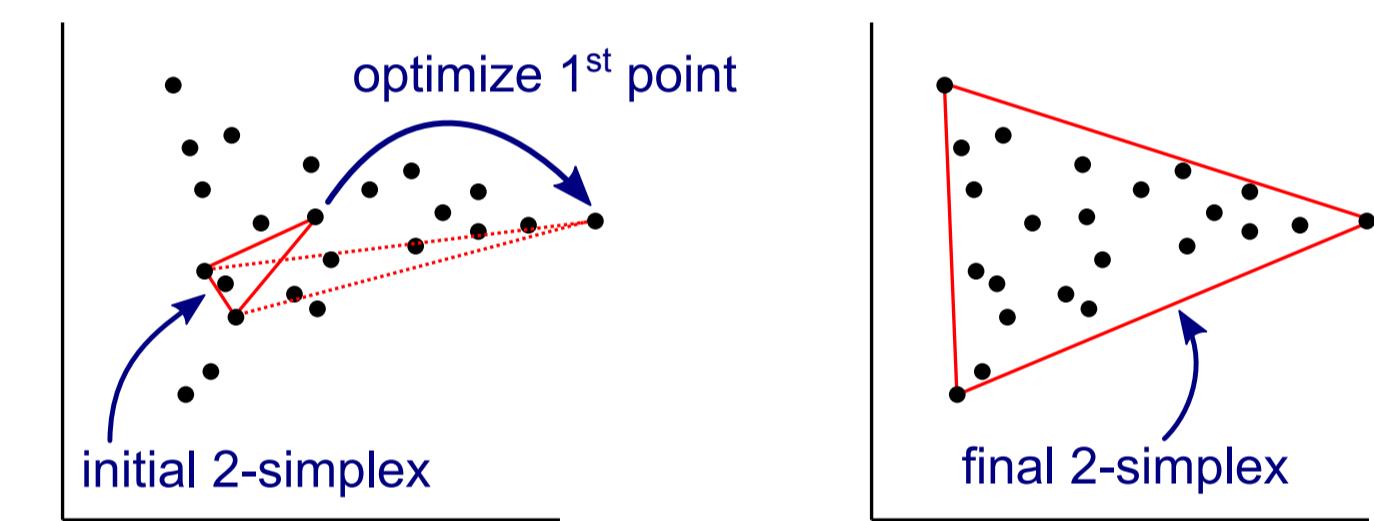
**Assumptions:**

- Data consists of *mixture spectra*
- Spectra of pure components are available somewhere in the data  $\mathbf{X}$
- Not too much noise on measurements (possibly after PCA)
- (Other methods relax assumptions 2 and 3)
- Number of pure components  $m$  ("chemical rank") provided by user input
- Abundances subject to non-negativity constraint

## N-FINDR Algorithm

**Heuristic:** find  $m$  spectra within data set that span  $(m - 1)$ -simplex with *largest volume*

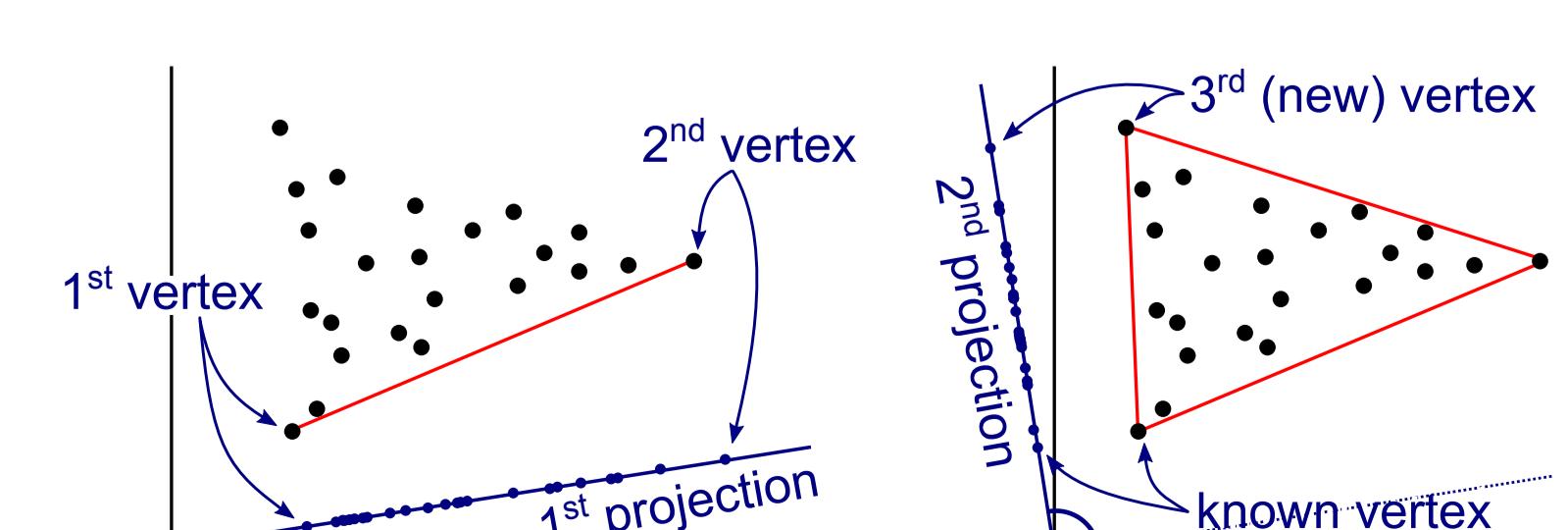
- Project  $\mathbf{X}$  into  $(m - 1)$ -dimensional space (typically by PCA)
- Initialize simplex with  $m$  arbitrary points
- Iteratively grow simplex:  
For each vertex point in turn:  
exchange by point that maximizes simplex volume (keeping the other  $m - 1$  points constant)  
Iterate/refine until convergence
- Return corresponding spectra of  $\mathbf{X}$  as endmembers
- predict abundances by non-negative least squares [nnls] on found endmembers



## VCA Algorithm

**Heuristic:** projection of points onto arbitrary direction will always have 2 of the  $m$  vertices as maximum and minimum.

- Project  $\mathbf{X}$  into  $(m - 1)$ -dimensional space if data is considered too noisy
- Project  $\mathbf{X}$  onto arbitrary direction
- Find first 2 vertices as min and max
- Project  $\mathbf{X}$  onto arbitrary direction orthogonal to all previously used directions
- Find next vertex as unknown min or max
- Repeat 4 and 5 until  $m$  vertices are found
- Return corresponding spectra of  $\mathbf{X}$  as endmembers
- predict abundances by non-negative least squares [nnls] on found endmembers



## AVIRIS Cuprite Data

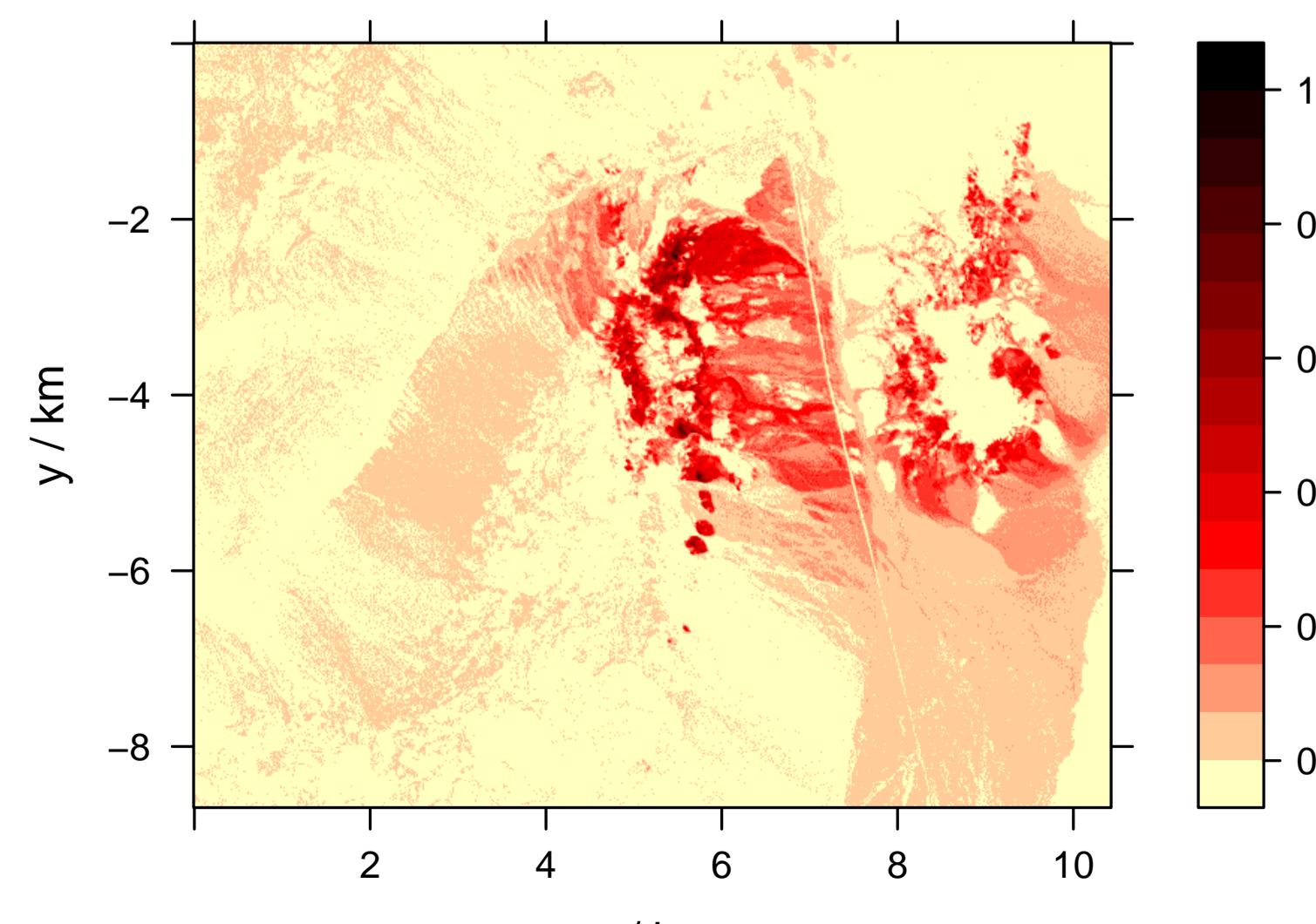
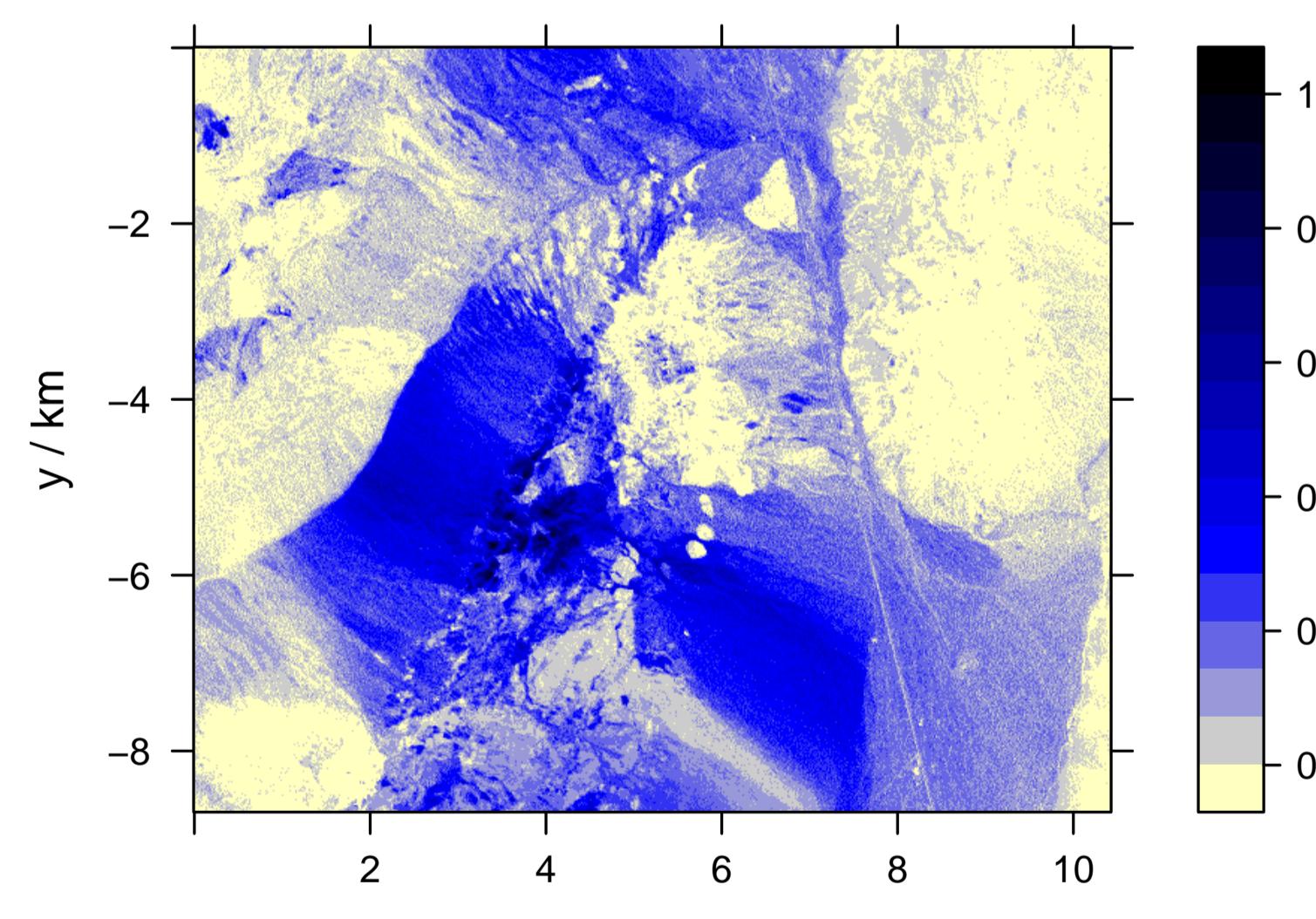
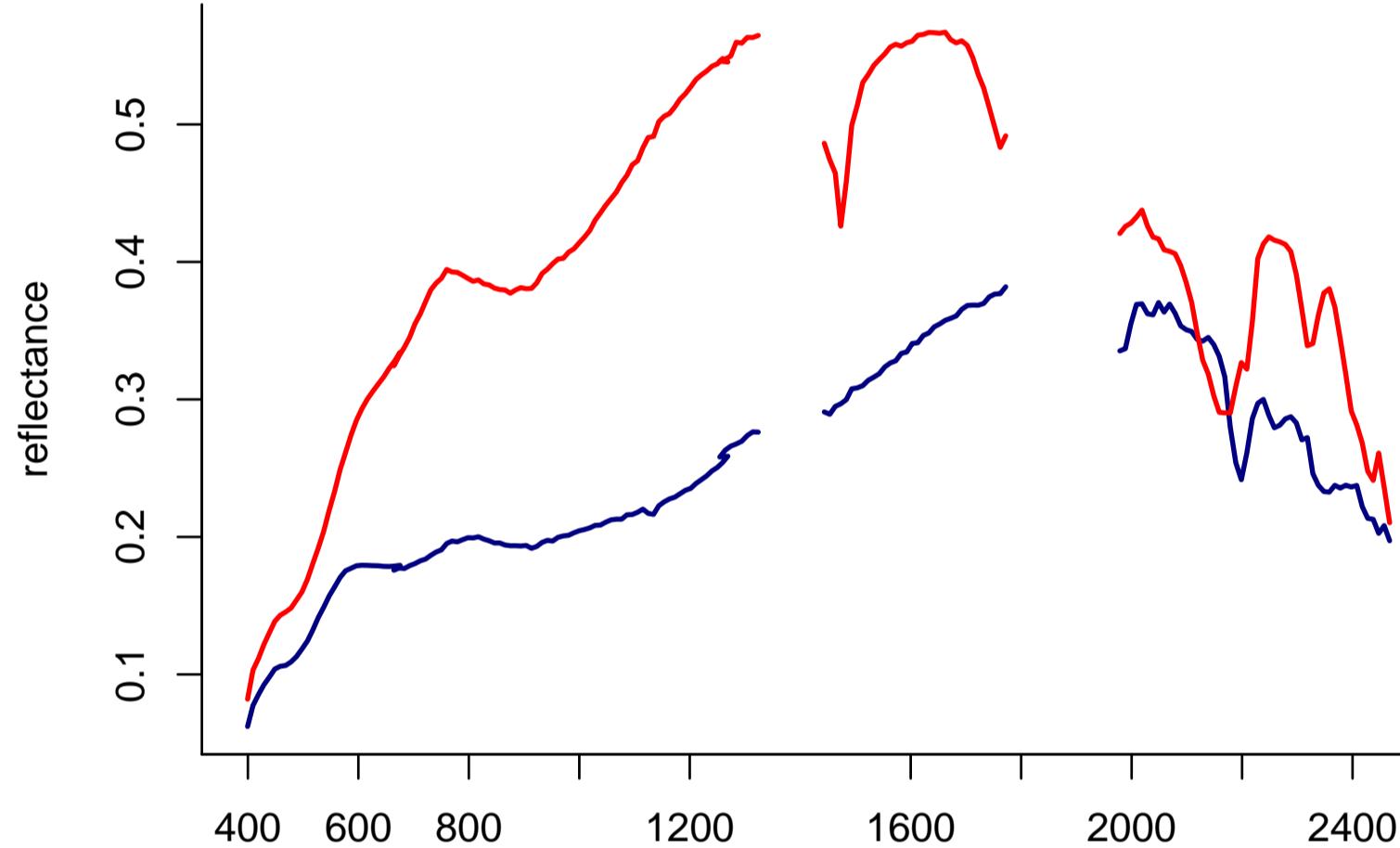
**Data Set:**

- Acquired by NASA's Airborne Visible/ InfraRed Imaging Spectrometer
- of mining region in the south of Nevada/USA
- $45 \times 10$  km (300 000 pixel subimage shown)
- 250 - 4 000 nm (224 spectral bands)
- Well-known ground truth

### N-FINDR with $m = 19$ endmembers

As example, we show 2 components identified as

- muscovite** (mica,  $KAl_2(AlSi_3O_{10})(F_OH)_2$ ), and
- alunite** (alumstone,  $KAl_3(SO_4)_2(OH)_6$ ).

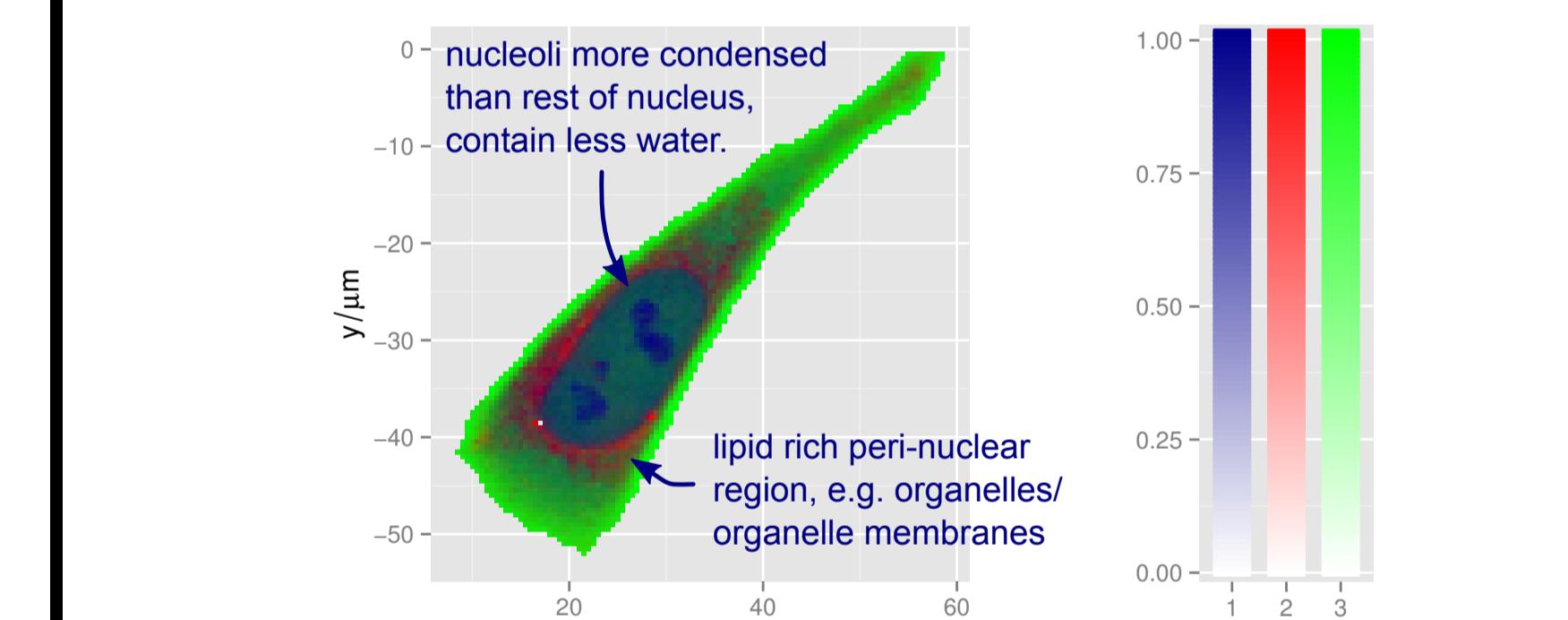
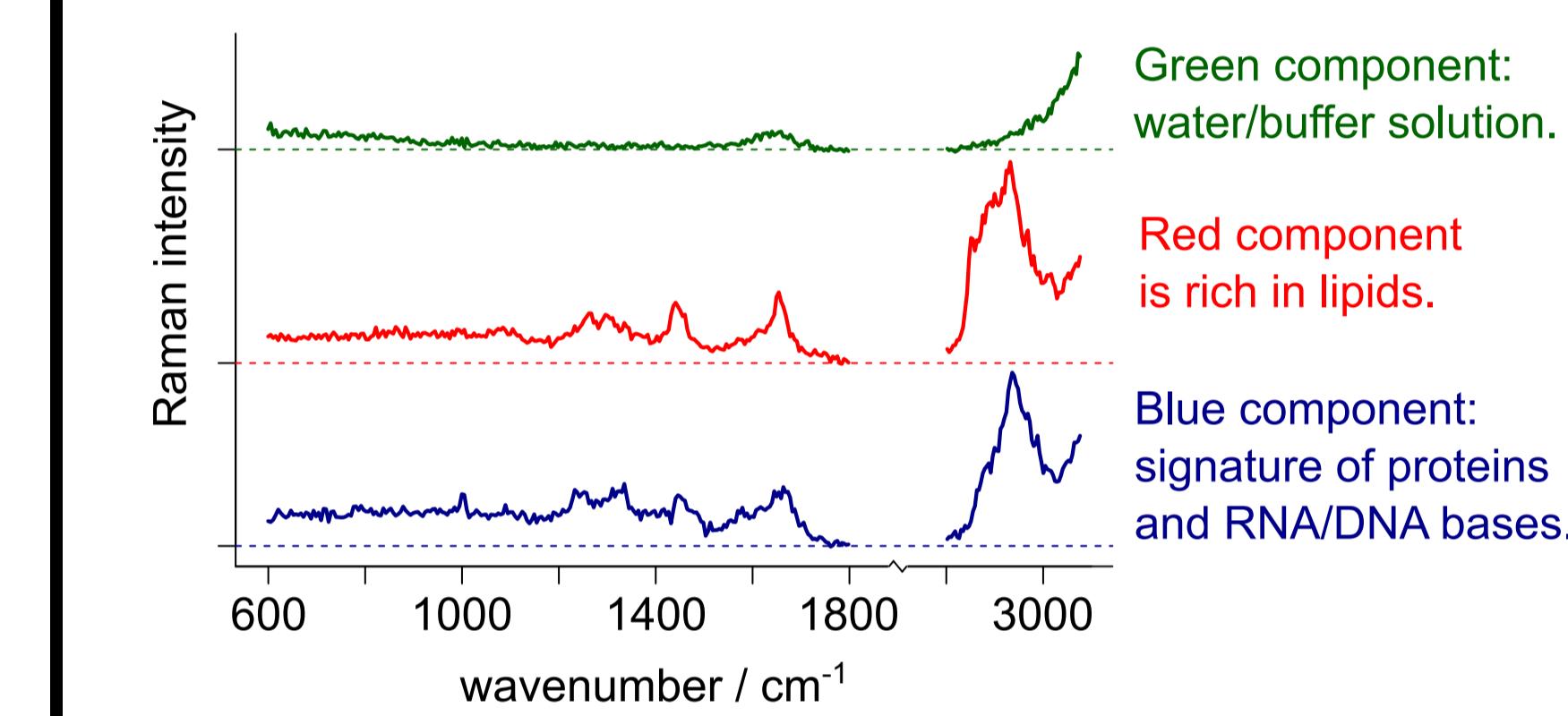


## Raman Image of HeLa Cell

**Data Set:**

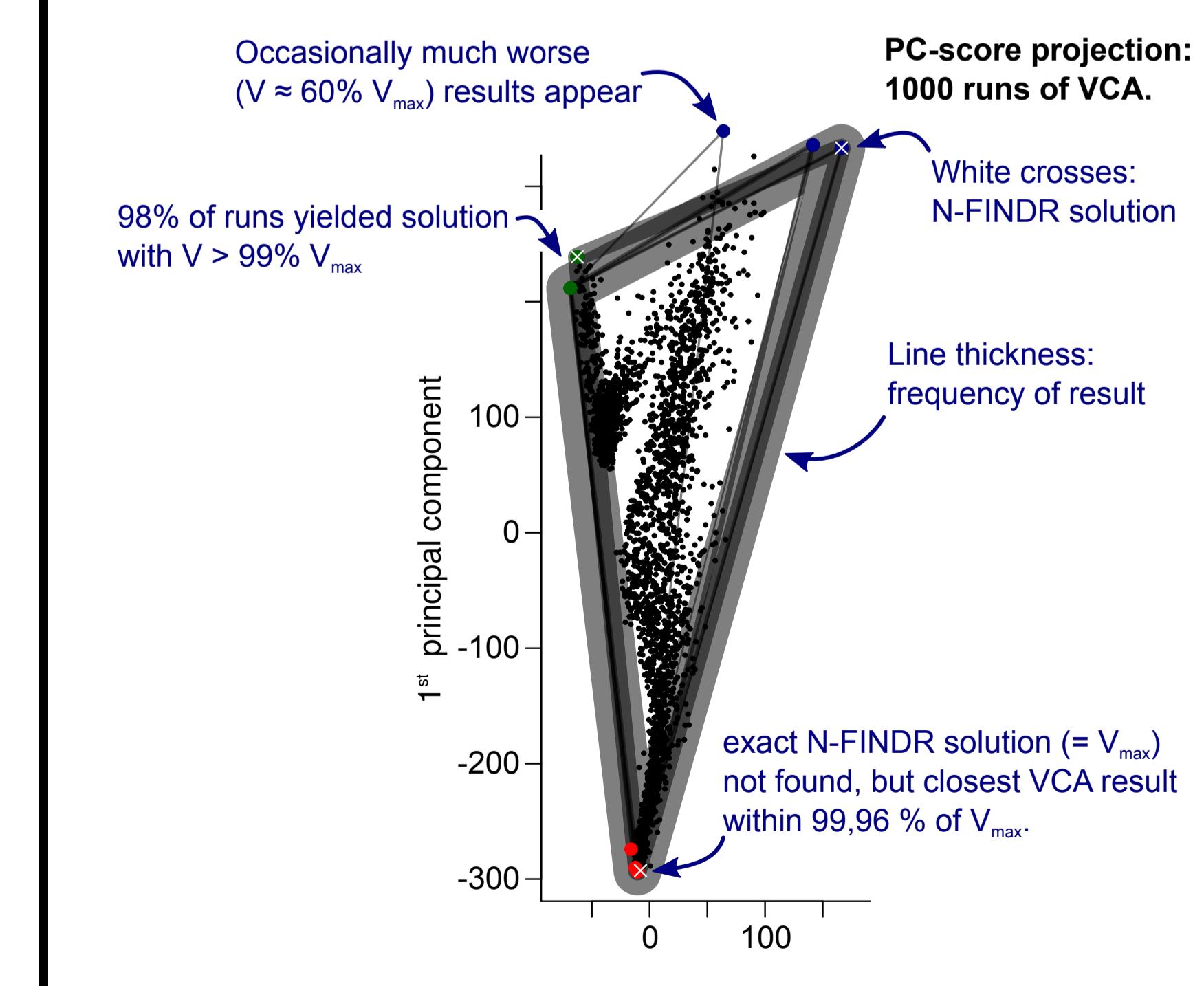
- Raman spectra of HeLa cell
- Excitation: 5 mW @ 488 nm, 0.5 s/spectrum
- Spectra: 600 – 1800 + 2800 – 3075 cm⁻¹, 314 bands (after pre-processing)
- Area:  $60 \times 60 \mu\text{m}$ , step size  $0.5 \mu\text{m}$
- For details see reference [HeLa Cell].

### N-FINDR with $m = 3$ endmembers



- Solution is stable: Identical results for 100 runs with random initialization

### VCA Results $m = 3$ endmembers



- VCA is expected to be less stable than N-FINDR: no refinement of tentative vertices
- VCA faster than Winter's N-FINDR, but advantage small for improved algorithms.

Conor McManus implemented N-FINDR [Winter, Dowler] and VCA [Nascimento, Lopez] algorithms as R package `unmixR`. He was supervised by Claudia Beleites, Simon Fuller and Bryan Hanson.

Claudia Beleites now maintains the package with help by Bryan Hanson.

The package is available at <http://github.com/Chathurga/unmixR>

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