## **General Overview**

The program phasor\_calculations.m is a supplement to the publication "Chemical complexity of the retina addressed by novel phasor analysis of unstained multimodal microscopy." [CHEMPH\_111091] In that publication, we introduced the super-phasor unmixing (SPU) method to unmix seven spectroscopic signals in the spectral range 300-690 nm. The performance of SPU was found to be significantly superior to fully constrained linear spectral unmixing especially in regions of high spectral overlap. This program can serve as a tool for implementing the super-phasor analysis. The reading of [CHEMPH\_111091] is highly encouraged before implementing phasor\_calculations.m as that publication provides much greater detail and context than these instructions. All equation numbers in these instructions reference [CHEMPH\_111091].

phasor\_calculations.m has the ability to do two things. The first is to convert spectral coordinates into phasor coordinates (eqs. 1 and 2) and perform a simple phasor analysis to find percent compositions (eq. 3). This function works with two and three pure components. The second function of phasor\_calculations.m is to perform super phasor analysis (eq 4). This function works with two and three super-phasors. Further details on the program's variables are addressed in Variable Specifications. A pseudo-run of the program for a seven component analysis is provided in Implementation.

# Variable Specifications

[phasor\_coords, percentages] = phasor\_calculations(pure\_spectra, exp\_spectrum, lambda\_min, lambda\_max)

## Input Variables

<u>pure spectra</u>: An n x 4, n x 6, 2 x 2 or 2 x 3 matrix used to hold the flourescence spectra expected to combine to form the mixed experimental spectrum.

- An n x 4 matrix should be used when two pure spectra are involved and an n x 6 matrix should be used for three pure spectra. These matrices should take the form of  $[\lambda, I, \lambda, I, \lambda, I]$  across the 4-6 columns with n rows.
- The 2 x 2 and 3 x 2 matrices are used for super phasor analysis where each index represents super-phasor G and S coordinates. These matrices should be of the form [G, S] and have two columns with each row containg a set of super-phasor coordinates.

<u>exp\_spectrum</u>: A m x 2 matrix used to hold the measured mixed flourescence spectrum. This matrix should take the form  $[\lambda, I]$  with m rows.

*lambda\_min*: the minimum wavelength for calculations

lambda\_max: the maximum wavelength for calculations

### **Output Variables**

<u>phasor\_coords</u>: a 4 x 2 or 3 x 2 matrix that returns the various calculated phasor coordinates. The columns in this matrix are of the format [G, S]. The first 2-3 rows are the phasor coordinates for the pure spectra and are returned according to their input order in the variable pure\_spectra. The last row is always reserved for the set of phasor coordinates belonging to exp\_spectrum.

<u>percentages</u>: a  $3 \ge 1$  or  $2 \ge 1$  matrix that returns the percent composition associated with each pure component. The order is once again determined by the order in pure\_spectra.

#### **Implementation**

spec1 =	$[\lambda_a]$	Ia	$\lambda_b$	$I_b$	$\lambda_c$	$I_c$ ]	(nx6)
spec2 =	$[\lambda_d]$	I <sub>d</sub>	$\lambda_e$	$I_e$ ]			(nx4)
spec3 =	$[\lambda_f]$	$I_f$	$\lambda_h$	$I_h$ ]			(nx4)

 $expSpec = \begin{bmatrix} \lambda_z & I_z \end{bmatrix}$ (nx2)

(coords1, percent1) = phasorcalculations(spec1, expSpec, 300, 500)

$$coords1 = \begin{bmatrix} a_a & s_c \\ G_b & s_l \\ G_c & s_c \\ G_{z1} & s_{z2} \end{bmatrix}$$
$$percent1 = \begin{bmatrix} a \\ b \\ c \end{bmatrix}$$

(coords2, percent2) = phasorcalculations(spec2, expSpec, 500, 650)

$$coords2 = \begin{bmatrix} G_d & S_d \\ G_e & S_e \\ G_{22} & S_{22} \end{bmatrix}$$
$$percent2 = \begin{bmatrix} d \\ e \end{bmatrix}$$

(coords3, percent3) = phasorcalculations(spec3, expSpec, 650, 800)  $coords3 = \begin{bmatrix} G_f & S_f \\ G_h & S_h \\ G_{z3} & S_{z3} \end{bmatrix}$   $percent3 = \begin{bmatrix} f \\ h \end{bmatrix}$   $super = \begin{bmatrix} G_{abc} & S_{abc} \\ G_{de} & S_{de} \\ G_{fh} & S_{fh} \end{bmatrix}$ 

 $G_{abc} = a \cdot G_a + b \cdot G_b + c \cdot G_c$  \*the other G and S values follow the same pattern

\*the formation of *super* is not included in the program and must be done by the user

(coords4, percent4) = phasorcalculations(super, expSpec, 300, 800)

$$coords4 = \begin{bmatrix} G_{abc} & S_{abc} \\ G_{de} & S_{de} \\ G_{fh} & S_{fh} \\ G_{z4} & S_{z4} \end{bmatrix}$$
$$percent4 = \begin{bmatrix} abc \\ de \\ fh \end{bmatrix}$$

 $total contributions_a = abc \cdot a$  \* the other total contribution values follow the same pattern.