

# Tutorial : Hyperspectral Analysis on 3D SIM/4D LSM Image Data

## Introduction

This tutorial is to help users analyze spectral information of granules (such as, lipofuscin, melano-lipofuscin, and etc.) with 3D high-resolution structured illumination microscopy (SIM) and 4D confocal multispectral laser scanning microscopy (LSM).

The required programs to follow the procedure in this tutorial are

- 1) 3D Slicer – <http://www.slicer.org>
- 2) FIJI – <https://fiji.sc>
- 3) ITK-SNAP – <http://www.itksnap.org>

## Workflow

We will describe the overall workflow of hyperspectral analysis, starting from 1) reading 4D LSM image, 2) granule segmentation on 3D SIM data, 3) co-registration between 4D LSM and 3D SIM data, and 4) hyperspectral analysis of granules.

### 1. Read 4D LSM

### 2. Granule segmentation on 3D SIM

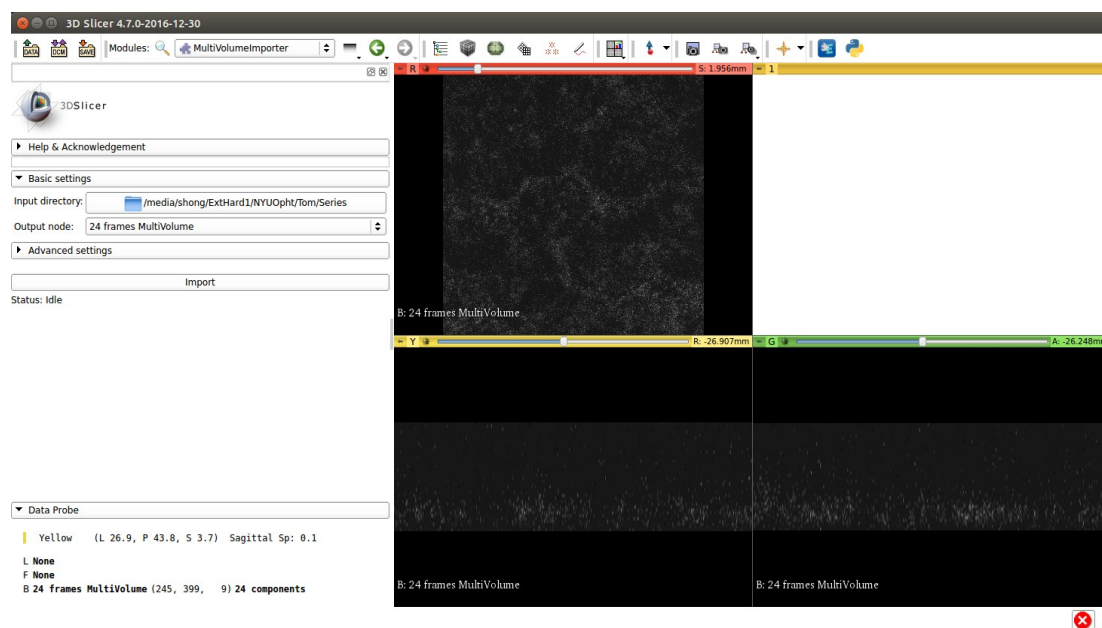
### 3. Co-Registration : SIM / LSM

Please see a previous tutorial for segmentation/registration processes using FIJI and ITK-SNAP

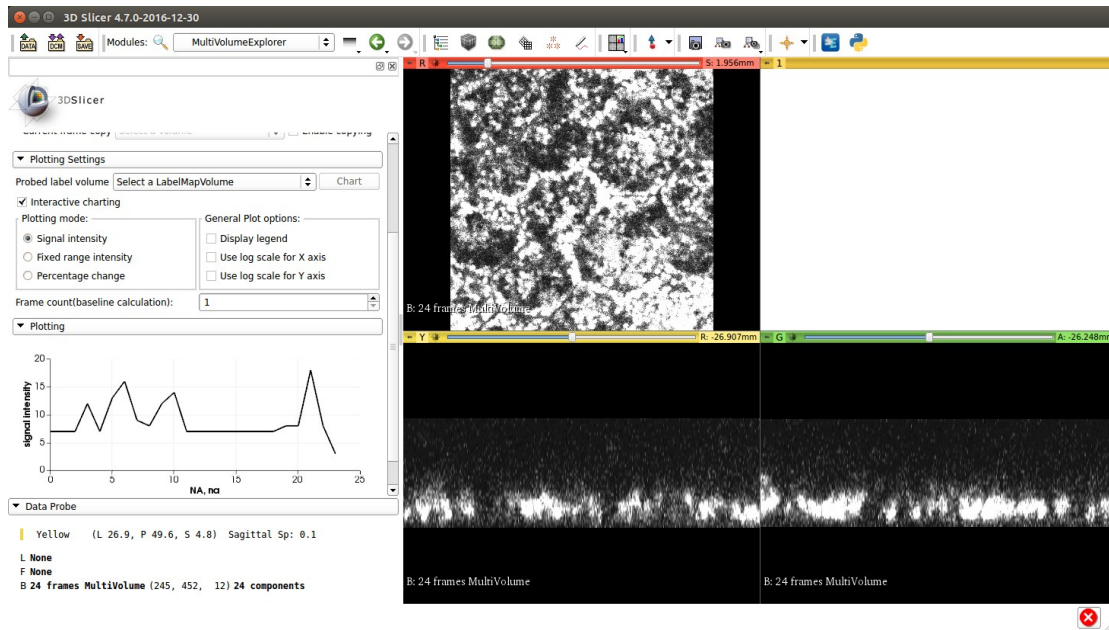
### 4. Convert concatenated 4D LSM to an image series

### 5. Hyperspectral Analysis

- 1) Open 3D Slicer
- 2) Go to Extension → MultiVolume Support → Multivolume Importer
- 3) Set Input Directory to the folder of an image series
- 4) Create new MRMLMultiVolume at Output node
- 5) Import (May take a few minutes)
- 6) Imported volume will be displayed as # frames MultiVolume ( e.g. 24 frames MultiVolume)



- 7) Go to Extension → MultiVolume Support → Multivolume Explorer
- 8) Adjust Window/Level by drag-n-drop on one of viewports
- 9) Set Input multiVolume to # frames MultiVolume (imported from Step 5))
- 10) Click Play next to Current frame number to play a video to display a sequence of image channels
- 11) The plot of hyperspectral information will be displayed on the left panel under “Plotting” while you navigate on the viewports.



- 12) Load a label map generated from the segmentation step in “Plotting Settings → Probed label Volume”
- 13) Click Chart next to “Probed label Volume”

