

Tutorial for analyzing data from dual-color ratiometric imaging using SMAP

Tested on Matlab 2021b.

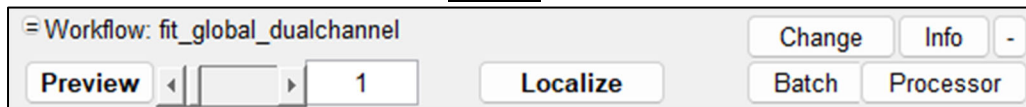
Future issues will be fixed in the develop branch at <https://github.com/jries/SMAP>.

Generate the PSF model (.h5) file using uiPSF.

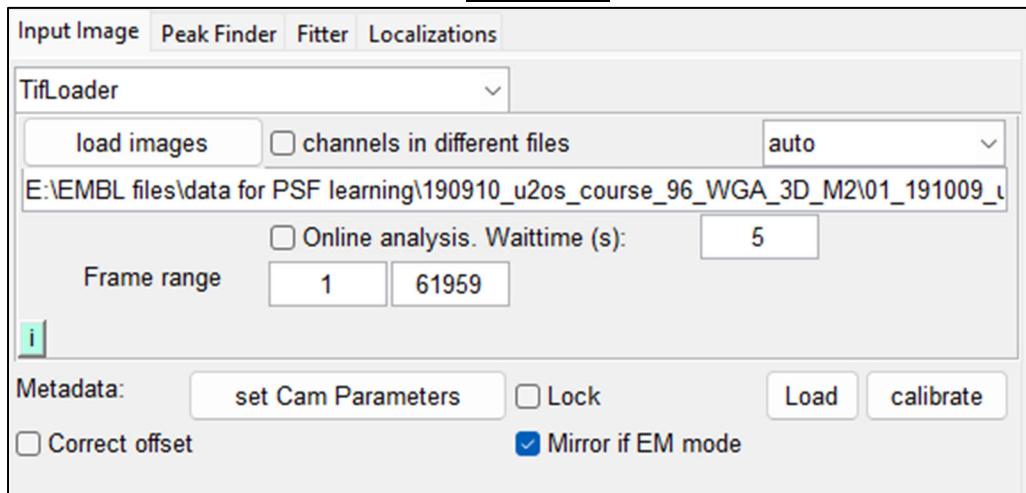
- If the data is in .tif format, set 'swapxy = true'. This is because there is a permutation in SMAP when loading .tif files.
- Set 'ref_channel = 0', this does depend on which channel SMAP consider as the reference channel, usually it should be set to zero.

Then follow the steps below for localization in SMAP.

1. Open MATLAB, set the working directory to C:\Users\Ries Lab\git\SMAP\
2. In Matlab command window, run: SMAP
3. In SMAP GUI, open tab Localize, click **Change** and select the file 'fit_global_dualchannel.txt'.



4. Go to tab Localize -> Input Image, click **load images** and select the first .tif file in the data folder.



5. Edit parameters defined in **set Cam Parameters** if metadata was not available. Sometimes, the bead data and the SMLM data are mirrored when they were collected at different EM mode settings, in this case, check 'Mirror if EM mode'. If insitu PSF model is used, uncheck 'Mirror if EM mode'.
6. Open tab Localize -> Peak Finder, click **load T**, select the .h5 file from uiPSF.

Input Image Peak Finder Fitter Localizations

DoG s:

☐ correct sCMOS variance

dynamic (factor)

☐ minimum distance (pix)

E:\EMBL files\data for PSF learning\bead data\190910_beads3Ddua

preview mode: image-bg

☐ frame dependent transformation

ROI to include

ROI to exclude

Clear ROIs

rectangle

☒ exclude rim

7. Open tab Localize -> Fitter, click **Load 3D cal**, select the same .h5 file. Check 'Global fit' and select 'ch1' for 'main x,y'. Check 'RI mismatch' if the PSF model was generated from bead data.

Input Image Peak Finder Fitter Localizations

Spline Iterations: ROI size (pix)

Load 3D cal or_M2\psfmodel_smap1_voxel_multi.h5 ROIs/fit:

☒ RI mismatch: ☒ Global fit

☐ pixelsize X,Y (um): main x,y: ch1

☐ Multi color: fix ratio to: Weights r t

☐ sCMOS correction ☐ get asymmetry

	I	x
x	<input checked="" type="checkbox"/> 1	
y	<input checked="" type="checkbox"/> 1	
z	<input checked="" type="checkbox"/> 1	
N	<input type="checkbox"/> 1	
Ra	<input type="checkbox"/> 1	

8. In Localize -> Peak Finder, click **Preview** and adjust 'dynamic (factor)' based on the output image.

Input Image Peak Finder Fitter Localizations

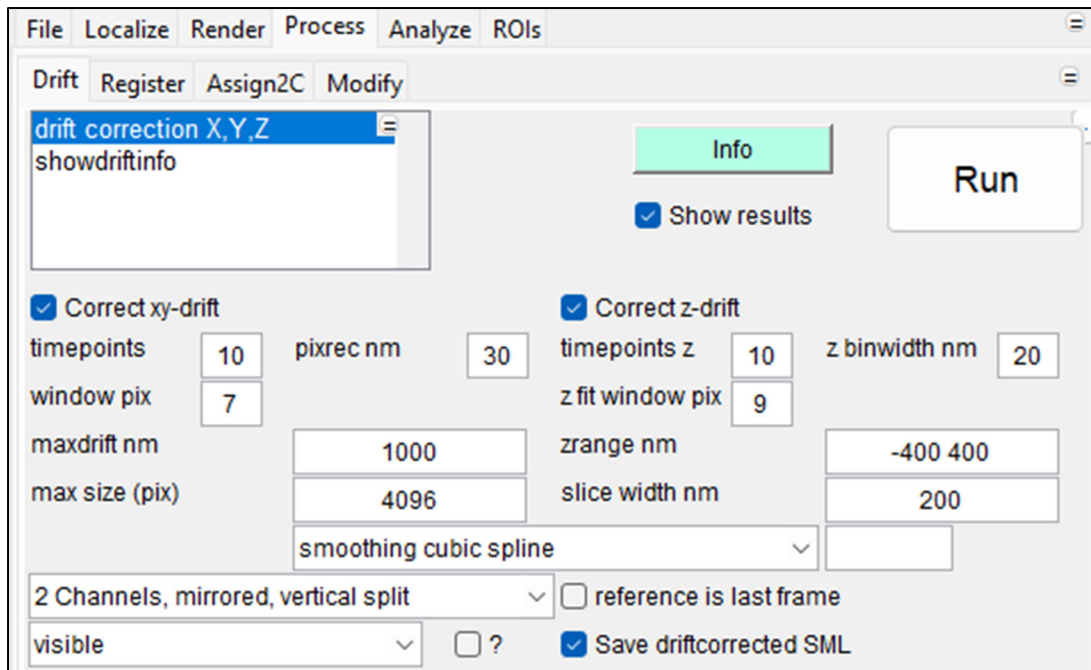
DoG s:

☐ correct sCMOS variance

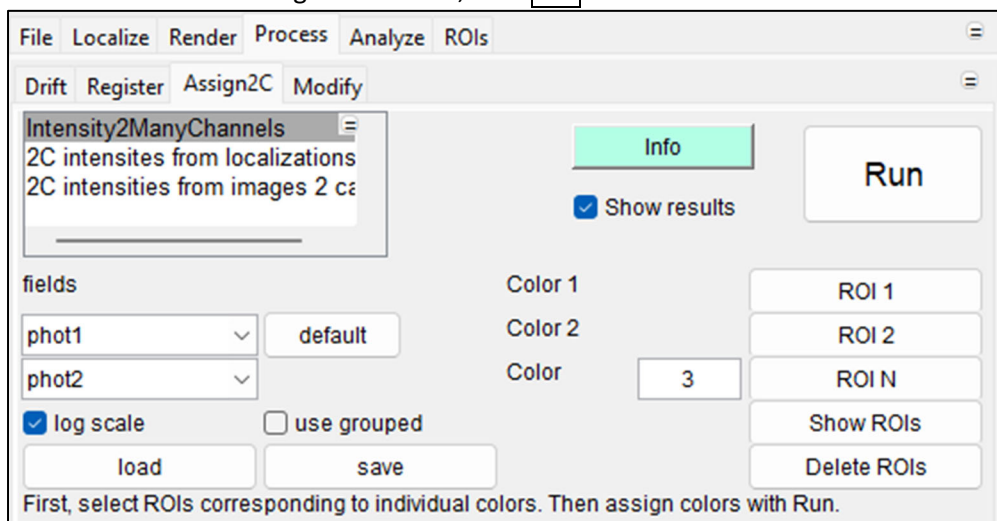
dynamic (factor)

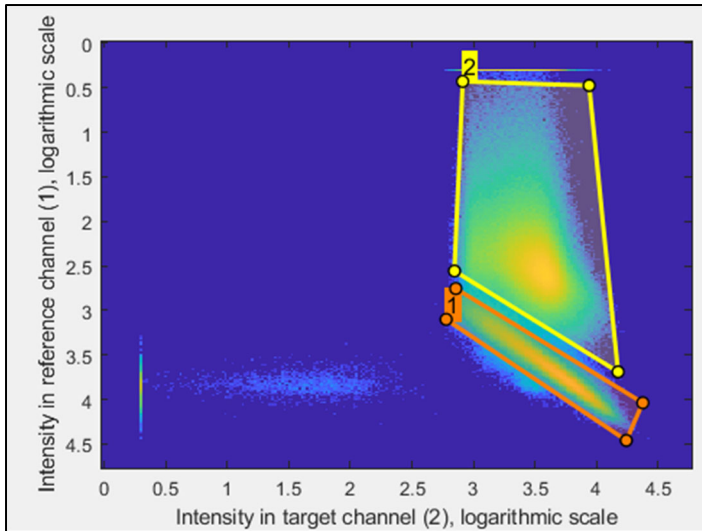
☐ minimum distance (pix)


9. Then click **Localize**.
10. After localization is finished, in panel 'format', click **Reset** at the upper right of the SMAP GUI.
11. Go to tab Process -> Drift, and set 'pixerec nm' to 30 and 'z binwidth nm' to 20. In the bottom dropdown menu, select correct splitting for your data, e.g. '2 Channels, mirrored, vertical split' Then click **Run**.

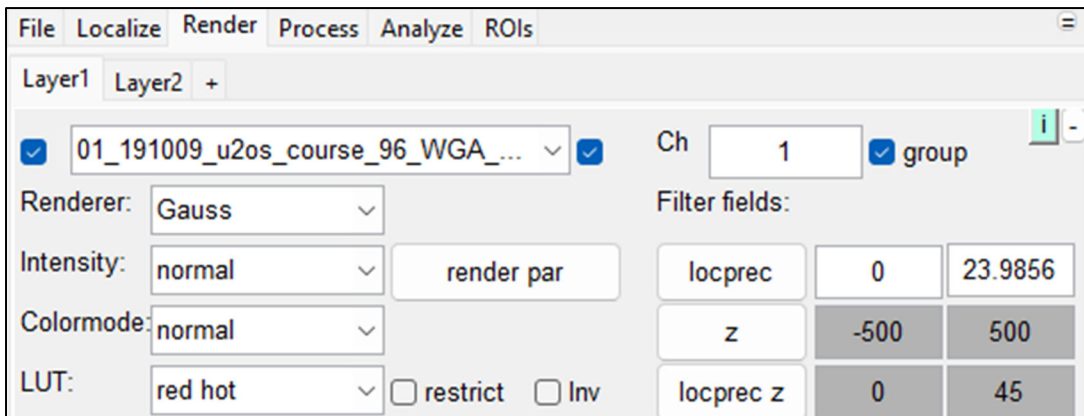


12. Go to tab Process -> Assign2C, select 'Intensity2ManyChannels'. Click **Show ROIs**, then click **ROI 1**, draw the region for color channel 1 in the popup Figure, then click **ROI 2**, draw the region for color channel 2. After region selection, click **Run**.





13. In tab Render, click  to create one more layer and set 'Ch' to 1 for Layer1 and to 2 for Layer2. For different layer, set 'LUT' to different colormap.



14. Now you can render a dual-color super-resolution image.