A step-by-step process for the axial to radial transformation of 2D localizations for free Janelia Fluor $^{\rm @}$ 646 (JF646) dye.

Section Outline

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Section I

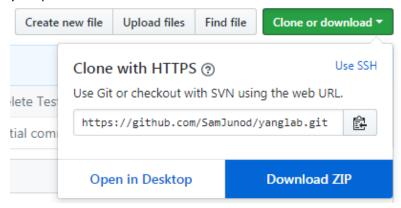
Software Download

Python 3

- 1) Install Python 3
 - i. Since these scripts make use of several Python 3 libraries (tkinter, csv, random, os, sys, numpy, scipy, and math), the simplest way to install Python 3 with these required libraries is through the Anaconda distribution of Python 3 which can be found here (https://www.anaconda.com/download/).
 - ii. Download the appropriate installer for your operating system (32-bit or 64-bit Windows or Mac)
 - iii. Once the installer has finished downloading, open it and follow the click-through instructions to install the Anaconda distribution of Python 3. An example of the first step of the installer for a 64-bit Windows system is shown below.



- 2) Download the script files
 - i. Follow the URL link below.
 - i. https://github.com/SamJunod/yanglab
 - ii. Select the "Clone or Download" button and download the ZIP file. Extract the ZIP file to your preferred file location.



MatLab

- 1) Go to MatLab website, MATLAB MathWorks MATLAB & Simulink, and download the MatLab software.
 - i. URL: https://www.mathworks.com/products/matlab.html

OriginLab

- 2) Go to OriginLab website and download the Origin software.
 i. URL: https://www.originlab.com/demodownload.aspx

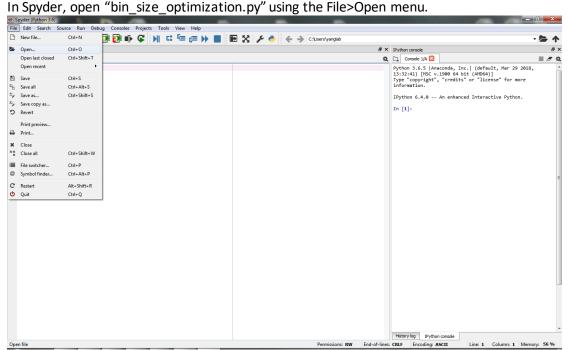
Section II

Bin Optimization

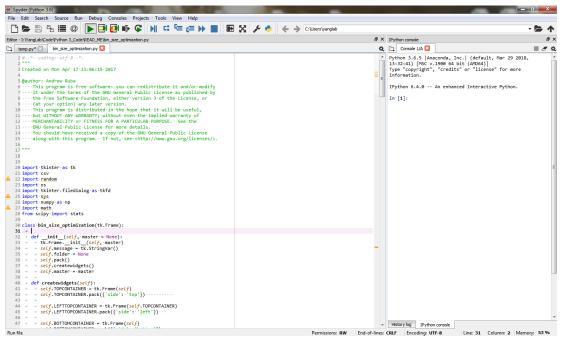
1) Packaged in the Anaconda distribution of Python 3 is an integrated development environment for Python 3 called "Spyder." Open Spyder- it should have an icon on your desktop or an icon in the program list under the start menu for Windows.



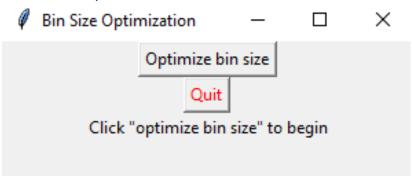
2) In Spyder, open "bin_size_optimization.py" using the File>Open menu.



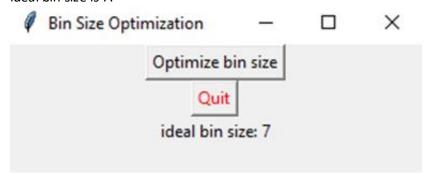
3) Then click the green "Run file" arrow in the toolbar.



4) Once the script begins to run, a pop-up window will appear. Select the Optimize bin size button and then select your data in csv format.



5) After the data is selected the script will run and output the ideal bin size. For JF646 dye, the ideal bin size is 7.

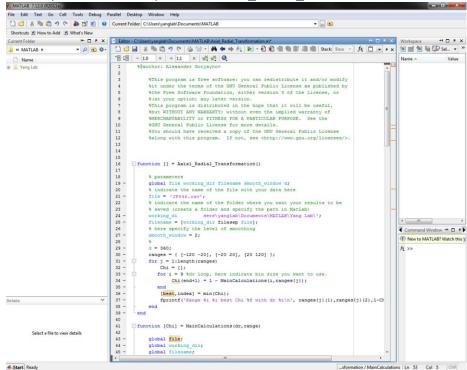


6) Select the "Quit" button to close the pop-up window.

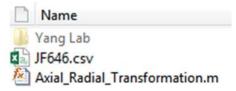
Section III

Axial to Radial Transformation

1) Start-up MatLab and Open the "Axial Radial Transformation.m" file.



2) The MatLab script (Axial_Radial_Transformation.m) and the 2D localizations (JF646.csv) will have to be in the same folder.



- 3) In the MatLab script, under the Editor box, make the following edits:
 - i. File name (line 21, file = 'file name.csv';)

ii. Working directory(line 24, working_dir = 'folder location of data';)

```
% indicate the name of the folder where you want your results to be
% saved (create a folder and specify the path in MatLab)
24 - working_dir = 'C:\Users\yanglab\Documents\MATLAB\Yang Lab\';
```

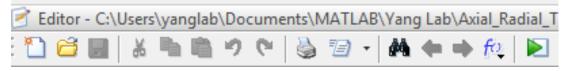
iii. Axial ranges (line 30, ranges = { [axial ranges], [for data] };

```
30 - ranges = { [-120 -20], [-20 20], [20 120] };
```

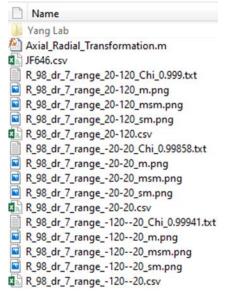
iv. Binsize (line 33, for i = **optimal bin size**). This is the bin size determined from section II.

```
33 - for i = 7 %dr loop, here indicate bin size you want to use.
```

4) Select "Save" and run the script by selecting the green arrow in the toolbar under the "Editor" box.



5) The program will write several items to the folder selected in the MatLab script.



- 6) Open the csv files created by the program. The values within each of the csv file are the raw results from the MatLab script. The file name of the csv document corresponds to the total radius of the program, the bin size(dr), and the range of the axial segment.
- 7) The csv file will have 3 columns of radial information (histogram count, raw matrix equation values, and smoothed matrix equation values). Copy the third column (smoothed matrix equation values) to a separate excel file.

1	Α	В	С	A	А	В	С
1	47.333	2.8633	2.8633	1	(-120 to -20)	(-20 to 20)	(20 to 120)
2	47.333	2.0464		2			
				3	2.8633		
3	49	3.4173		4	2.7757		
4	49	3.9335	3.9997	5	3.1324		
5	47.667	4.6483	4.0714	6	3.9997		
6	43.667	3.6324	3.8972	7	4.0714		
7	41.333	3.4109	3.6004	8	3.8972		
8	38.667	3.758	3.1013	9	3.6004		
9	34	2.135	3.0399	10	3.1013		
10	34	3.2266	2.428	11	3.0399		
				12	2.428		
11	30.333	1.9224		13	2.9323		
12	31.667	3.648	2.9696	14	2.9696		
13	26.667	3.3384	4.1267	15	4.1267		
14	20	5.3936	2.9107	16	2.9107		
15	0	0	0	17	0		
				1.8			

8) Repeat the last step for the other axial ranges from the other excel files.

Α	В	С
(-120 to -20)	(-20 to 20)	(20 to 120)
2.8633	23.232	1.8556
2.7757	14.346	2.5687
3.1324	7.9862	3.3688
3.9997	3.5107	3.4927
4.0714	3.3281	4.092
3.8972	2.4	3.5678
3.6004	2.1932	4.2182
3.1013	1.1375	3.5082
3.0399	1.1346	3.7557
2.428	1.0058	3.4099
2.9323	0.95779	3.6367
2.9696	0.5119	3.486
4.1267	1.293	4.8753
2.9107	1.1762	3.5981
0	0	0
	(-120 to -20) 2.8633 2.7757 3.1324 3.9997 4.0714 3.8972 3.6004 3.1013 3.0399 2.428 2.9323 2.9696 4.1267 2.9107	(-120 to -20) (-20 to 20) 2.8633 23.232 2.7757 14.346 3.1324 7.9862 3.9997 3.5107 4.0714 3.3281 3.8972 2.4 3.6004 2.1932 3.1013 1.1375 3.0399 1.1346 2.428 1.0058 2.9323 0.95779 2.9696 0.5119 4.1267 1.293 2.9107 1.1762

9) These values will have to be normalized to the number of bins in each subsection. For instance, with a bin size of 7nm the section that ranges from -120nm to -20nm is comprised of 14.29 subregions (bins). This value is calculated by taking the number the length of the axial segment (100nm) and dividing by the bin size (7nm). Likewise, the subsection that ranges from -20nm to 20nm is comprised of 5.71 subregions, and the subsection that ranges from 20nm to 120nm is comprised of 14.29 subregions. The values in the -120nm to -20nm column will be divided by 100/7. The data from -20nm to 20nm is divided by 40/7, and the data from 20nm to 120nm is divided by 100/7.

(-120 to -20)	(-20 to 20)	(20 to 120)
0.200431	4.0656	0.129892
0.194299	2.51055	0.179809
0.219268	1.397585	0.235816
0.279979	0.6143725	0.244489
0.284998	0.5824175	0.28644
0.272804	0.42	0.249746
0.252028	0.38381	0.295274
0.217091	0.1990625	0.245574
0.212793	0.198555	0.262899
0.16996	0.176015	0.238693
0.205261	0.16761325	0.254569
0.207872	0.0895825	0.24402
0.288869	0.226275	0.341271
0.203749	0.205835	0.251867
0	0	0

10) Find the maximum value of the whole data set.

	Max	4.0656
0	0	0
0.203749	0.205835	0.251867
0.288869	0.226275	0.341271
0.207872	0.0895825	0.24402
0.205261	0.16761325	0.254569
0.16996	0.176015	0.238693
0.212793	0.198555	0.262899
0.217091	0.1990625	0.245574
0.252028	0.38381	0.295274
0.272804	0.42	0.249746
0.284998	0.5824175	0.28644
0.279979	0.6143725	0.244489
0.219268	1.397585	0.235816
0.194299	2.51055	0.179809
0.200431	4.0656	0.129892
(-120 to -20)	(-20 to 20)	(20 to 120)
(-120 to -20)	(-20 to 20)	(20 to 120)

11) Normalize the values in each column with the maximum value by dividing each value by the maximum value. For JF646, this maximum value is 4.0656.

(-120 to -20)	(-20 to 20)	(20 to 120)
0.04929924	1	0.031949
0.04779098	0.61751033	0.0442269
0.05393251	0.34375861	0.0580028
0.06886536	0.15111484	0.060136
0.07009986	0.14325499	0.0704545
0.06710055	0.10330579	0.0614291
0.06199036	0.09440427	0.0726274
0.05339704	0.04896264	0.0604029
0.05233988	0.04883781	0.0646643
0.04180441	0.04329373	0.0587104
0.05048726	0.04122719	0.0626154
0.05112948	0.02203426	0.0600207
0.071052	0.05565599	0.0839411
0.05011536	0.05062844	0.0619508
0	0	0

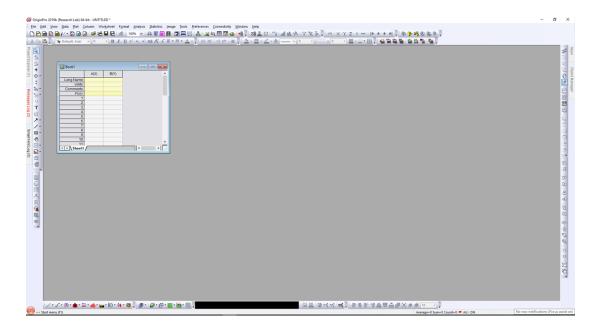
12) Then add a radial dimension column "R (nm)". This column will start at half the bin size then increase by the bin size. For JF646, with an optimal bin size of 7, the column will start at 3.5 nm then increase by 7 nm for each density measurement.

R (nm)	(-120 to -20)	(-20 to 20)	(20 to 120)
3.5	0.04929924	1	0.031949
10.5	0.04779098	0.6175103	0.0442269
17.5	0.05393251	0.3437586	0.0580028
24.5	0.06886536	0.1511148	0.060136
31.5	0.07009986	0.143255	0.0704545
38.5	0.06710055	0.1033058	0.0614291
45.5	0.06199036	0.0944043	0.0726274
52.5	0.05339704	0.0489626	0.0604029
59.5	0.05233988	0.0488378	0.0646643
66.5	0.04180441	0.0432937	0.0587104
73.5	0.05048726	0.0412272	0.0626154
80.5	0.05112948	0.0220343	0.0600207
87.5	0.071052	0.055656	0.0839411
94.5	0.05011536	0.0506284	0.0619508
101.5	0	0	0

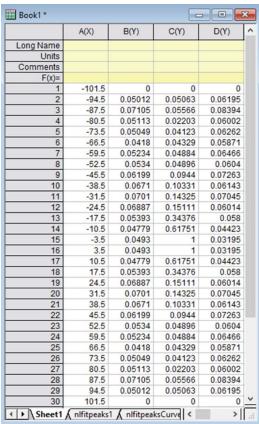
13) Then create a mirror image for each column, for the negative R (nm) rows.

R (nm)	(-120 to -20)	(-20 to 20)	(20 to 120)
-101.5	0	0	0
-94.5	0.05011536	0.0506284	0.0619508
-87.5	0.071052	0.055656	0.0839411
-80.5	0.05112948	0.0220343	0.0600207
-73.5	0.05048726	0.0412272	0.0626154
-66.5	0.04180441	0.0432937	0.0587104
-59.5	0.05233988	0.0488378	0.0646643
-52.5	0.05339704	0.0489626	0.0604029
-45.5	0.06199036	0.0944043	0.0726274
-38.5	0.06710055	0.1033058	0.0614291
-31.5	0.07009986	0.143255	0.0704545
-24.5	0.06886536	0.1511148	0.060136
-17.5	0.05393251	0.3437586	0.0580028
-10.5	0.04779098	0.6175103	0.0442269
-3.5	0.04929924	1	0.031949
3.5	0.04929924	1	0.031949
10.5	0.04779098	0.6175103	0.0442269
17.5	0.05393251	0.3437586	0.0580028
24.5	0.06886536	0.1511148	0.060136
31.5	0.07009986	0.143255	0.0704545
38.5	0.06710055	0.1033058	0.0614291
45.5	0.06199036	0.0944043	0.0726274
52.5	0.05339704	0.0489626	0.0604029
59.5	0.05233988	0.0488378	0.0646643
66.5	0.04180441	0.0432937	0.0587104
73.5	0.05048726	0.0412272	0.0626154
80.5	0.05112948	0.0220343	0.0600207
87.5	0.071052	0.055656	0.0839411
94.5	0.05011536	0.0506284	0.0619508
101.5	0	0	0

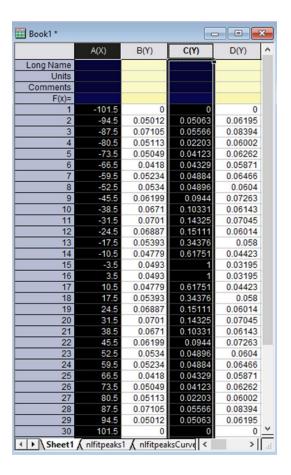
14) To determine the peak density at different radial distances, you will then transfer the data set into Origin.



15) Copy the dataset into a new worksheet. With the radial dimension values as (X) and the density values as (Y).



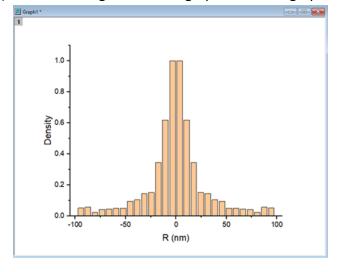
16) Select the column containing the radial dimension values (For this example, A(X)) and then while holding the "Ctrl" button on the keyboard select one of the density value columns (For this example, B(Y), or C(Y), or D(Y)).



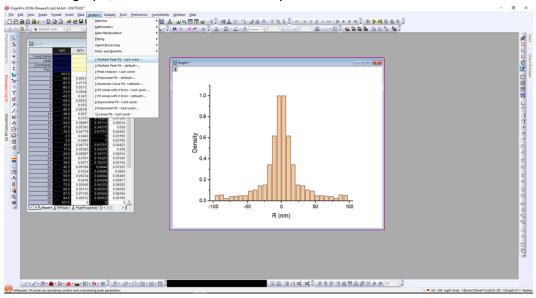
17) Then select the red vertical column graph button on the toolbar at the bottom of the Origin program.



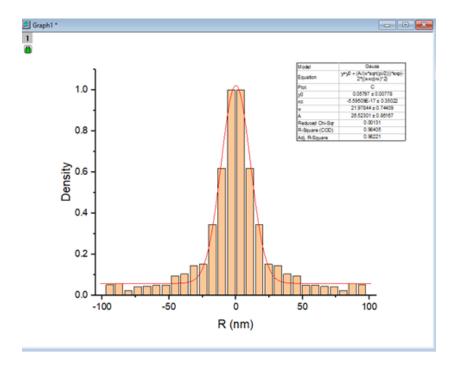
18) After selecting the vertical graph column a graph will appear.



19) Select the graph, then on the toolbar Analysis>Fit Gaussian



20) Origin will fit a Gaussian to the selected graph. The xc will be the mean of the peak density distribution. This value is the transport route with largest density distribution within that axial range. The density distribution of JF646, within the axial range of -20nm to 20nm will produce a peak mean of -5.59509E-17 ± 0.35022.



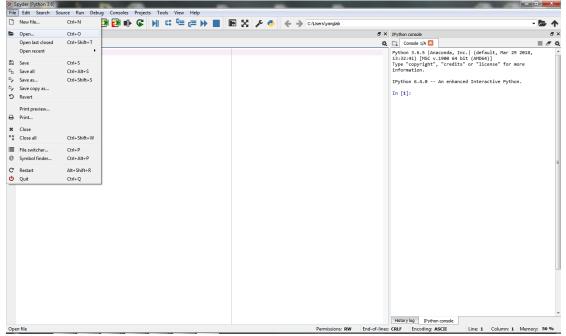
Section IV

Point Number Simulation

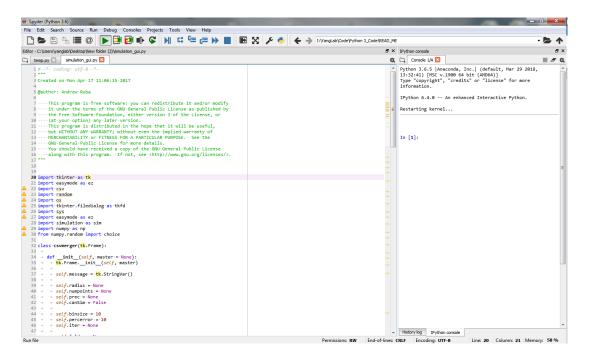
1) Similarly to section II, Open Spyder – it should have an icon on your desktop or an icon in the program list under the start menu for Windows.



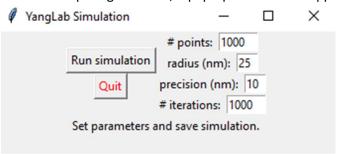
2) Once Spyder starts up, open "simulation_gui.py" using the File>Open menu.



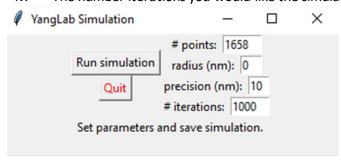
3) Once "simulation_gui.py" is open, click the green "Run file" arrow in the toolbar.



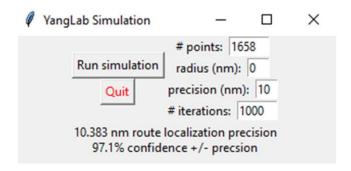
4) Once the script begins to run, a pop-up window will appear.



- 5) Input the following values;
 - i. The number of points (# points:).
 - ii. The peak radius found from Origin as explained in section III (radius (nm):).
 - iii. The precision of the your single-molecule localizations (precision (nm):).
 - iv. The number iterations you would like the simulation to run (# iterations:).



6) Click "Run simulation" after entering the proper integer values for the simulation parameters you would like to run. The optimal bin size will be dynamically calculated according to the parameters and the reproducibility rate and route localization precision will be written into the gui message area.



7) Select the "Quit" button to close the pop-up window.