# Decode-PAINT: 9-color decoding sample dataset user guide

# **Before starting analysis**

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- 1. Download Picasso following instructions (as of 10/10/2022, ver 0.4.11)
  - https://github.com/jungmannlab/picasso
- 2. Modify Picasso Render source code
  - o picasso>gui>render.py (NOT render.py in
  - In def align(self)

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sp = lib.ProgressDialog("Shifting channels", 0, len(self.locs), self)
sp.set_value(0)
for i, locs_ in enumerate(self.locs):
locs_.y == shift[0][i]
```

```
self.update_scene()
else:
    max_iterations = 4
    convergence = 0.001 # Thhat is 0.001 pixels ~0.13nm
    shift_x = []
    shift_y = []
shift_z = []
    display = False
    # 10/10/2022 added
    shift_x_combined = []
    shift_y_combined = []
    progress = lib.ProgressDialog("Aligning images...", 0, max_iterations, self)
    progress.show()
    progress.set_value(0)
    for iteration in range(max_iterations):
        completed = "True
        progress.set_value(iteration)
        shift = self.shift_from_rcc()
        sp = lib.ProgressDialog("Shifting channels", 0, len(self.locs), self)
        sp.set_value(0)
```

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shift_y.append(np.mean(temp_shift_y))
                              if len(shift) == 3:
                                  shift_z.append(np.mean(temp_shift_z))
                               iteration += 1
                               self.update_scene()
                              # 10/10/2022 added
                              if len(shift_x_combined) > 0:
                                  shift_x_combined = [x1 + x2 \text{ for } (x1, x2) \text{ in } zip(shift_x_combined, temp_shift_x)]
                                  shift_y_combined = [y1 + y2 for (y1, y2) in zip(shift_y_combined, temp_shift_y)]
                              else:
                                  shift_x_combined = temp_shift_x
                                  shift_y_combined = temp_shift_y
                              # Skip when converged:
                              if completed:
                                  break
                          progress.close()
                          # Plot shift etc
                          if display:
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                                   plt.subplot(1, 1, 1)
                                   plt.plot(shift_x, "o-", label="x shift")
plt.plot(shift_y, "o-", label="y shift")
plt.xlabel("Iteration")
                                   plt.ylabel("Mean Shift per Iteration (Px)")
                                   plt.legend(loc="best")
                                   fig1.show()
                              # 10/10/2022 added
                              for i in range(len(shift_x_combined)):
                                   print(shift_x_combined[i], end=',')
                              print('\n')
                              for j in range(len(shift_y_combined)):
                                   print(shift_y_combined[i], end=',')
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                              print('\n')
                    @check pick
                     def combine(self):
                         channel = self.get_channel()
                         picked_locs = self.picked_locs(channel, add_group=False)
                         out_locs = []
                         r_max = 2 * max(
                              self.infos[channel][0]["Height"], self.infos[channel][0]["Width"]
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```

In def \_undrift\_from\_picked\_coordinate(self, channel, picked\_locs, coordinate):

```
drift_mean = np.ma.average(drift, axis=0, weights=1 / msd)
drift_mean = drift_mean.filled(np.nan)

# Linear interpolation for frames without localizations
def nan_helper(y):
    return np.isnan(y), lambda z: z.nonzero()[0]

nans, nonzero = nan_helper(drift_mean)
drift_mean[nans] = np.interp(nonzero(nans), nonzero(nans), drift_mean[nans])

# 10/10/2022 added
drift_init = drift_mean[0]
drift_mean -= drift_init

return drift_mean

def _undrift_from_picked(self, channel):
    picked_locs = self.picked_locs(channel)
status = lib.StatusDialog("Calculating drift...", self)

drift_x = self._undrift_from_picked_coordinate(channel, picked_locs, "x")
drift_y = self._undrift_from_picked_coordinate(channel, picked_locs, "y")
```

- These modification allows Picasso Render to:
  - Align/register localizations to the initial frame (frame 0)
  - Show XY registration shift coordinates
- Install picasso

> python setup.py install

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## Sample 1: 9-color decoding (Fig. 1n)

#### Step 1. Process raw localization data

- Raw image file: 190509 X r1 021.nd2
  - The file is a concatenation of multiple z stacks:
    - Frames 1–100: z = 0 nm (glass surface)
    - Frames 101–10100: z = 0 nm
    - Frames 10101–10200: z = 0 nm
    - Frames 10201–20200: z = 170 nm
    - Frames 20201–20300: z = 0 nm
    - Frames 20301–30300: z = 340 nm
    - Frames 30301-30400: z = 0 nm
    - Frames 30401–40400: z = 510 nm
    - . ...
    - Frames 90901–91000: z = 0 nm
    - Frames 91001–101000: z = 1530 nm
  - The file needs to be split into z stack series for localization with Picasso Localize
- 1. Process raw images with Picasso Localize to localize single molecules
  - Box Size: 17
  - Convergence Criterion: 0

- o Max. Iterations: 0
- o Min. Net Gradient: 7500
- PSF file for astigmatism 3D analysis: 190509\_psf.yaml
- Output files: ./locsfile/raw\_locs/190509\_X\_r1\_0##\_locs.hdf5
  - 190509\_X\_r1\_001, 003, 005, ...: Imaging fiducial markers on the surface for further registration
  - 190509\_X\_r1\_002, 004, 006, ... : Actual localization data in 170 nm steps
- 2. Register the surface fiducial marker localization files (190509\_X\_r1\_001, 003, 005, ...\_locs.hdf5) to generate a text file with drift registration coordinates
  - Open all surface fiducial marker localization files 190509\_X\_r1\_001, 003, 005,
     ...\_locs.hdf5 in Picasso Render

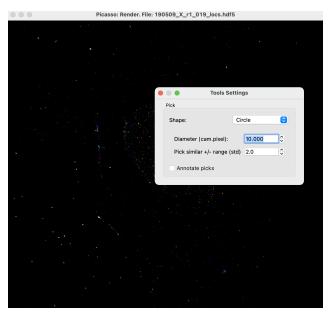


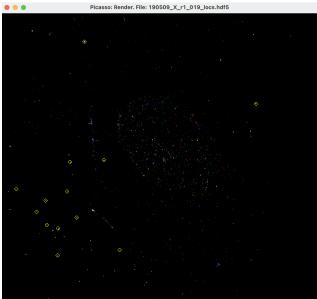
Apply Postprocess>Align channels (RCC or from picked) without any picks



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- Manually pick ~10 fiducial markers in the image using Tools>Pick
- Optional:) Use *Tools>Pick similar* to pick more fiducials automatically if there are many





- Apply Postprocess>Align channels (RCC or from picked) with picks
- Copy the last three lines, which are x, y, and z drift against the first file

Create a text file with the following lines with comma (,) delimitor: 190509 r1 drift.txt

- Line1, file names to be undrifted
- Lines 2–4, the three lines above: x, y, and z shift

```
190509_r1_drift.txt

1 190509_X_r1_002_locs,190509_X_r1_004_locs,190509_X_r1_006_locs,190509_X_r1_008_locs,190509_X_r1_010
    _locs,190509_X_r1_012_locs,190509_X_r1_014_locs,190509_X_r1_016_locs,190509_X_r1_018_locs,190509_X
    r1_020_locs
2 0.0,-2.65508419573,-4.96590882987,-3.72427673992,-5.11782836467,-3.37165220082,-2.4585662514,-1.867
    06850529,-2.83761903644,-3.69600827936
3 0.0,12.6123923302,16.9455328703,20.6623250723,23.9738042831,28.0911989331,29.6879745632,31.13522518
    87,29.3304789171,29.094566606
4 0.0,-41.8267650604,-51.8772878647,-81.8539729595,-91.797389257,-108.208583069,-90.1446902633,-90.88
    02564859,-70.562548542,-75.8468509197
```

- 3. Undrift localization files (190509\_X\_r1\_002, 004, 006...\_locs.hdf5) using undrift.py
  - Copy 190509\_r1\_drift.txt to the folder containing data to be undrifted: ./locsfile/raw\_locs/
  - Run undrift.py
    - > python undrift.py -f ./sample\_9col/locfiles/raw\_locs/190509\_r1\_drift.txt
  - Output files: ./locsfile/undrifted/190509 X r1 0## locs undrifted.hdf5
- 4. Further undrift each localization file using Picasso Render built-in function (*Postprocess>Undrift by RCC* and or *Postprocess>Undrift from picked*)
  - Output files: ./locsfile/aligned/190509\_X\_r1\_0##\_locs\_undrifted\_render.hdf5
- 5. Merge all files into one single 3D localization file using concatenate.py
  - Run concatenate.py
    - > python concatenate.py -d ./sample\_9col/locfiles/aligned/ -s 170 -o 190509\_X\_r1\_merged
  - Output file: 190509\_X\_r1\_merged.hdf5
- 6. Link localizations using Picasso Render built-in function (Postprocess>Link localizations)
  - Output file: 190509\_X\_r1\_linked.hdf5
- 7. Crop the localization data to 256 x 256 px using crop.py
  - Run crop.py
    - > python crop.py -f ./sample\_9col/locfiles/190509\_X\_r1\_linked.hdf5 -x 230 -y 220 -s 256
  - Output file: 190509\_X\_r1\_cropped.hdf5
- 8. Process localization data with HDBSCAN using hdbscanlocs.py
  - Run hdbscanlocs.py
    - Set HDBSCAN parameter 'minimum cluster size' to 30
    - > python hdbscanlocs.py -f ./sample\_9col/190509\_X\_r1\_cropped.hdf5 -c 30
  - Output file: ./hdbscan/hdbscan 30 None 0/190509 X r1 hdbscan 30 None 0.hdf5

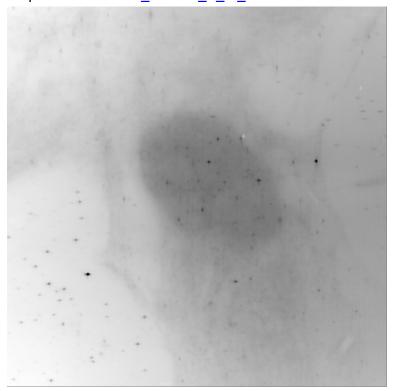
### Step 2. Process SABER diffraction limited image

Raw image file: ./diff/190509\_saber1\_r1.tif, 190509\_saber2\_r1.tif, 190509\_saber3\_r1.tif

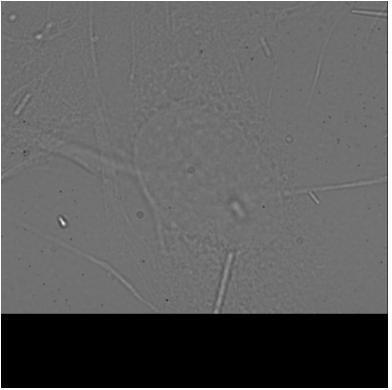
- o 5 channels: 488, 565, 647, DAPI, and bright field
- Registration reference file: 190509\_X\_r1\_021.nd2, Frame 1–100 (the first set of z = 0 nm (glass surface) image)
- 1. Generate a reference image for registration
  - Open ./diff/190509\_X\_r1\_021.nd2, Frame 1–100 with ImageJ/Fiji
  - Generate the average projection
  - Invert the intensity

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o Output file: ./diff/AVG\_190509\_X\_r1\_021.tif



- o The black spots outside of the cell are fiducial gold nanoparticles
- 2. Register SABER images to localization data based on fiducial markers visible in the bright field images
  - Open ./diff/190509\_saber1\_r1, r2, r3.tif and find the surface plane in the bright field channel



- Find a few gold nanoparticles observed in both bright field and registration reference
- o Register the SABER images based on fiducial markers manually or using Fiji Plugin
- Generate max Z projections
- Merge all SABER images into one single file: ./diff/190509\_X\_r1\_saber.tif
- 3. Generate a SABER binary mask file

images

- Crop 190509\_X\_r1\_saber.tif to align it to the localization file (256 x 256 px, origin = (220, 230))
- Apply Fiji Image>Adjust>Threshold>Yen
- o Output file: ./diff/190509 X r1 saber mask.tif

## Step 3. Decode localizations

- 1. Run decode.py
  - Input files:
    - Localization:
      ./hdbscan/hdbscan 30 None 0/190509 X r1 hdbscan 30 None 0.hdf5
    - SABER mask: ./diff/190509\_X\_r1\_saber\_mask.tif

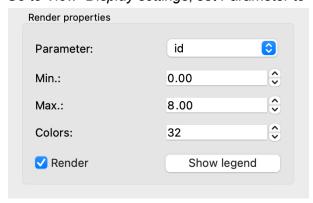
python decode.py -f
./sample\_9col/hdbscan/hdbscan\_30\_None\_0/190509\_X\_r1\_hdbscan\_30\_None\_0.hdf5 -m
./sample\_9col/diff/190509\_X\_r1\_saber\_mask.tif

Output file: ./hdbscan/hdbscan\_30\_None\_0/decoded/190509\_X\_r1\_hdbscan\_30\_None\_0\_decoded. hdf5

2. Open the decoded file in Picasso Render



Go to View>Display settings, set Parameter to 'id', and check Render



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