Chapter 2

3D Reconstruction

In this exercise we will use different image registration methods to build a 3D view from aligned slices, correct 3D drift, correct chromatic aberration and stitch a 3D mosaic. We will use different FIJI plugins and inspect the results with napari.

2.1 Alignment of Sequential Histological Sections

We will use the FIJI plugin stackreg[10] in this exercise. It uses the phase correlation based registration approach. It registers neighboring slices of the stack pairwise. The registration starts from the slice selected when the plugin is run and advances from there in both directions. The initially selected slice, called the anchor, is the one that will not be transformed.

Open the image rat-brain.tif. In order to make the 3D display easier, invert the contrast, then transfer the image to napari. You can see that the slices are not aligned.

In FIJI duplicate the stack, select the slice number 9 and run stackreg, using the rigid-body transformation. To find and run plugins you can use the ImageJ search-bar.

Compare the registered image to the original in napari. To transfer a second image to napari without replacing the first, you can use Get Labels instead of Get Image and then use the layers context menu to convert the labels to an image.

Align the original stack again with the rigid-body transformation, this time using slice 1 as anchor. What happens?

Align the stack using the affine transformation. What do you think about the result?



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Figure 2.1: Sequential histological sections unaligned, aligned with the rigidbody transformation and the affine transformation.

2.2 3D Drift Correction

In this exercise we will align a moving 3D object between different time-points. Open the image LargeDrift.tif[7]. This is a synthetic image that could represent a moving cell. Run the Correct 3D Drift script[5] in FIJI. As in the last exercise, Correct 3D Drift uses the phase correlation based registration approach. Compare the result with the original time-series image in napari.

2.3 3D Stitching

To have a large field of view and a high resolution in the same time we can take overlapping image stacks at a number of positions and stitch them together. You find 3 channels of the drosophila larva image[6] in the folders C1, C2 and C3. We will use the FIJI plugin Grid/Collection Stitching[6] to do the job. We will calculate the transformation on channel three, while stitching it in the same time and then apply the same transformation to stitch the two other channels.

 Run the Grid/Collection Stitching in FIJI. Select Unknown position



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Figure 2.2: Drift corrected and original time-series as a temporal-color-coded projection.

and All files in directory in the first dialog. Set the directory to the C3directory and make sure that Linear blending and Fuse and display are selected. Run the stitching and wait until the fused result image for channel 3 is displayed.

The plugin has written the transformations of the individual stacks to a file TileConfiguration.registered.txt in the C3-folder. Copy that file to the C1 and C2 folders. Open the copied files and replace C3 with C1 and C2 respectively. Run the Grid/Collection Stitching-plugin again. This time select Positions from file and Defined by TileConf in the first dialog. Uncheck the Compute overlap option in the second dialog and run the plugin on the C2 folder. When finished repeat the procedure for the C1 folder. You should now have the 3 stitched and merged images of the three channels open. Merge the channels in FIJI using Image>Color>Merge Channels.

We do not know the voxel size of the image and it is not in the image metadata. In order to nevertheless achieve a resonable 3D display, set the x and y size to 1 and the z size to 5 in FIJI via the Image>Properties menu. Transfer the image to napari to examine the result of the stitching.



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Figure 2.3: MIP projections of the individual drosophila larva stacks.



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Figure 2.4: The stitched mosaic of the drosophila larva stacks.