- Keywords: myenteric plexus, submucosal plexus, immunostain, immunofluorescence, enteric neuron, ileum, macrophage, T cell
- Note: The plexuses are fragile. At each wash, use a micropipette and a dissecting microscope to carefully remove the solution from the well not suck up the plexus.
- 1. Cut out two small regions of each plexus sheet in PBS at 4 degrees
- 2. Add each tissue section to separate wells in a 96 well dish.
- Add 10% normal donkey serum (Jackson Immunoresearch, Cat #017-000-121; West Grove, PA) and 1% Triton-X in PBS. Leave in blocking solution for 1 hour at room temperature on a rotator.
- Incubate the tissue overnight at room temperature on a rotator with primary antibodies in 10% normal donkey serum, 1% Triton-X in PBS.
 - a. Sections were stained with antibodies against tyrosine hydroxylase (TH) (1:500, Millipore-Sigma, Cat # AB152; Burlington, MA) and CD3 (1:400, Bio-Rad Laboratories, Cat #MCA2690; Hercules, CA), ANNA1 (1:32,000; kind gift by the Gershon laboratory (Margolis et al., 2016)), Iba1 (1:500, WAKO, Cat # 019-19741; Richmond, VA), CD68 (1:1000, Abcam, Cat # ab53444; Waltham, MA).
- 5. On Day 2, remove the primary antibody solution from each well and wash three times with PBS-Tween (0.1%) for 10 minutes each.
- 6. Incubate in secondary antibody (1:1000) in blocking solution for two hours at room temperature, covered on a rotator.
- 7. Wash again three times for 10 minutes each in 0.1% PBS-Tween
- 8. Mount on slide with vectashield medium with DAPI
- 9. Imaging:

- a. For imaging enteric neurons, for each plexus collect 2-3, 2x2 tile z-stack 640.17x
 640.17 μm confocal images at 20x magnification
- b. For imaging macrophages, for each plexus collect 2-3, 2x2 tile z-stack 390.09 x
 390.09 μm confocal images at 20x magnification

10. Analysis:

- Count the number of ANNA1⁺, TH⁺ cells, and IBA1⁺ cells were counted for each stacked image using Fiji
- b. Within the SP, threshold the TH⁺ signal, then analyze mean fluorescent intensity (MFI) and the area of the TH signal. Keep the thresholding consistent across each image, animal, and condition within each experiment
- c. Within each animal, sum the the number of ANNA1⁺, TH⁺ cells, and IBA1⁺ cells across all images separately then divide by the acquisition area.
- d. For each experiment, normalize to the CFA only condition.