

# Standard Operating Procedure Imaging RosetteArray<sup>™</sup> Well Plates

**Overview:** This document summarizes the microscope settings needed to:

- 1) Acquire Z-stack images of immunostained tissues within RosetteArray<sup>™</sup> plates/kits;
- 2) How to Use RosetteDetect<sup>™</sup>

#### **Equipment Required:**

- This SOP is written for a Nikon A1R Confocal Microscope, and it should be amended for other imaging platforms.
- Computer(s) with Image Capture Software, Internet Access, and spreadsheet processing software

#### **Materials Required:**

Immunostained 96-well RosetteArray<sup>™</sup> plate (DAPI (405 nm), N-Cadherin (488 nm), Pax6 (561 nm)) mounted with Glass Antifade Mountant Solution (Catalog #: P36984, ThermoFisher Scientific) and cultured using Neurosetta's <u>Standard Operating Procedure</u>

#### **Protocol Overview:**

- 1. Image Capture Settings/Parameters
- 2. How to Use RosetteDetect<sup>™</sup>

# **Protocol**:

### 1. Image Capture Settings/Parameters

High-quality inputs generate high-quality outputs. Clients should use a confocal microscope to capture Z-stack images of their RosetteArray<sup>™</sup> samples. Appropriate image capture settings/parameters are crucial for generating images that are amenable to RosetteDetect<sup>™</sup> processing and analysis.

- Using a negative control well, adjust the Z position of a 20x objective to get a resolved, digital view of a single rosette structure within a tissue of interest. Center the tissue within your frame. A violet corrected 20x objective is recommended but a normal 20x objective is sufficient to magnify your view of the tissue. You may use Perfect Focus to center your defined Z on the lumen within the tissue.
- 2. Set the digital frame zoom size to 2 (for forebrain tissues) or 3 (for spinal tissues), These values may be different for non-Nikon confocal microscope systems.
- 3. Set the digital frame resolution to 512x512 pixels.

- 4. Set the pinhole size to 1.5 2. This pinhole size may differ depending on the type of confocal microscope system you work with.
- 5. If you are using a resonance-like mode of acquisition, you may need to apply line averaging to your capture settings to get a resolved view of cell nuclei and N-Cadherin expression. For clients working with Nikon Confocal Microscope systems, I recommend 8x Line Averaging.
- Apply laser power and gain (photomultiplier tube sensitivity) values to your 405 nm, 488 nm, and 561 nm – or comparable PMT/channels-to produce a multi-signal fluorescent image without excessive pixel oversaturation.

NOTE: Users need to centralize tissues within the X and Y bounds of the capture frame (see Figure 1).

7. Generate a Z-stack image. The Z-stack image should feature 5 separate slices that span a Z range of 8-10 µms. For Forebrain tissues, the step height of the slices should be 2.5 ums. For spinal tissues, the step height of the slices may be 2-2.5 µms. Users should program their Z-stack acquisition tool (e.g., symmetric mode on Nikon A1R confocal microscopes) so that the third/center slice features the lumen of the rosette structure. The second and fourth adjacent slices may also feature portions of the rosette structure. RosetteDetect<sup>™</sup> supported file formats include: TIFF, ND2 (Nikon), LIF (Leica), OIB (Olympus), and CZI (Zeiss).

## 2. How to Use RosetteDetect<sup>™</sup>

Neurosetta's RosetteDetect<sup>™</sup> is a web-based image analysis software designed to provide

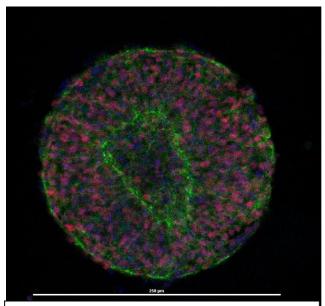


Figure 1: A 250-µm diameter forebrain tissue featuring a single neural rosette structure was captured with a violet-corrected, 20X objective (third slice). The tissue is centered in the capture frame and does not touch the boundaries of the image frame. N-Cadherin (green), Pax6 (red), Cell Nuclei (blue). Scale bar = 250 µms.

valuable insights into your RosetteArray<sup>™</sup> images. After imaging RosetteArray<sup>™</sup> plates using recommended procedures, simply upload your images into the <u>RosetteDetect<sup>™</sup> portal</u> and our Al-powered software will provide a tabulated readout of the relative cell count (nucleic acid stain), neural induction (Pax6 stain), and rosette formation and morphology (N-cadherin stain) for each micropatterned tissue. For more detailed instructions see Neurosetta's <u>Standard</u> <u>Operating Procedure</u>.