

Dec 19, 2023

Immunohistochemistry data processing

DOI

dx.doi.org/10.17504/protocols.io.x54v9p5omg3e/v1

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Protocol Citation: Tae-Un Han 2023. Immunohistochemistry data processing, protocols.io

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Protocol status: Working

We use this protocol and it's working

Created: December 19, 2023



Last Modified: May 31, 2024

Protocol Integer ID: 92522

Keywords: ASAPCRN, high throughput quantification of immunohistochemistry data, immunohistochemistry data, immunohistochemistry, mouse brain, high throughput quantification, data

Funders Acknowledgements:

ASAP

Grant ID: ASAP-000458

Abstract

This protocol has been used for high throughput quantification of immunohistochemistry data of mouse brain sections

Materials

- 1. Zeiss AxioScan.Z1 slide scanning microscope system (Carl Zeiss Inc., Thornwood, NY, USA) with a Plan-Apochromat 20x/0.8 objective lens.
- 2. Hamamatsu Orca flash 4.0 camera
- 3. The Zeiss ZEN blue 2.3 software
- 4. MediaCybernetics' Image-Pro 11 software package (Rockville, MD, USA).

Troubleshooting



- Widefield fluorescent images are collected for GFP and DAPI fluorescent channels using a Zeiss AxioScan.Z1 slide scanning microscope system with a Plan-Apochromat 20x/0.8 objective lens.
- 2 All images are acquired using a Hamamatsu Orca flash 4.0 camera with an average tile count of 165 tiles per brain section.
- The Zeiss ZEN blue 2.3 software package was used for collection and stitching of the 2-color (DAPI & GFP) tiled images.
- Widefield fluorescent images are then post-processed using MediaCybernetics' Image-Pro 11 software package (Rockville, MD, USA).
- 5 Every stitched image is processed using a protocol modified for each antibody. Initially, the image is masked to solely include the tissue in areas of interest.
- Threshold segmentation is used for each antibody staining to separate actual signal from background auto-fluorescence.
- 7 Smart Segmentation is used to separate GFP-expressing puncta or fibrils from DAPIstained nuclei in tissue evaluated with each antibody.
- 8 The Count/Size function is designed to extract the percent of the tissue sample that stains positive for GFP.