

Overcoming challenges in neuroscience: Multiplex immunofluorescence and autofluorescence isolation in brain tissue using multispectral imaging

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BACKGROUND

Disorders that affect the central nervous system are highly complex and often the result of many players interacting with one another. The ability to visualize and quantify multiple antigens and complex phenotypes within the same tissue section could lead to better treatment development and more accurate predictive outcomes.

Brain tissue (FFPE) is known to have an inherent autofluorescence signal that has made traditional multiplexing a challenge. Using Akoya's OPAL-TSA staining technology and spectral unmixing with autofluorescence isolation, we have been able to overcome some of these challenges in order to enhance the signal-to-noise ratio and to visualize the tissue environment more completely.

METHODS

Tissue Preparation and Staining

Tissue was fixed in PLP and sectioned, blocked, and embedded in paraffin. 10 micron slides were cut using a microtome and dehvdrated. After de-paraffinizing, the slides were stained manually. using the following steps: 1) 15 minute antigen retrieval using a microwave protocol; 2) 30 minute 3% DKS block; 3) 1 hour primary antibody incubation; 4) 30 minute secondary antibody incubation; 5) 10 minute Opal dye incubation. All steps were performed at room temperature. After staining was finished, the slides were coverslipped and sealed, and then stored at 4°C.



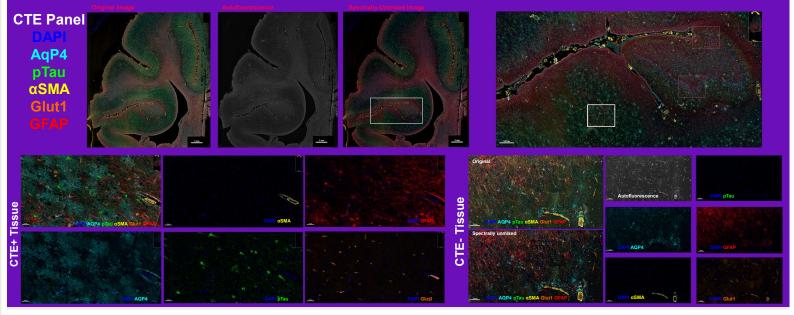
Vectra Polaris Scanning

The Vectra Polaris was set up using the MOTiF workflow with each channel autoexposed. At the under the same conditions as the multiplex image but rehydrated slide was scanned and used to separate the inherent autofluorescence of the tissue of Sudan Black or other harsh chemical treatments.

inForm Software Analysis

inForm software was used to create an image anal spectrally unmix Opal dyes and autofluorescence or annotated in the Phenochart software. Next, an was created to analyze pixel-based signal throug and thresholding. Finally, a project was created based signal using tissue segmentation, segmentation, and two rounds of phenoty phenotypes were identified and analyzed with phenoptrReports.

RESULTS – MULTISPECTRAL WHOLE-SLIDE IMAGING – HUMAN BRAIN TISSUE



RESULTS – INFORM IMAGE ANALYSIS

and the second		Pixel-Based Analysis: GF	AP, AQP4, pTau	Phe		Average Signal to Noise Ratio					
Star A Star					AP+, AdP4+		out AF removal	With AF removal			
					au+ AgP4+ au+, Gut1+	API	26:1	999:1	38.42		
i i	and the second second				· · ·	al 480	14:1	867:1	61.93		
		Contraction of the second s				al 520	41:1	651:1	15.88		
	and the second					al 570	229:1	999:1	4.36		
1						al 620	41:1	194:1	4.73		
F whole slide scan	CONTRACTOR STATES	and the second second			° Or	al 690	43:1	53:1	1.23		
the same time and ages, an unstained to determine and sue without the use ts.							CONCLUSIONS The addition of two markers over traditional FL- IHC allowed for more complete and complex				
s	Tissue Segmentation	Adaptive Cell Segmentation	Phenotyping, Round 1: AqP4, Glut1	Phenotyping, Round 2: GFAP, pTat	8		typing of cells		I COMPICX		
nalysis algorithm to e on tissue samples an analysis project pugh colocalization						for an the ne	increased sig	fluorescence s nal-to-noise rat al treatment (up Opal 480 chann	io without to 60x		
d to evaluate cell- n, adaptive cell otyping. Complex	and the plant						se of inForm s and cell-based	oftware allows I analysis.	for both		

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