

## Analysis Sequence "LysoTr-AV-updated-1550\*"

Input Image	Input		
	<b>Channel group</b> : 1 <b>Sequences</b> : ALL <b>Flatfield Correction</b> : Advanced Brightfield Correction <b>Stack Processing</b> : Individual Planes		
Find Nuclei	Input	Method	Output
	<b>Channel</b> : DAPI <b>ROI</b> : None	<b>Method</b> : B Common Threshold : 0.4 Area : > <u>20</u> $\mu\text{m}^2$ Splitting Coefficient : <u>3.1</u> Individual Threshold : 0.4 Contrast : > 0.1	Output Population : Nuclei
Calculate Intensity Properties (2)	Input	Method	Output
	<b>Channel</b> : DAPI <b>Population</b> : Nuclei <b>Region</b> : Nucleus	<b>Method</b> : Standard Mean Sum	Property Prefix : Intensity Nucleus DAPI
Select Population	Input	Method	Output
	<b>Population</b> : Nuclei	<b>Method</b> : Filter by Property Intensity Nucleus DAPI Mean : < <u>6000</u>	Output Population : Nuclei Selected
Find Image Region	Input	Method	Output
	<b>Channel</b> : 561nm <b>ROI</b> : None	<b>Method</b> : Absolute Threshold Lowest Intensity : $\geq$ <u>1550</u> Highest Intensity : $\leq$ inf Area : > 0 $\mu\text{m}^2$	Output Population : Image Region Output Region : Image Region
Calculate Intensity Properties	Input	Method	Output
	<b>Channel</b> : 561nm <b>Population</b> : Image Region <b>Region</b> : Image Region	<b>Method</b> : Standard Mean Sum	Property Prefix : LysoTr 561nm Intensity
Define Results	Results		
	<b>Method</b> : List of Outputs <b>Population</b> : Nuclei		

Number of Objects

**Population : Image Region**

LysoTr 561nm Intensity Sum : Sum

**Method : Formula Output**

Formula :  $a/b$

Population Type : Objects

Variable a : Image Region - LysoTr 561nm Intensity Sum Sum

Variable b : Nuclei - Nuclei Selected Sum

Output Name : LysoTr Intensity Sum / Neuron

**Object Results**

Population : Nuclei : ALL

Population : Nuclei Selected : None

Population : Image Region : ALL