Morphology Driven High-Plex Spatial Analysis of Tissue Microenvironments

For SITC Spatial Biology Award

September 2019



GeoMx™ DSP Enables Spatial, High-Plex Protein & RNA Profiling

High-Plex Mixtures of Proprietary Reagents

Protein reagentsOligo-labeled antibodies



RNA reagentsOligo-labeled probe

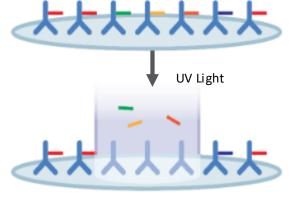
Target Complementary
Sequence Indexing Oligo
Sequence
Target RNA

Profile Regions of Interest on FFPE slide

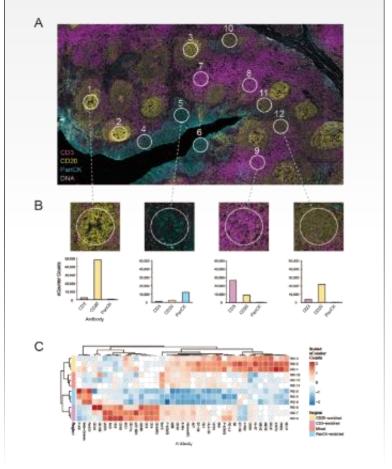


Illuminate Region of Interest, as Small as a Single Cell

Label FFPE Slide with Probe Mix



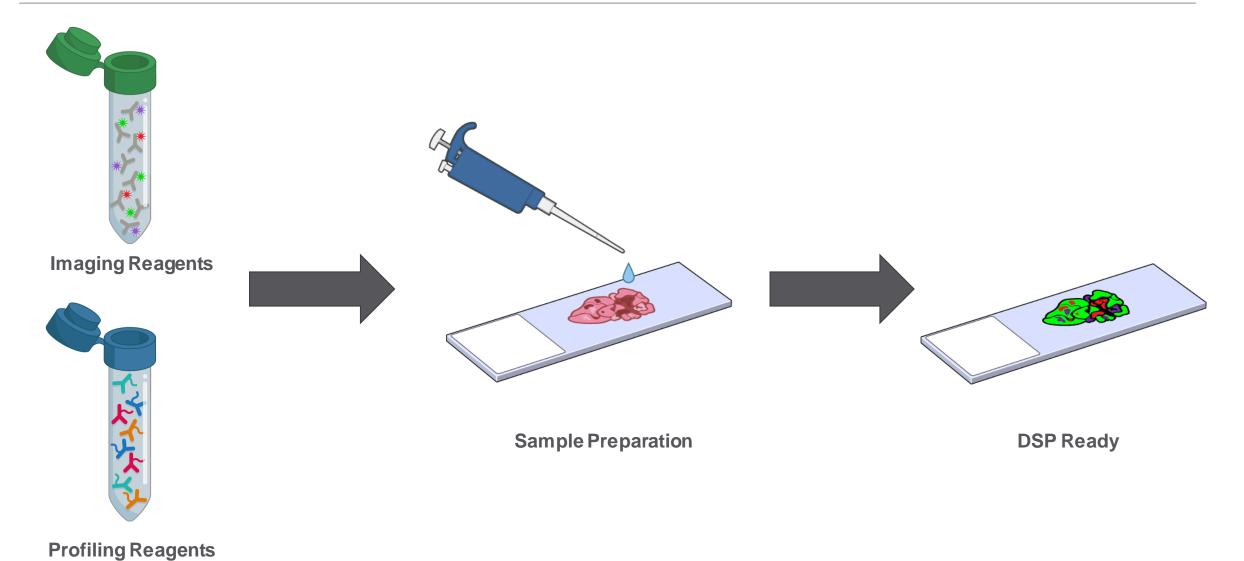
Rich Data Sets of Biology, Region by Region







Imaging and Profiling in One Assay

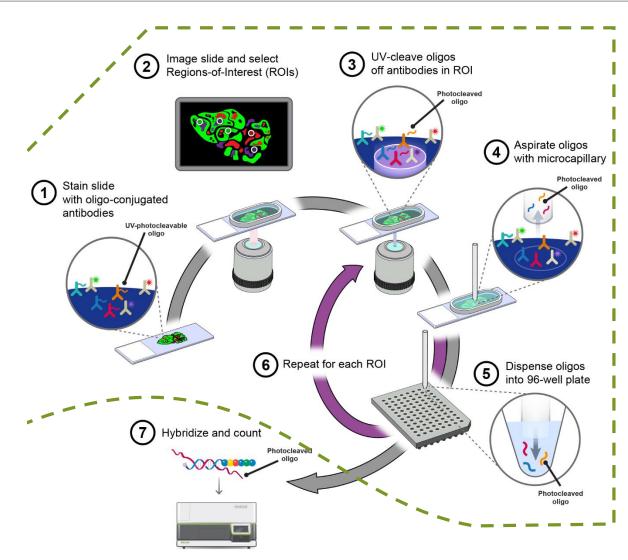




GeoMx DSP Enables Spatial, High-Plex Protein & RNA Profiling



GeoMx Digital Spatial Profiler Your GPS for Immuno-Oncology





DSP GeoMx has Numerous Advantages

Morphology driven profiling



Seamless integration with current pathology workflow

Multiplex

Many analytes on one tissue slice in a single pass

Multi-Analyte

High plex spatial analysis of both protein and RNA

Quantitative

Simple counts delivered to you in an elegant integrated software

High resolution

Limit of detection down to single cell

Non-destructive

Sample is only touched by light

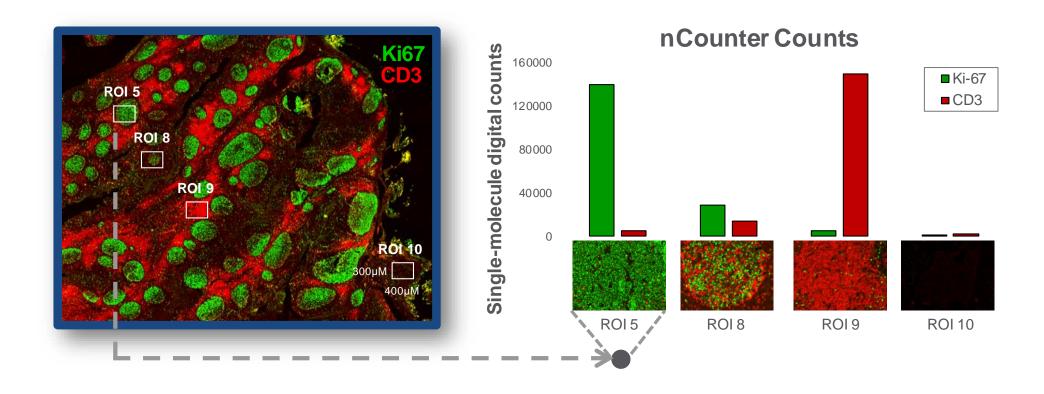
Throughput

Up to 12 sections per day



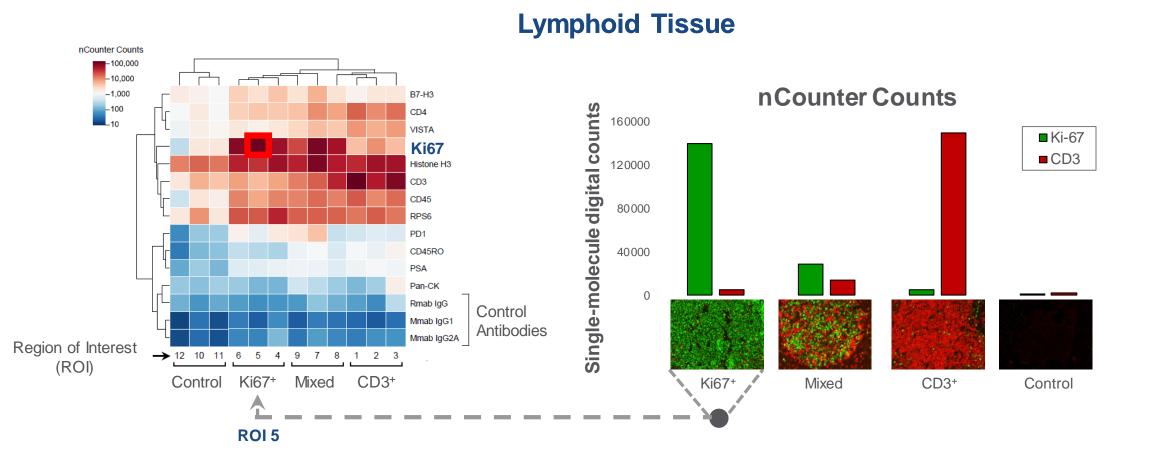
Single-molecule Counting of Photocleaved Oligos

Lymphoid Tissue



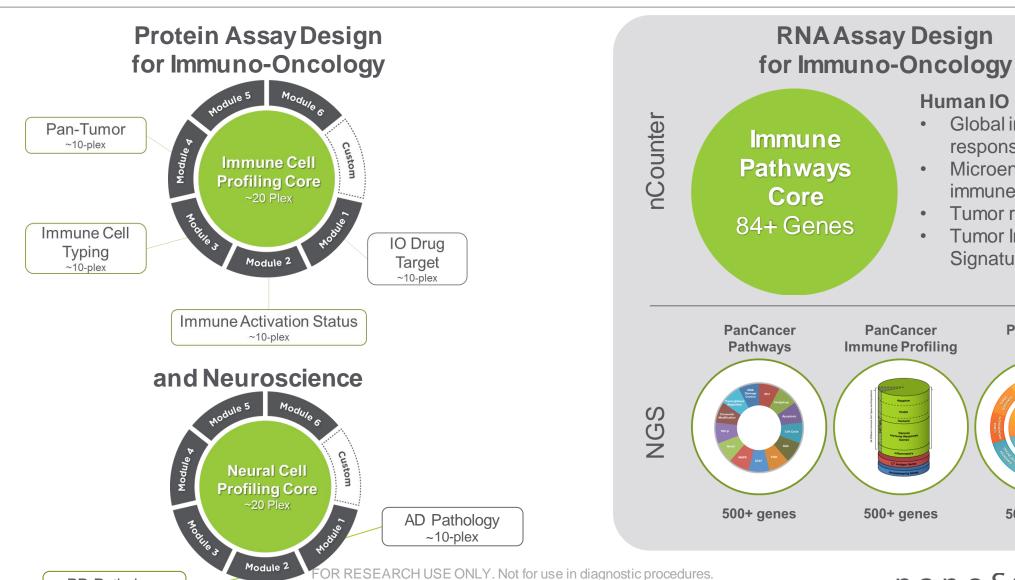


Single-molecule Counting of Photocleaved Oligos





Flexible and validated content design to fit a range of applications and plex needs



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Human IO

Global immune

Microenvironment

immune activity

Tumor reactivity

Tumor Inflammation

PanCancer

IO 360

500+ genes

response

Signature

PD Pathology

~10-plex

Currently Available IO Protein Content

CD66b

CD14

CD163

FAP-alpha

GeoMx™ Immune Cell Profiling Panel
Human Protein Core for nCounter®
PD-1
CD68
HLA-DR
Ki-67
Beta-2-microglobulin
CD11c
CD20
CD3
CD4
CD45
CD56
CD8
CTLA4
GZMB
PD-L1
PanCk
SMA
Fibronectin
Controls
Rb lgG
Ms lgG1
Ms lgG2a
Histone H3
S6
GAPDH

GeoMx™ IO Drug Target Panel Human Protein Module for nCounter®
4-1BB
LAG3
OX40L
Tim-3
VISTA
ARG1
B7-H3
IDO1
STING
GITR
GeoMx™ Immune Cell Typing Panel Human Protein Module for nCounter®
CD45RO
FOXP3
CD34

GeoMx™ Immune Activation Status Panel
Human Protein Module for nCounter®
CD127
CD25
CD80
ICOS
PD-L2
CD40
CD44
CD27
GeoMx™ Pan-Tumor Panel
Human Protein Module for nCounter®
MART1
NY-ESO-1
S100B
Bcl-2
EpCAM

	mmune Cell Profiling Pane
	tein Core for nCounter®
PD-1	
MHC II	
CD11b	
Ki-67	
F4/80	
CD11c	
CD19	
CD3e	
CD4	
CD45	
CD8a	
CTLA4	
GZMB	
PD-L1	
PanCk	
SMA	
Fibronectin	1
Controls	
Rb lgG	
Rt lgG2a	
Rt lgG2b	
Histone H3	}
S6	
GAPDH	

GeoMx™ IO Drug Target Panel Mouse Protein Module for nCounter®
AG3
X40L
īm-3
/ISTA
37-H3
ATE



Her2

PTEN

ER-alpha

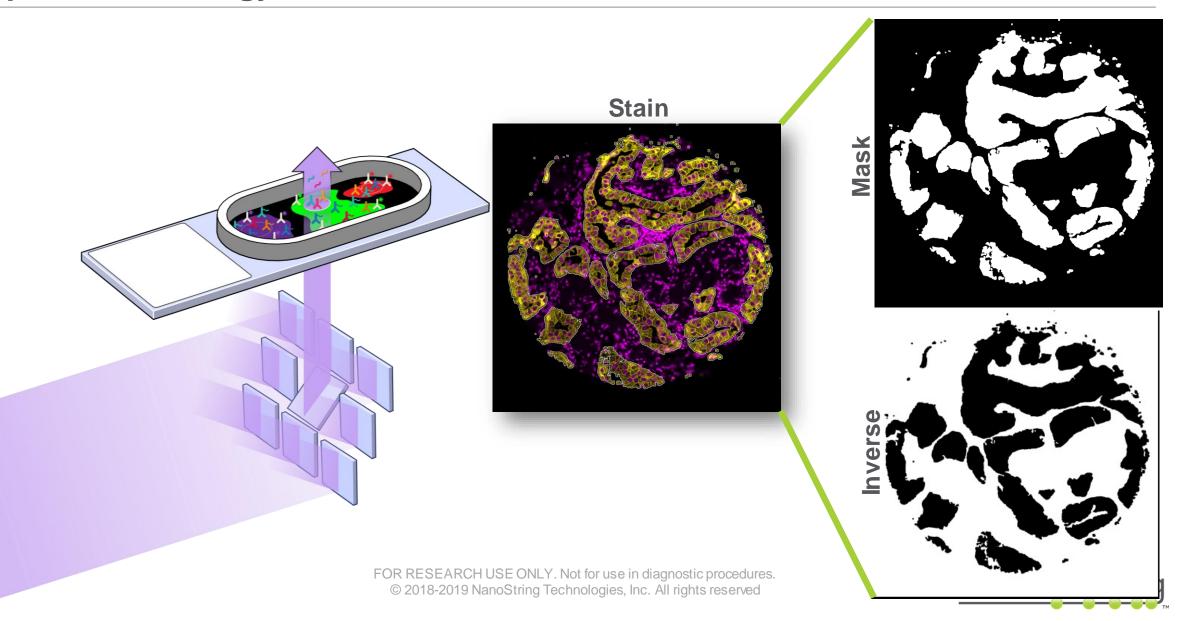
Currently Available IO RNA Content

GeoMx[™] Immune Pathways Panel Human RNA Core for nCounter®

AKT1	CD3E	CMKLR1	FOXP3	IFNAR1	ITGB8	PECAM1	TNFRSF9
ARG1	CD4	CSF1R	GZMB	IFNG	KRT	PSMB10	VEGFA
B2M	CD40	CTLA4	HAVCR2	IFNGR1	LAG3	PTEN	VSIR
BATF3	CD40LG	CTNNB1	HIF1A	IL12B	LY6E	PTPRC	Internal Reference
BCL2	CD44	CXCL10	HLA-DQ	IL15	MKI67	STAT1	Genes
CCL5	CD47	CXCL9	HLA-DRB	IL6	MS4A1	STAT2	OAZ1
CCND1	CD68	CXCR6	HLA-E	ITGAM	NKG7	STAT3	POLR2A
CD27	CD74	DKK2	ICAM1	ITGAV	pan-melanocyte	TBX21	RAB7A
CD274	CD86	EPCAM	ICOSLG	ITGAX	PDCD1	TIGIT	SDHA
CD276	CD8A	FAS	IDO1	ITGB2	PDCD1LG2	TNF	UBB



Optical Technology Enables Flexible ROI Selection

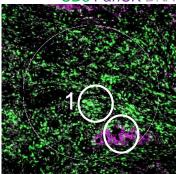


Five Unique Profiling Modalities Designed to Interrogate Tissue Samples

Geometric



CD3 PanCK DNA

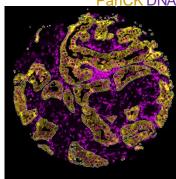


What is the heterogeneity of expression in different regions of my tissue?

Segmentation

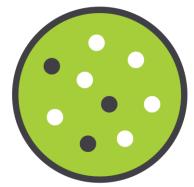


PanCK DNA



What is the expression profile of distinct biological compartments (e.g., Tumor-TME)?

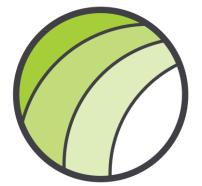
Rare cell

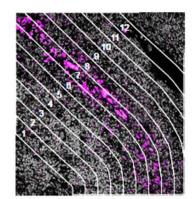


Ki67 GDS

What is the expression profile of a specific cell population in my tissue?

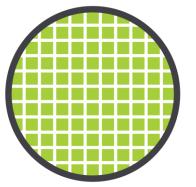
Contour

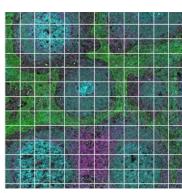




How does the immune environment change on either side of an infiltrate boundary?

Gridded

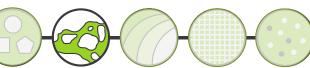




What novel targets are uncovered with deep mapping of a specific tissue region?



Integrating bulk and spatial profiling technologies for the discovery of RNA and protein biomarkers in muscle invasive bladder cancer



Background

- ~80% of patients with muscle-invasive bladder cancer (MIBC) fail to respond to immunotherapies.
- This suggest existence of complementary immune evasion mechanisms.
- There is a need for comprehensive immune profiling of patient tumor samples.

Experimental Questions/Goals

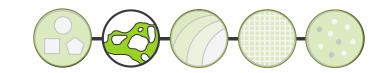
- Technical evaluation of bulk vs spatial genomics for assessing tumor microenvironment and defining MIBC molecular subtypes.
- Define mechanisms of immune evasion in PPARγ^{High} MIBC.
- Evaluate the role of TMB (tumor mutation burden) and tumor somatic mutations on immune evasion and PPARγ expression.
- Establish a solution for comprehensive immune profiling in clinical samples where tissue is limiting.



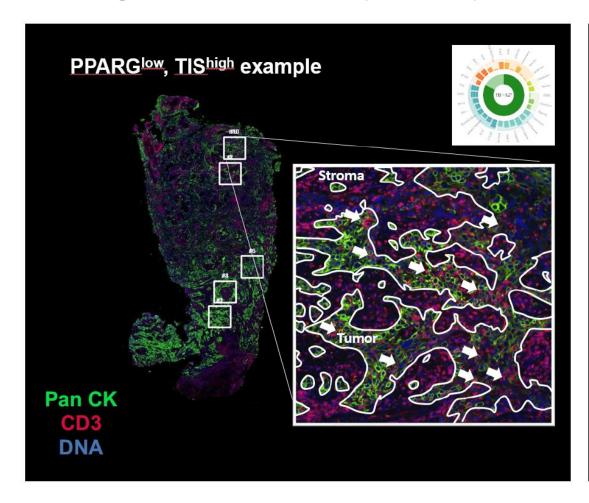
Victoria Rimkunas, PhD
Associate Director
H3 Biomedicine
(Now: Director at Repare Therapeutics)

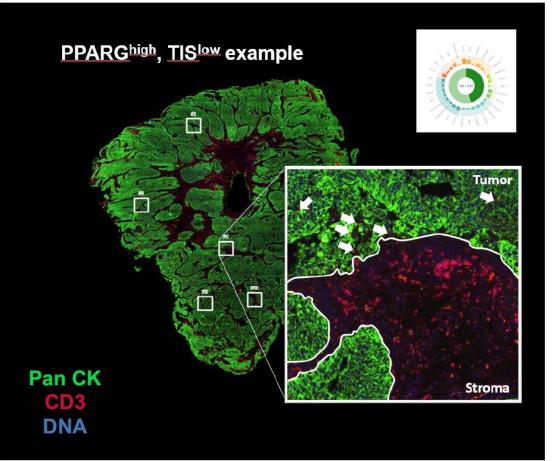


ROI Selection and Segmentation



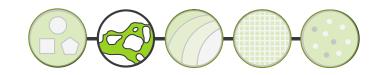
Auto-segmentation for tumor (Pan-CK+) and Stroma (Pan-CK-)



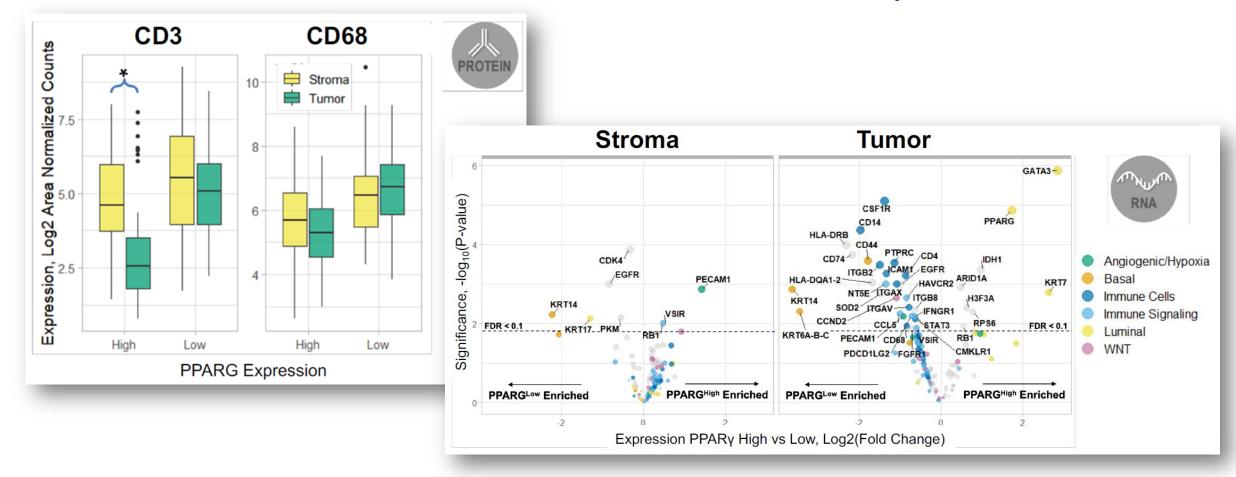




GeoMx DPS illustrates Immune Exclusion

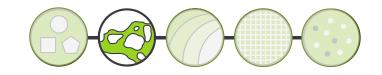


Immune Exclusion demonstrated on both Protein and mRNA level in PPARγHigh tumors

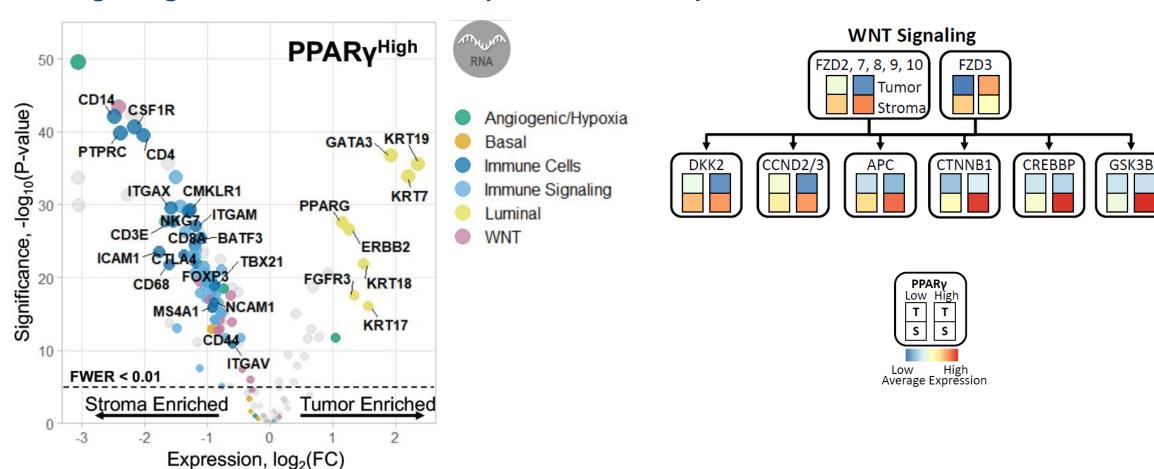




Pathway analysis



Wnt signaling is enriched in stromal compartment of PPARγ^{High} tumors





Tip for Success

General Tips

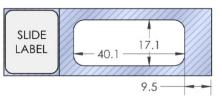
- Larger ROI give more robust counts
- 2 samples can be analyzed on 1 slide if they are both mounted within viewing area (white below)

Protein Profiling

- Robust data achieved with at least 50 cells per ROI
- Visualization Markers Used Successfully: CD45, CD3E, CD8, CD68, CD11c, S100B+HMB45 (pan-melanoma), panCK, pan-cadherin, and more (just ask)

RNA Profiling

- Robust profiling achieved with 200 cells per ROI
- Visualization Markers Used Successfully: CD8A, CD3E, CD68, PanCK AE1+AE3, DNA dye (e.g. Syto13)
- Others may also work but have not been tested



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Multi-Analyte

High plex spatial analysis of both protein and RNA

Quantitative

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High resolution

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