



GeoMx DIGITAL SPACE PROFILER: APPLICATIONS EXAMPLES

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Cancer Research Center of Lyon – Melanoma group

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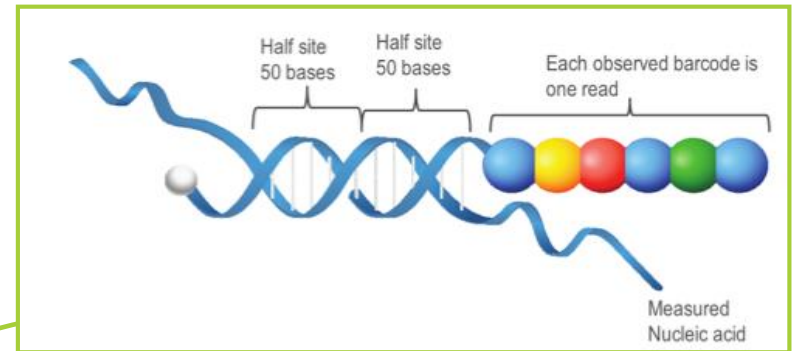
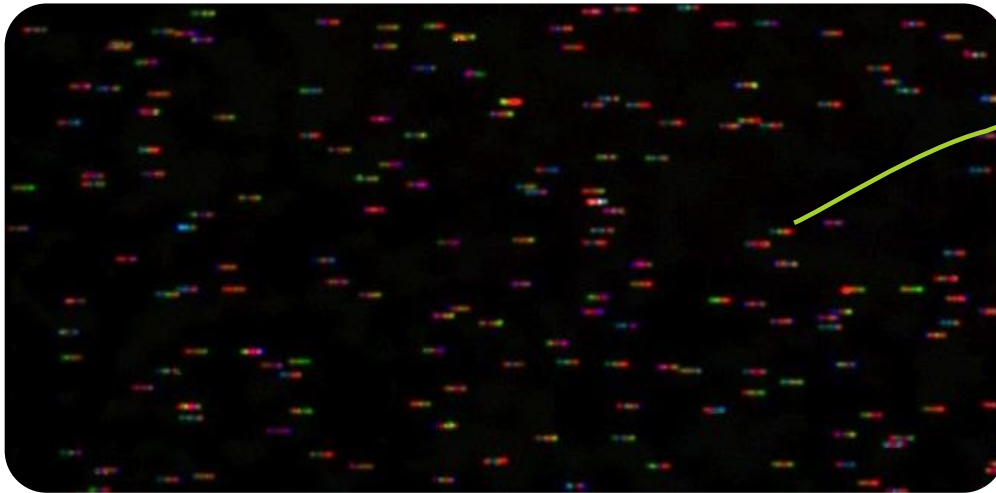
Digital Pathology & AI

London, 07/12/2018



① Digital counting of single molecules

- Probes up to 800 genes simultaneously
- **Digital gene expression** applied to biological pathways

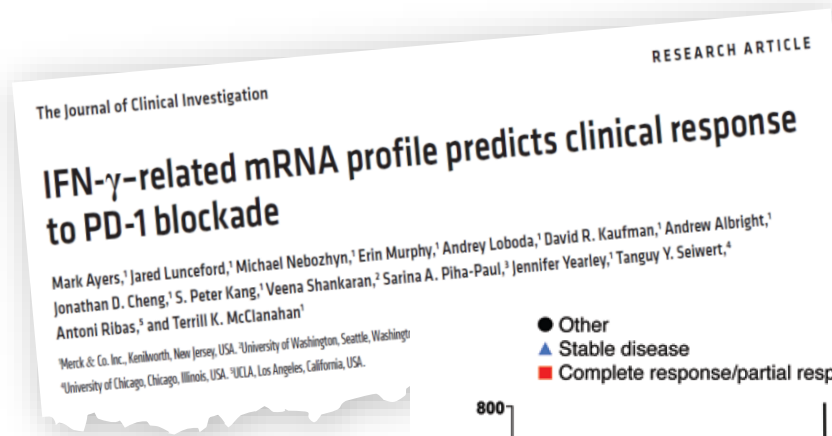


nanoString
TECHNOLOGIES

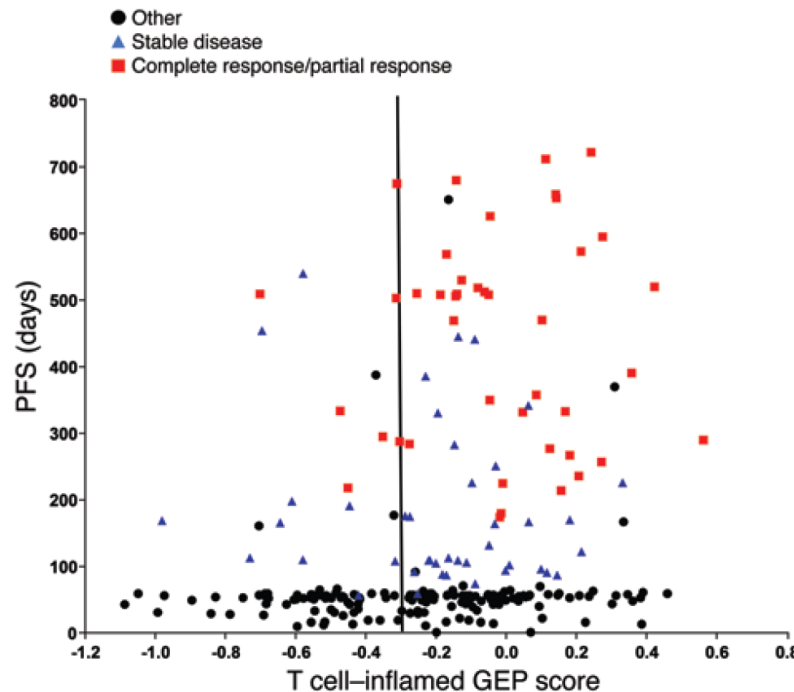
Single molecule fluorescent barcodes
each attached to an individual nucleic acid molecule

① Molecular imaging: for what?

BULK ANALYSIS



18 genes



Non-inflamed tumors
rarely respond

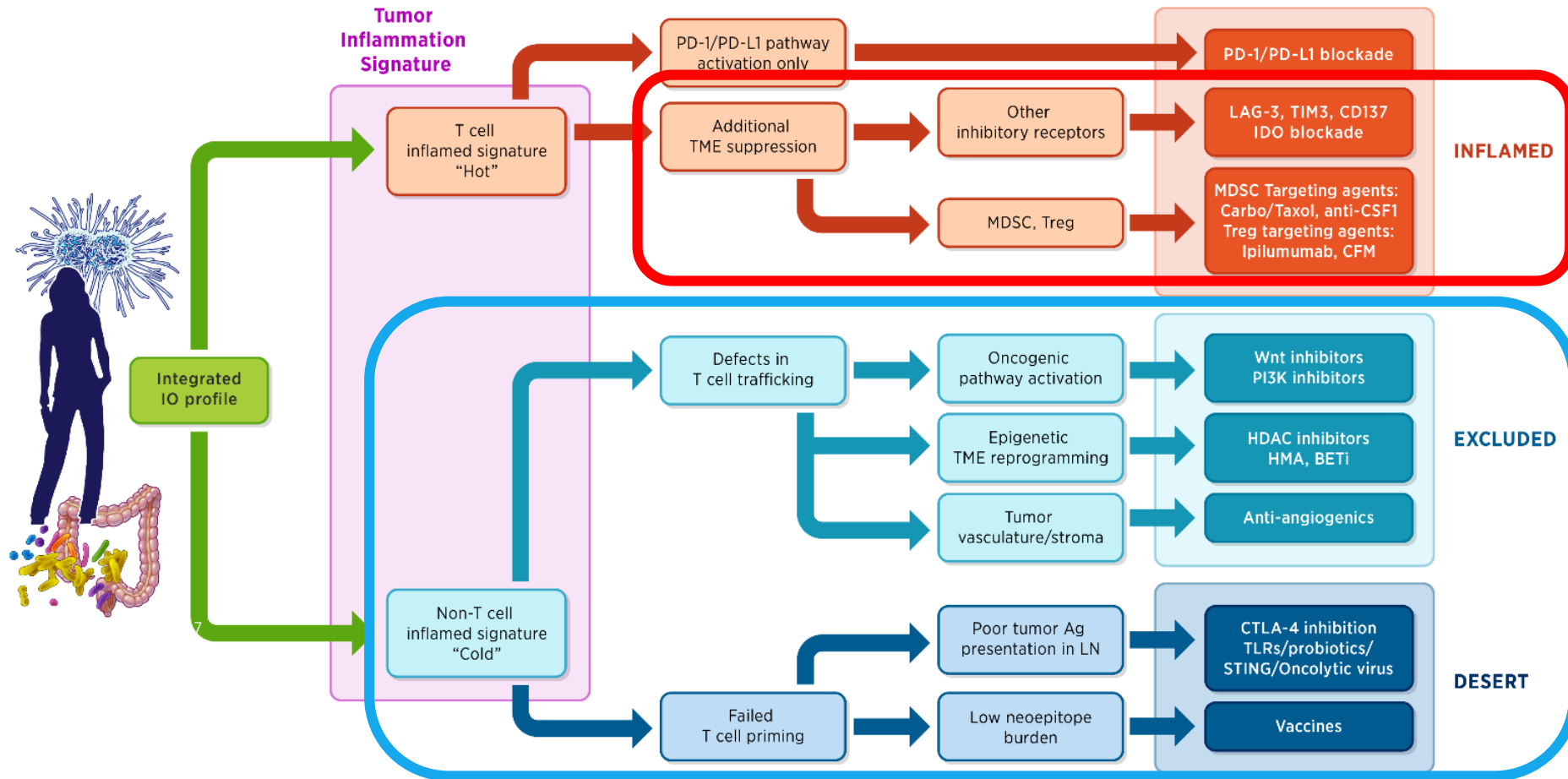
Almost all responses have
T-cell inflamed tumors

Not all inflamed tumors
respond

1 Molecular imaging: for what?

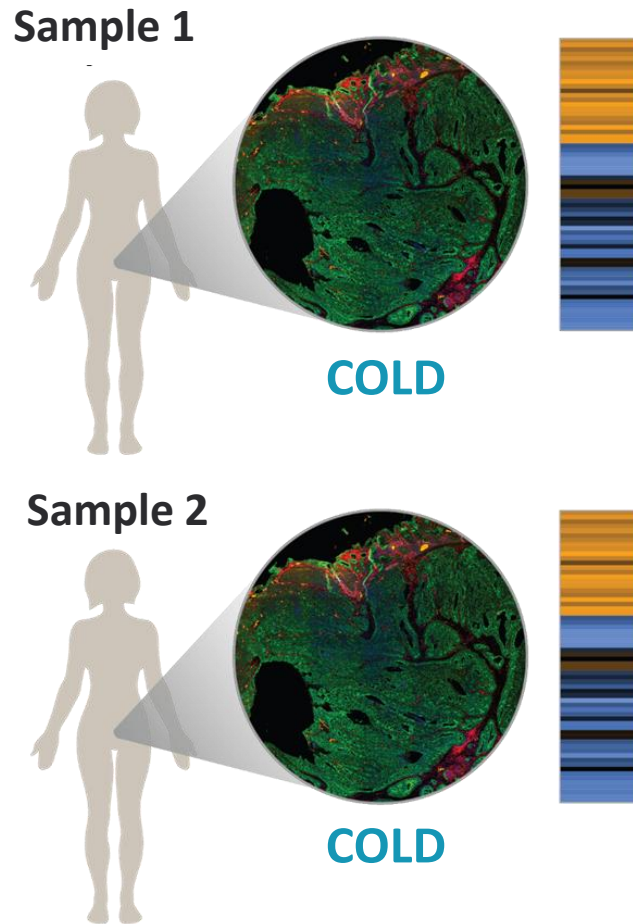
BULK ANALYSIS

770 genes



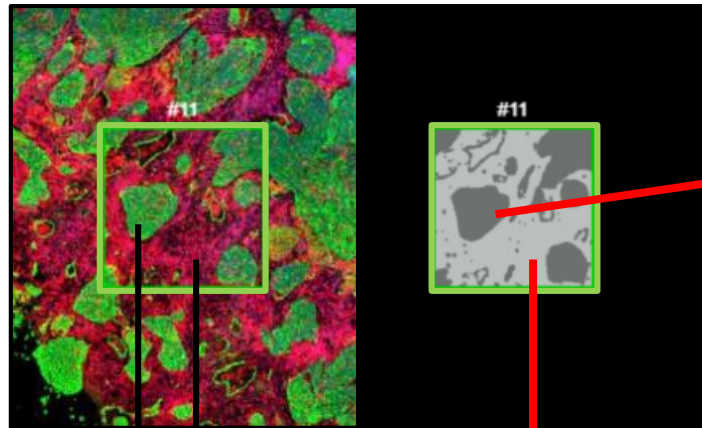
① Molecular imaging: for what?

SPATIALLY RESOLVED ANALYSIS



TRUE PERSONALIZATION of IMMUNOTHERAPY

① Molecular imaging: for what?



TME
Tumor

High levels of STAT3 expression in MDSCs have immunosuppressive effects and associated with poor prognosis

Treatment options may include removal of MDSCs using CSF-1R

High levels of STAT3 expression in tumor only indicates highly proliferative tumor associated with poor prognosis

Treatment options may include anti-STAT3 agents like JAK1/2 inhibitors to block STAT3



Jim Allison (MDAnderson)
Nobel Prize Medicine (2018)

Tumor vs Tumor microenvironment

Myeloid-derived suppressor cells (MDSCs)

① Molecular imaging: challenges

- Keeping the **tissue architecture to integrate spatial information** (and avoid microdissection)
- **Resolution** (down to single cell)
- High **multiplexing** (more is better)
- **Sensitivity & dynamic range** (for RNA+++)
- **Multiple analytes** (RNA + proteins)
- **Customization of panels / No panel** (discovery)
- **Cost-effectiveness**

2 GeoMx Digital Space Profiler

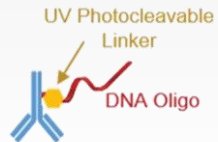


FONDATION
Hospices Civils de Lyon

High Plex Mixtures of Proprietary Reagents

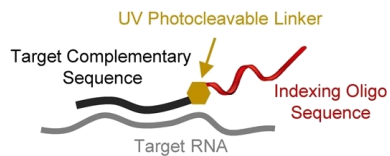
Protein reagents

Oligo-labeled antibodies



RNA reagents

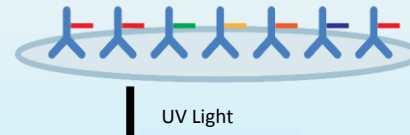
Oligo-labeled probe



Profile Regions of Interest on FFPE slide

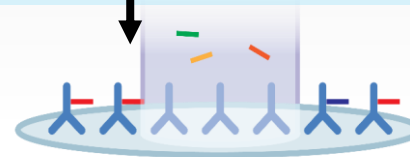
How it Works

Label FFPE Slide with Probe Mix



UV Light

Illuminate Region of Interest, as Small as a Single Cell

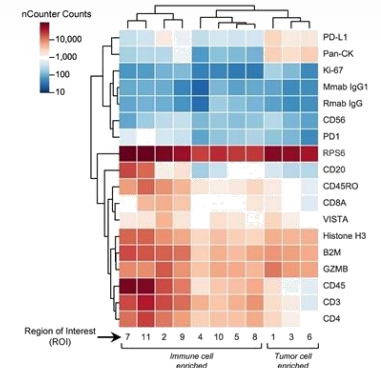
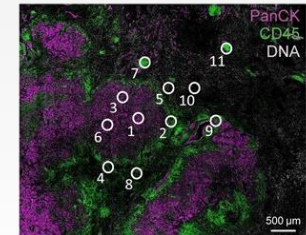


Analyze Barcodes on nCounter

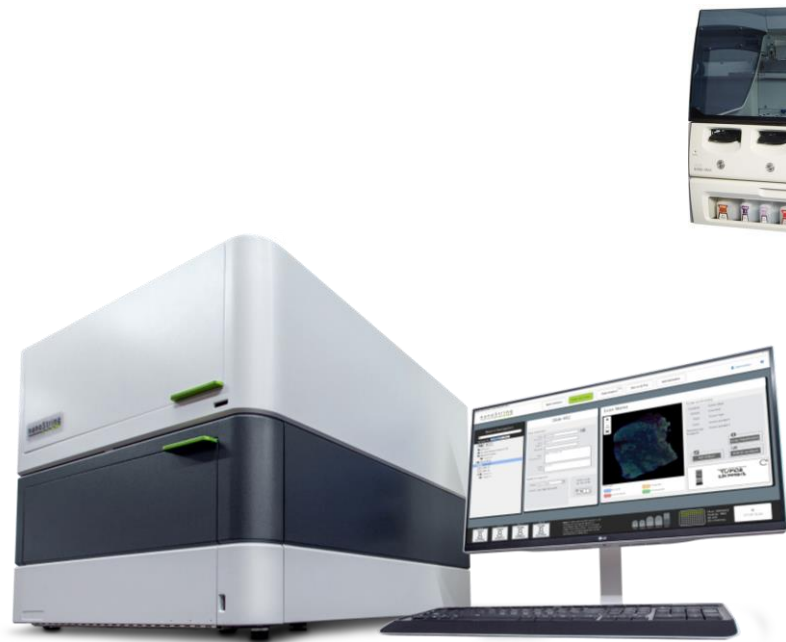


Rich Data Sets of Biology, Region by Region

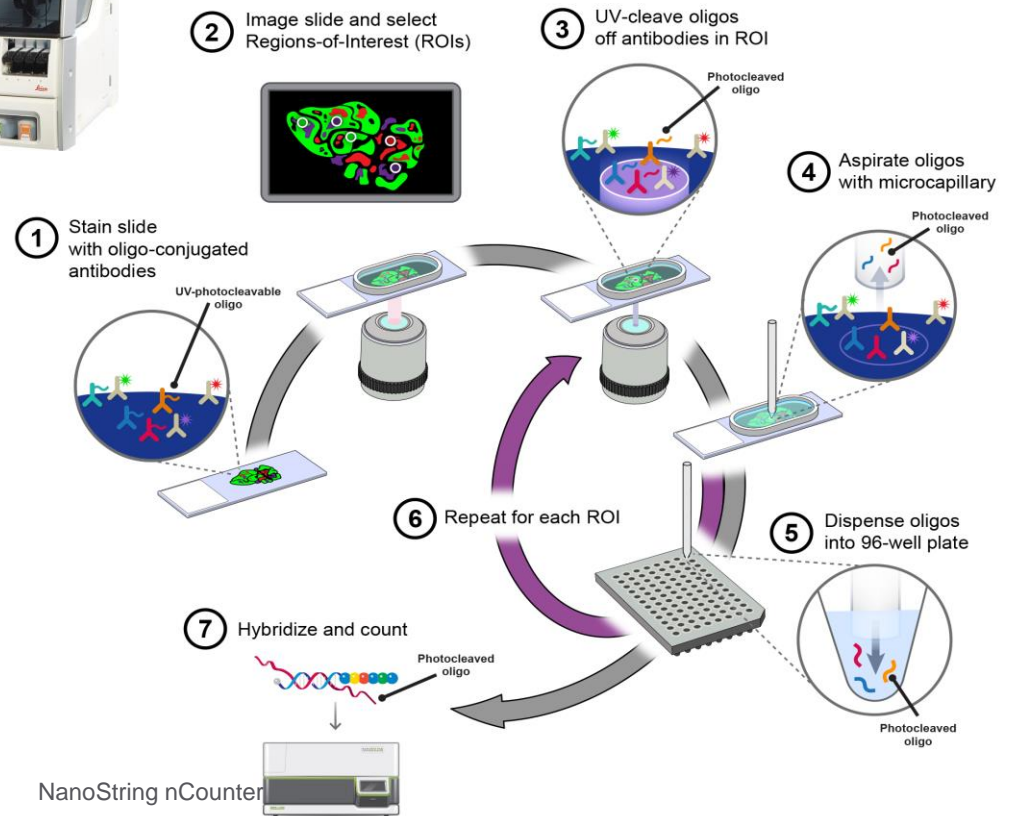
Lung Cancer (NSCLC)



2 GeoMx Digital Space Profiler



GeoMx™ Digital Spatial Profiler
Your GPS for Spatially-Resolved Biology



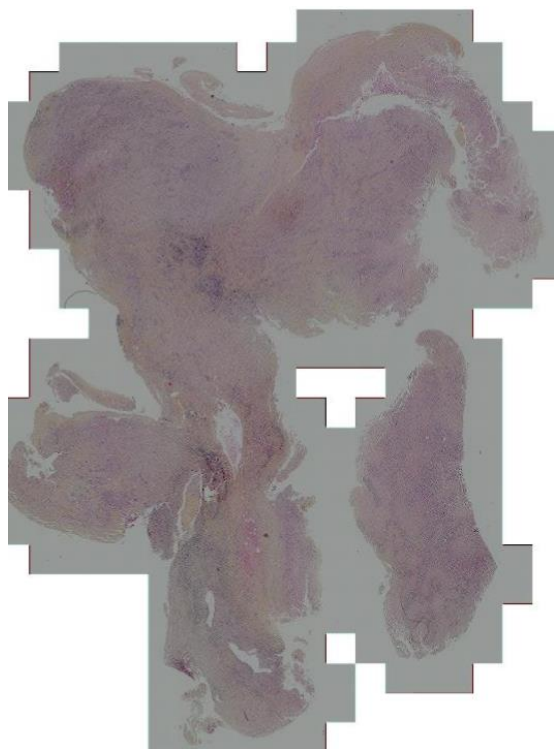
NanoString nCounter

Illumina NGS

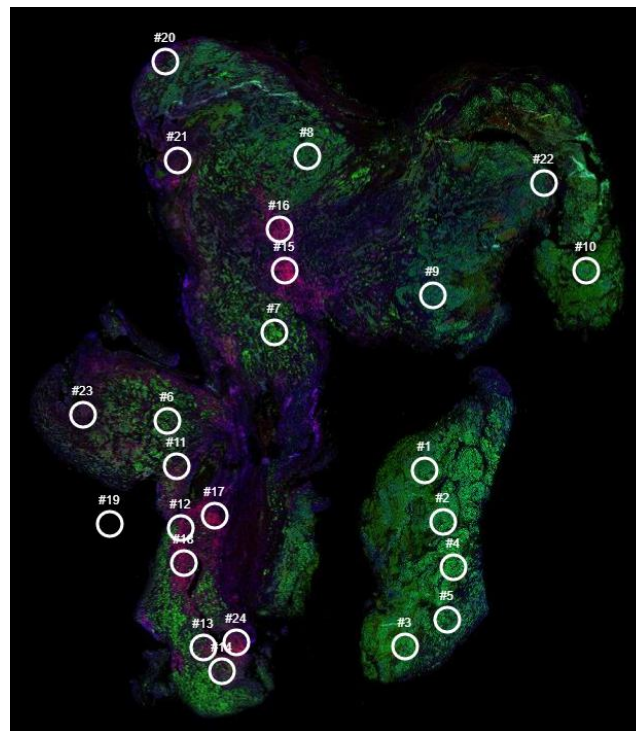


2 GeoMx DSP: Imaging

16SD07504



H & E



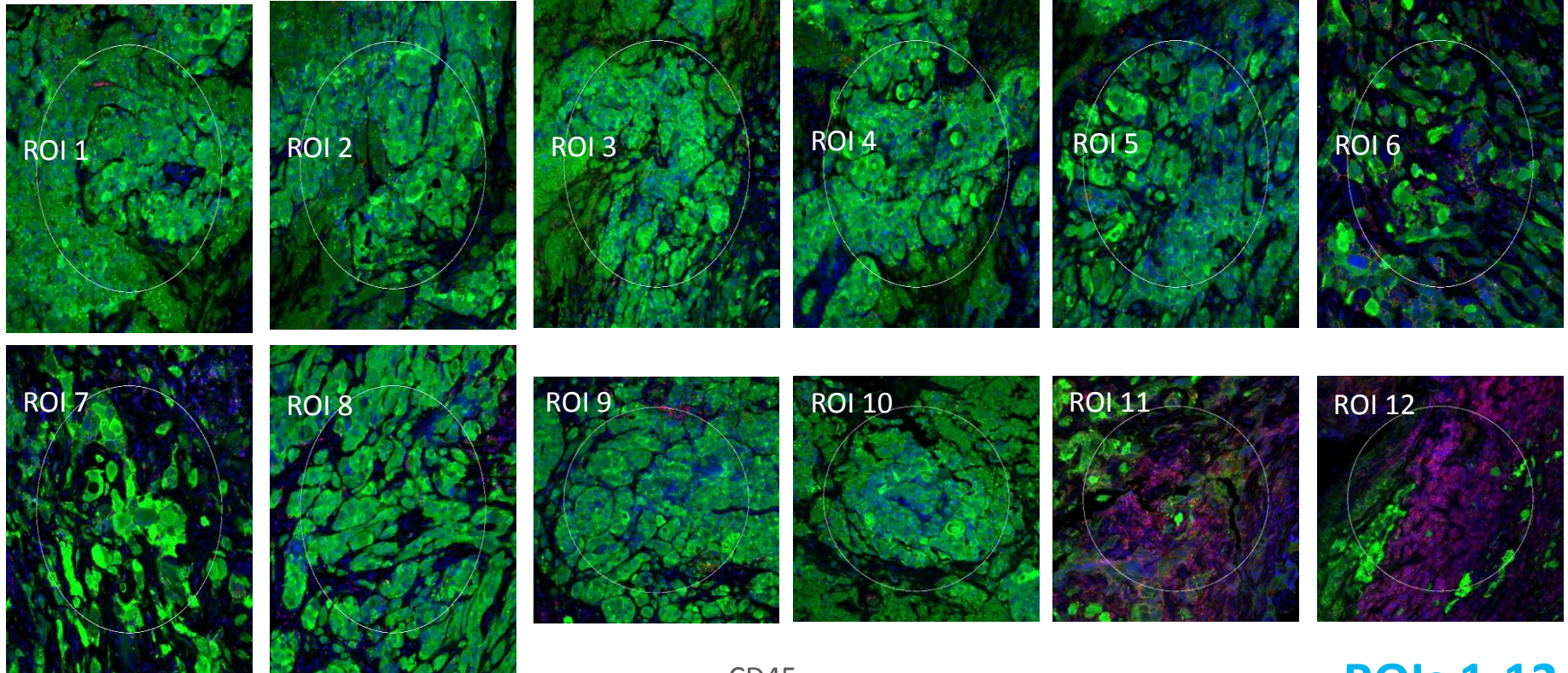
fluorescence

Visualization markers:

CD45 (red),
S-100B
(green),
DNA (blue)

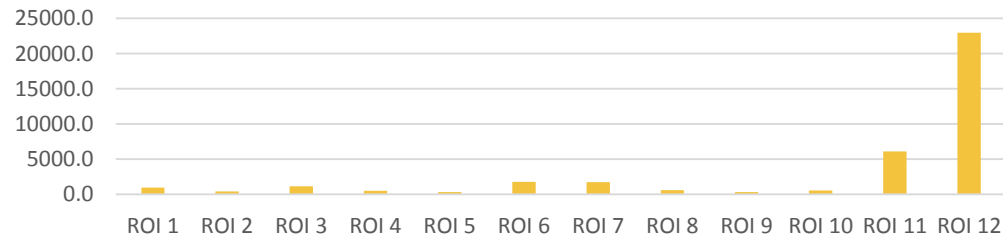
20X overlay

2 GeoMx DSP: Imaging

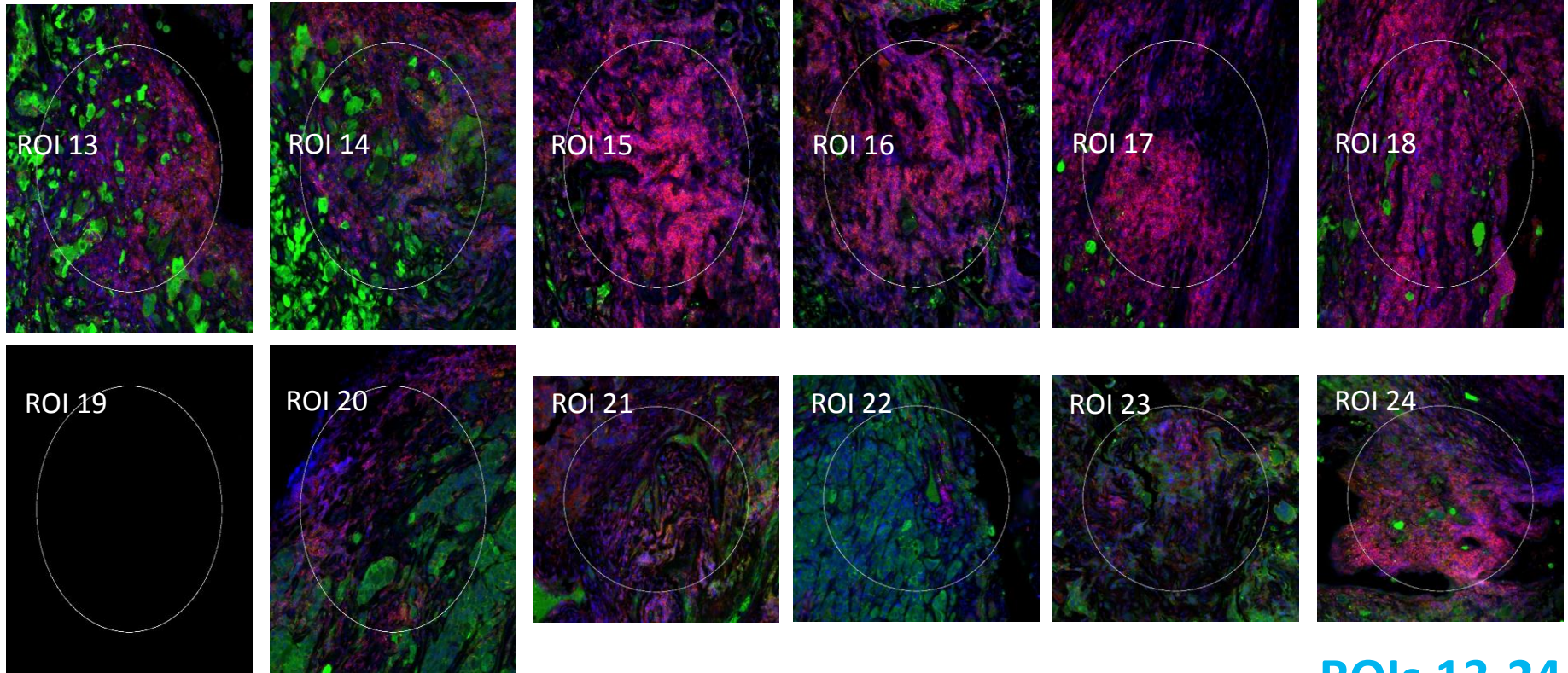


CD45

ROIs 1-12

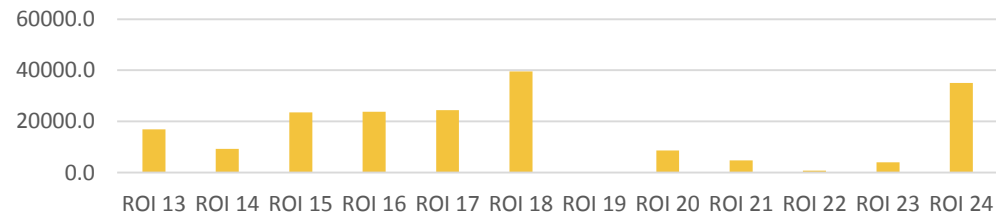


2 GeoMx DSP: Imaging

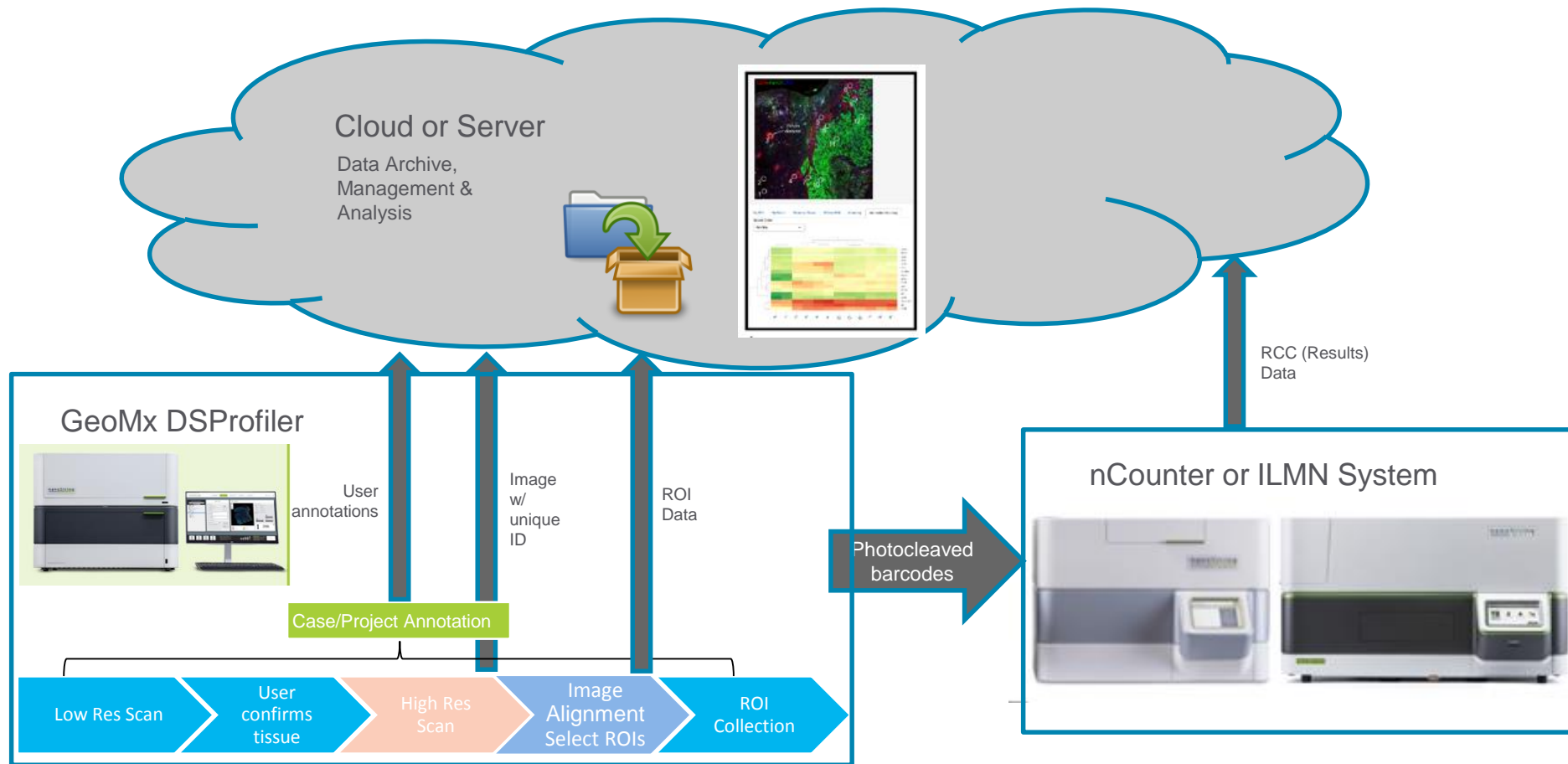


CD45

ROIs 13-24



2 GeoMx DSP: Software



2 GeoMx DSP: Software

nanoString DSPUser

Scan XYZ 123 B - Segments

- "Segments" are sub-regions of a particular biological target within your ROIs, for example immune cells or tumor cells.
- Define your own segments using the "Import Segment Mask" buttons on the ROIs panel, or use the tools below to automatically generate segments.
- To automatically generate segments, first click "Add Segment Definition".
- For each segment definition, choose whether each fluorescence channel should be included (+), excluded (-), or ignored (0) when generating the segment.
- Drag segment definitions up and down to change their definition order. Definition order effects which segments are generated first.
- Once your segment definitions are complete, click "Generate Segments".
- Once segments are generated, adjust their boundaries and coverage with the channel intensity thresholds grid.
- Segment changes will be updated and saved automatically.

Segment Definitions:

Segment	Red	Blue	Yellow	Color
Tumor	+	+	+	Red
Erode: 44 μm Dilate: 0 μm Hole Size: 60 μm^2 Particle Size: 50 μm^2				
Generation Order: 0				
Collection Order: 0				

Immune

Erode: 44 μm Dilate: 0 μm Hole Size: 60 μm^2 Particle Size: 50 μm^2

Generation Order: 1

Collection Order: 1

+ Add Segment Definition ☒ Show Advanced Parameters

Channel Thresholds:

ROI	R	G	B	Y
001	65	0	35	20
002	16	0	40	45
003	16	0	37	49
Modify All	-120	0	0	0

Revert Thresholds

30 μm

Slot Empty Slot Empty Slot Empty Slot Empty

localhost

Note: The NanoString DSP system is not certified for storage of personally identifiable information (PII) or patient health information (PHI). Ensure that all study, case, and slide identifiers are de-identified from individuals.

Water Imaging Buffer A Buffer B Waste

Plate: Analyte: RNA 95 wells 0 remaining

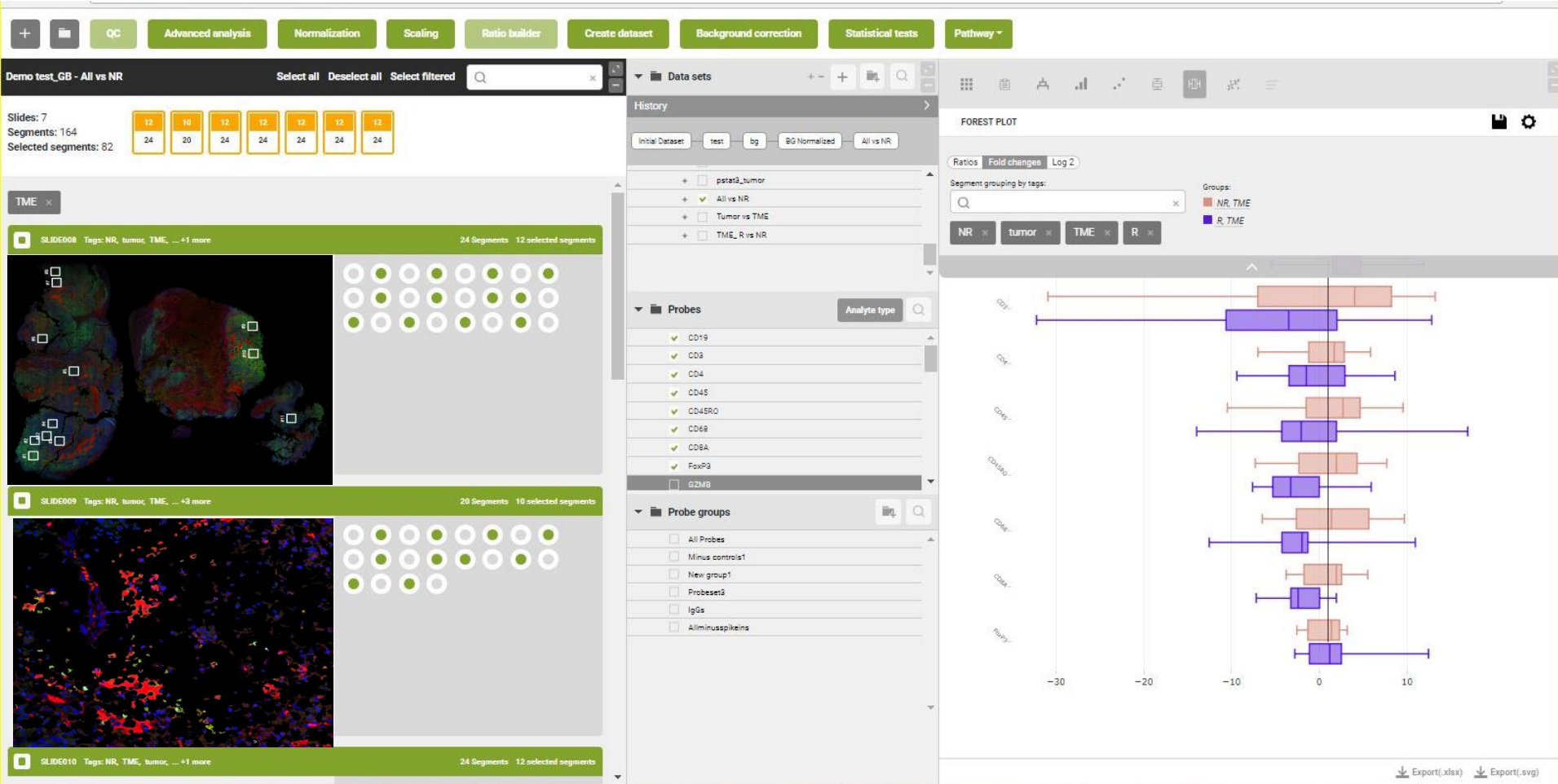
No Plate

STOP RUN

Automatic segmentation tumor / microenvironment



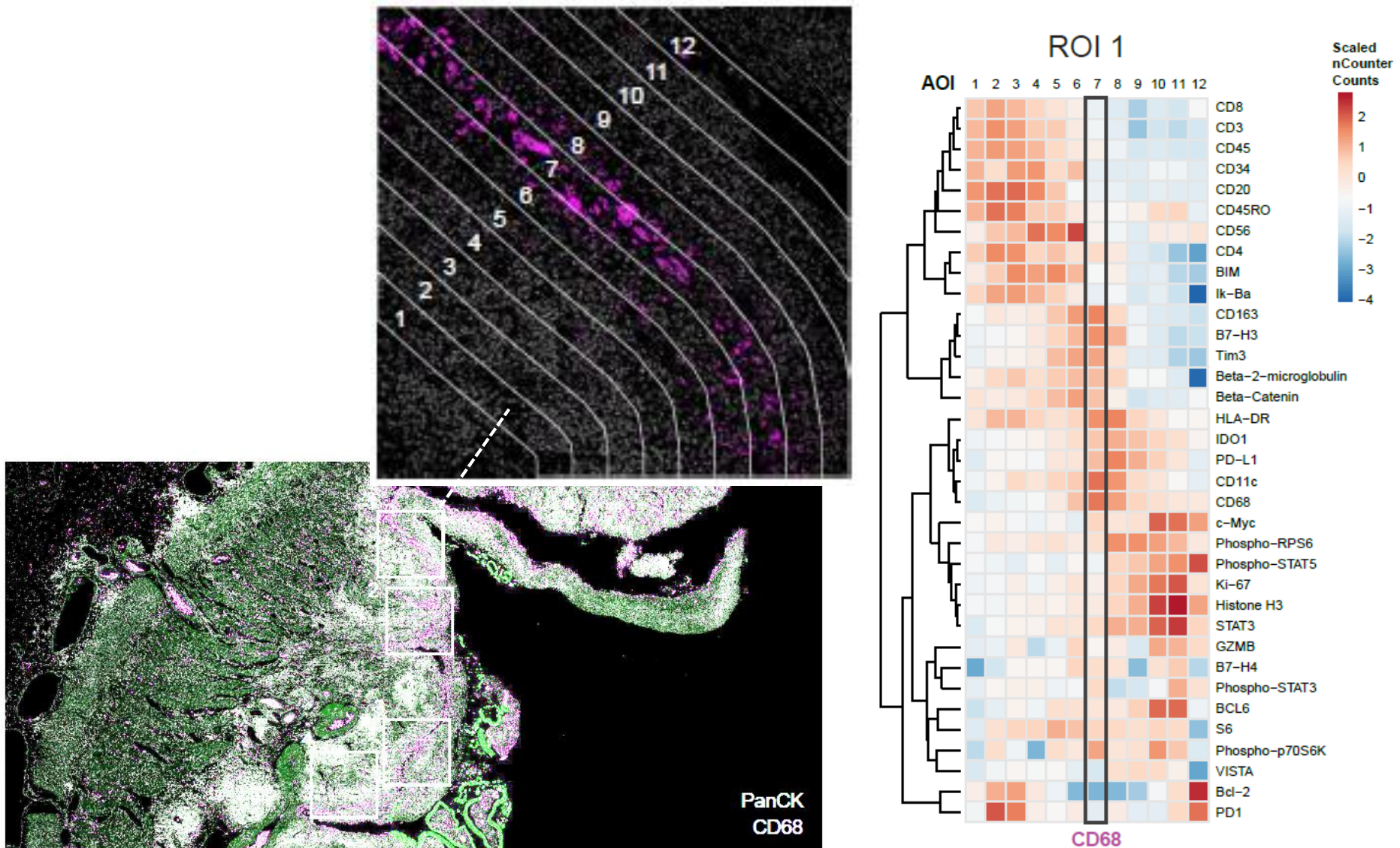
2 GeoMx DSP: Software



Comparison between ROIs

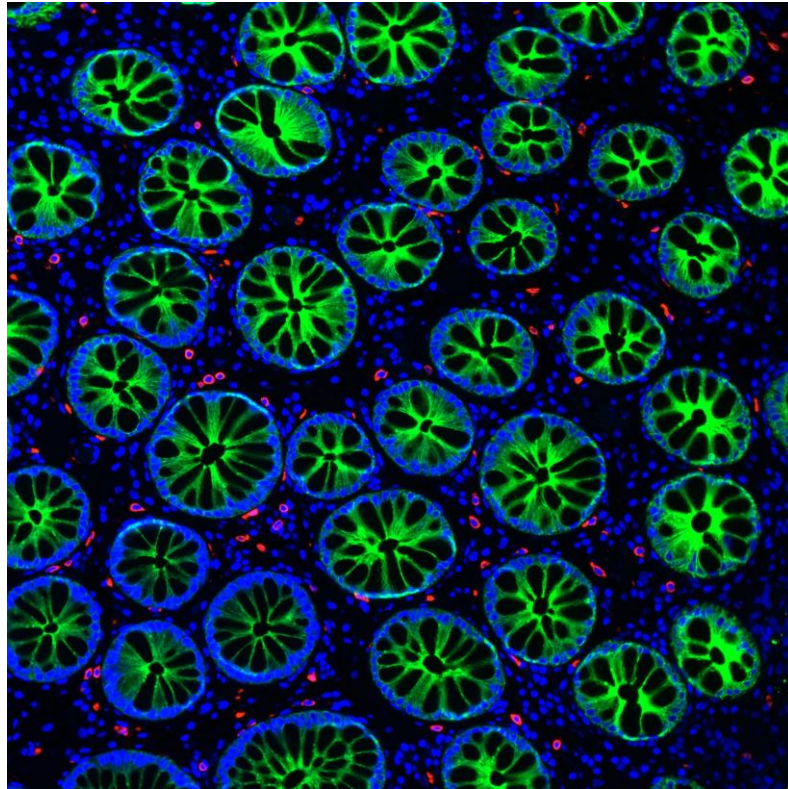


2 GeoMx DSP: tailored ROIs selection



2 GeoMx DSP: tailored ROIs selection

Profiling in the “shape” of Inflammatory Bowel Disease

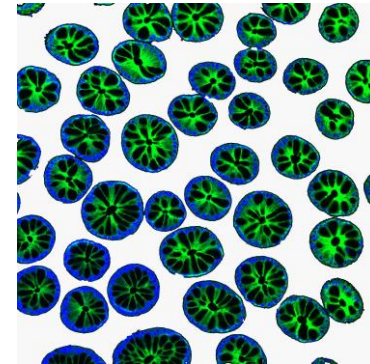
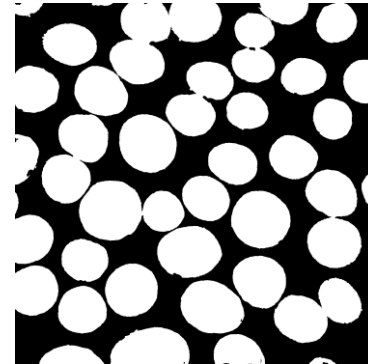


Blue = nuclear stain
Green = PanCytokeratin
Red = Tryptase, Mast-Cell Marker

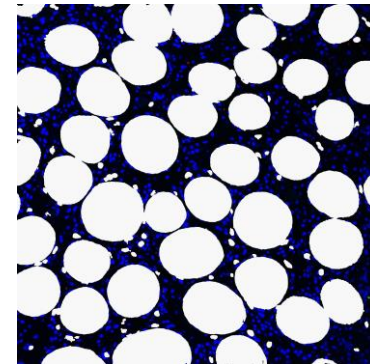
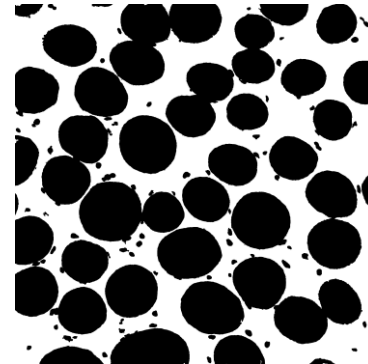
Mast-Cell
ROI mask



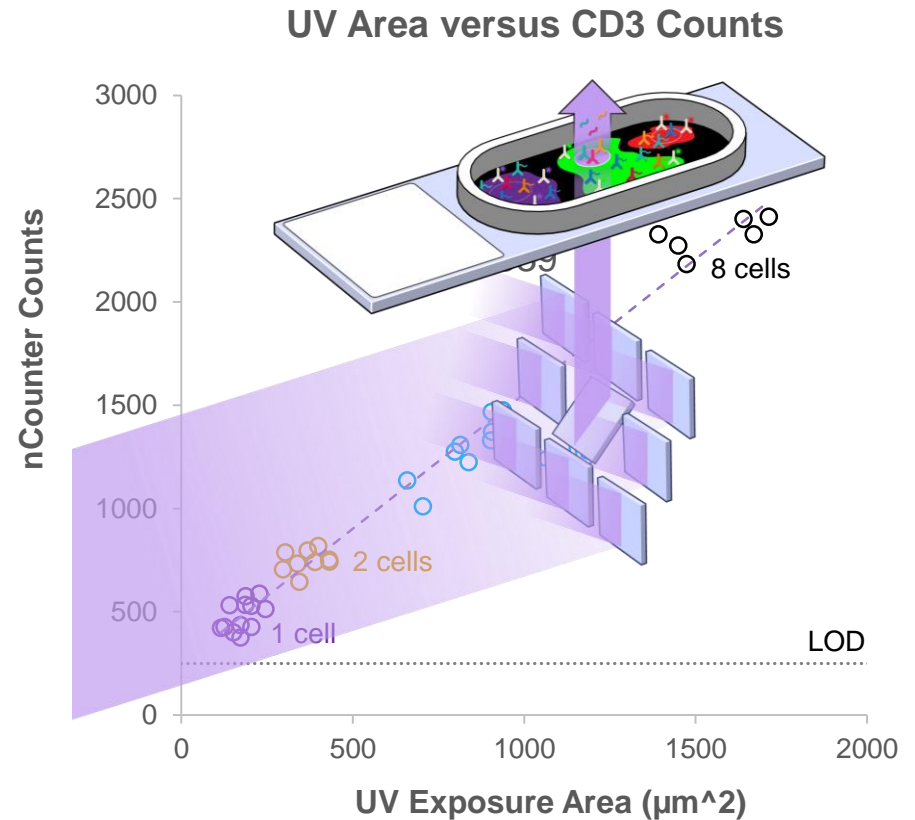
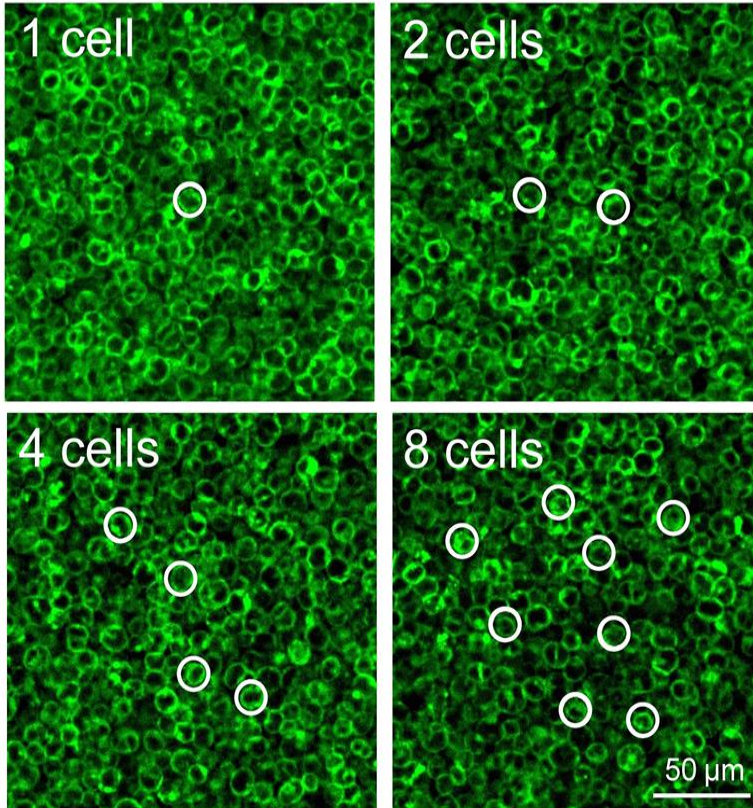
Intestinal Crypt
ROI mask



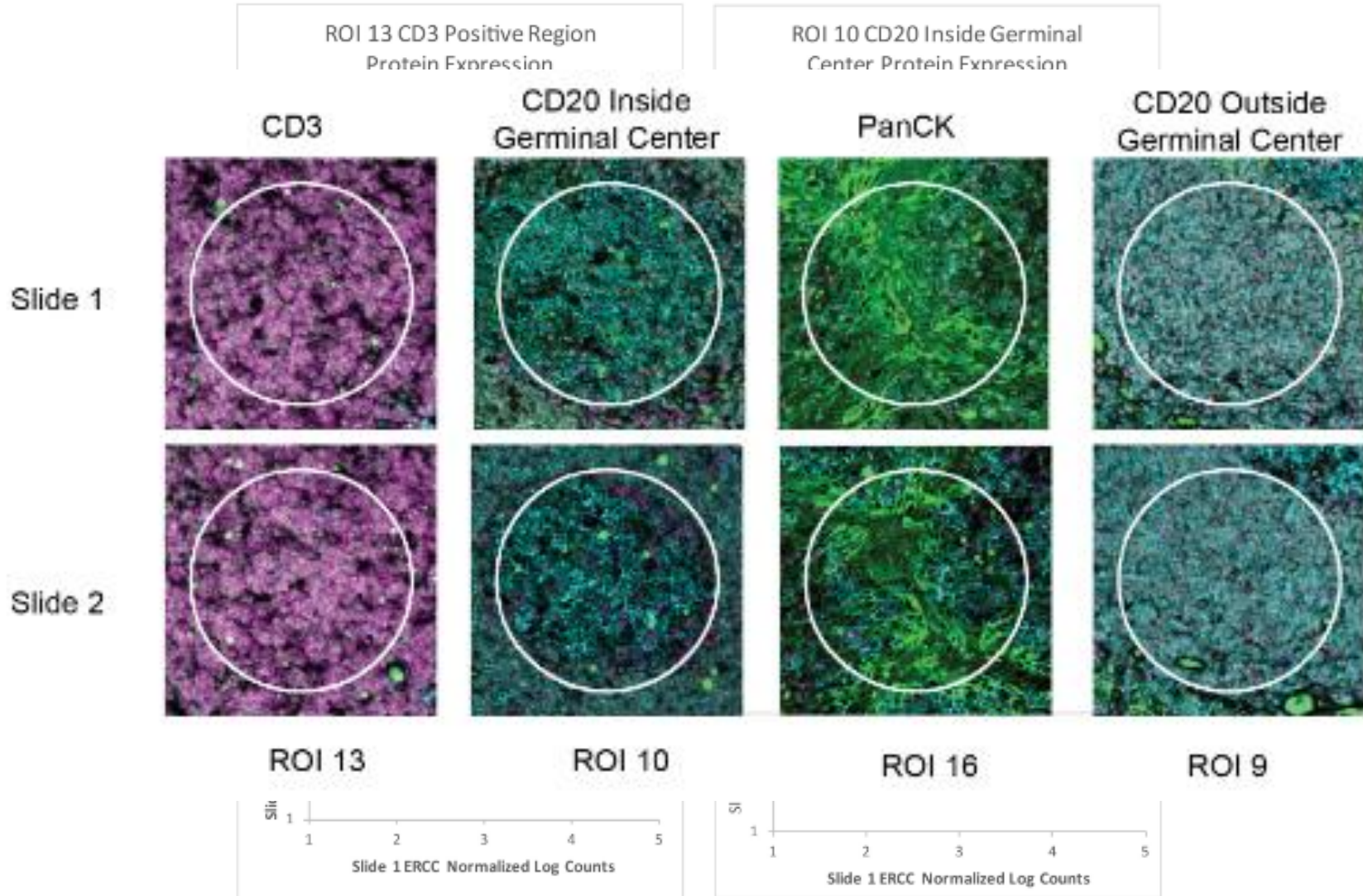
Connective tissue
ROI mask



2 GeoMx DSP: down to single cell

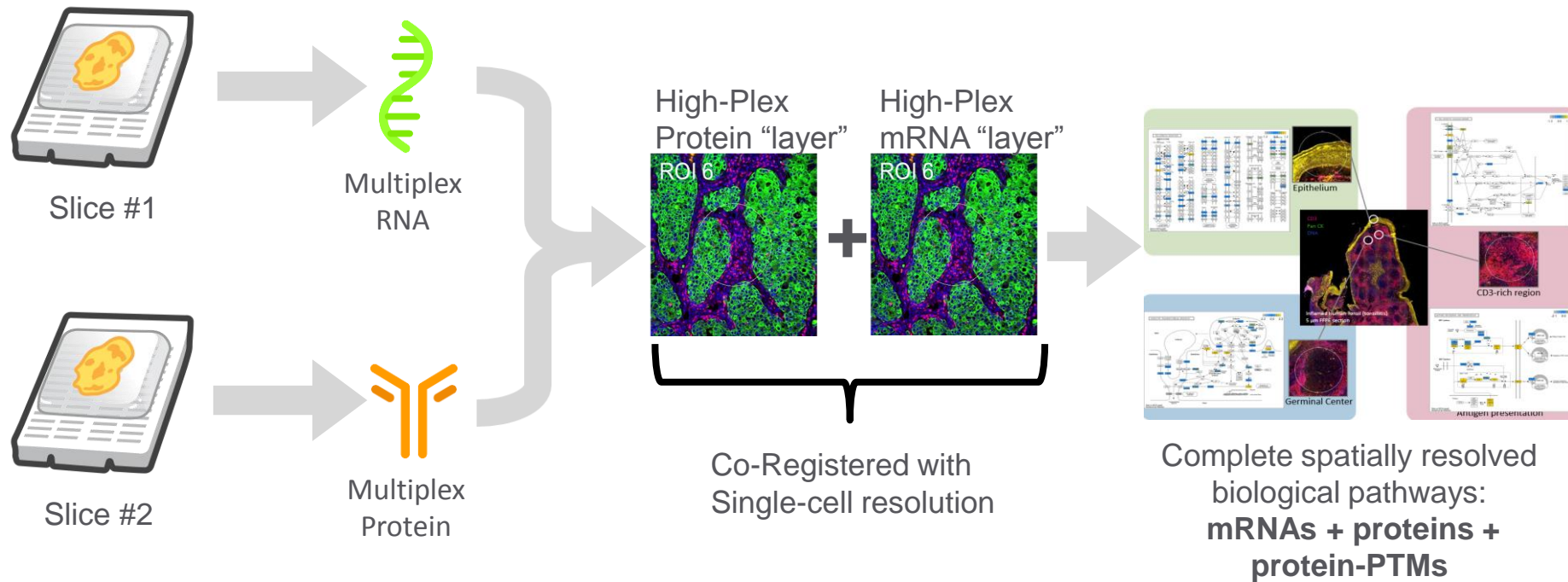


2 GeoMx DSP: Reproducibility



50-plex Immuno-Oncology protein panel on 5um FFPE tissue sections

2 GeoMx DSP: RNA & proteins



3 Examples in Immuno-Oncology

Core Cell Profiling

Beta-2-microglobulin
CD11c
CD20
CD3
CD4
CD45
CD56
CD68
CD8
CTLA4
GZMB
Histone H3
Ki-67
PD1
PD-L1
Pan-Cytokeratin
S6

Drug Target Module

4-1BB ARG1
B7-H4 B7-H3
LAG3 GITR
OX40L IDO1
Tim-3 STING
VISTA

Activation/Inhibition Module

CD127 PD-L2
CD25 CD40
CD80 CD40L
CD86 HLA-DR
ICOS CD27

Cell Typing Module

CD45RO Gamma
FOXP3 Delta TCR
CD34 CD14
CD66b FAPalpha

Tumor Module

MART1 EpCAM
NY-ESO-1 Her2/Erb
S100B B2
Bcl-2

Trafficking Module

CXCR3 CXCL9
CD31 E selectin
CXCL10 L-selectin
CXCL11 P-selectin

Cytokine Module

Interferon gamma
IL-17
IL-6
TGF beta
TNFalpha

■ Human IO Proteins panel



3 Examples in Immuno-Oncology

- 96-gene-plex (~ 1000 mRNA probes) for RNA detection

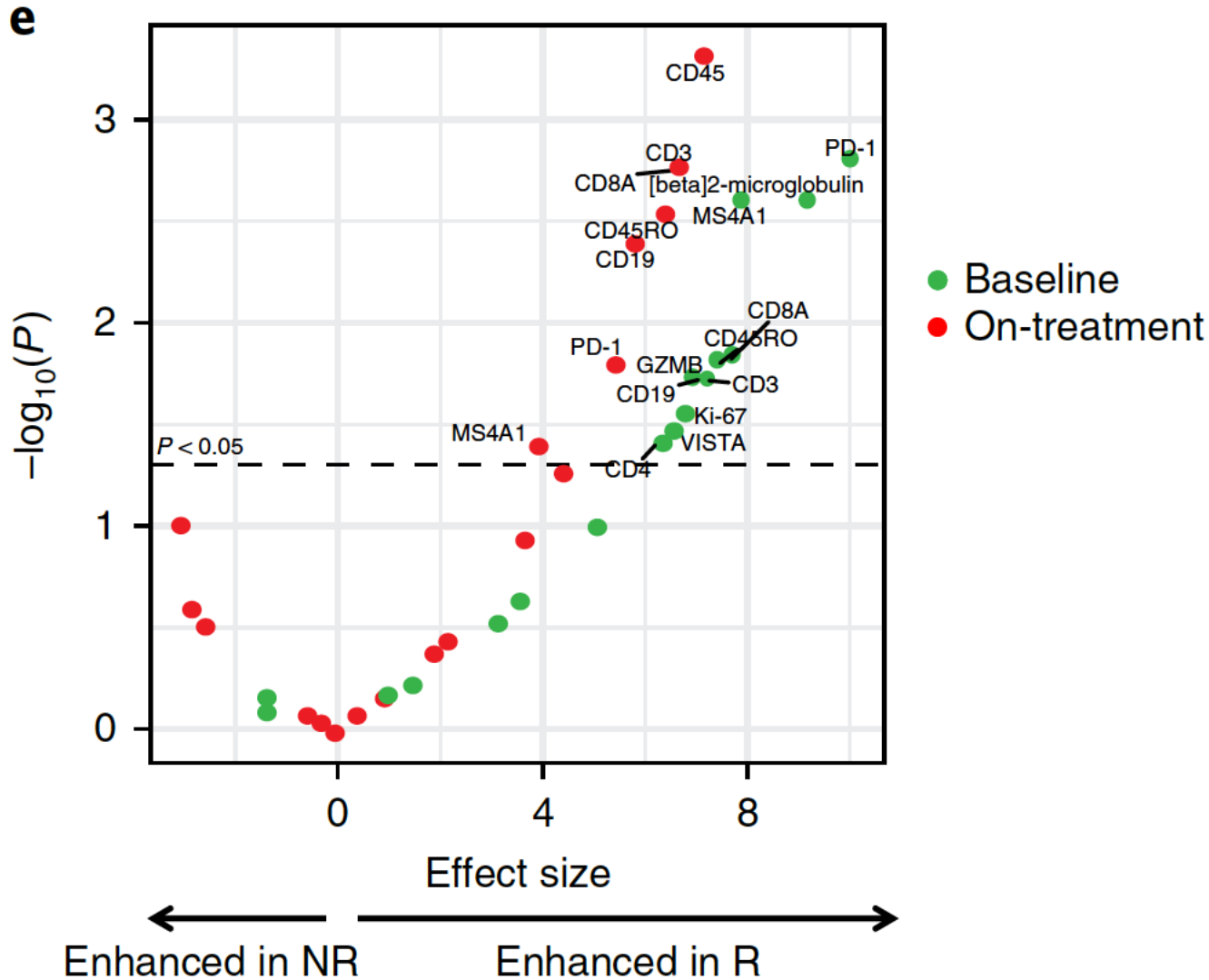
CCL5	LAG3	CD3E	CXCL10	IFNGR1	Multi-CK	VSIR (VISTA)	OAZ1
CD27	NKG7	CD4	DKK2	IL12B	pan-Melanoma	Custom 1	POLR2A
CD274 (PDL1)	PSMB10	CD40	EPCAM	IL15	PDCD1	Custom 2	RAB7A
CD276 (B7-H3)	PDCD1LG2	CD40LG	FAS	IL6	PECAM1	Custom 3	SDHA
CD8A	STAT1	CD44	FOXP3	ITGAM (CD11B)	PTEN	Custom 4	Neg 1
CMKLR1	TIGIT	CD47	GZMB	ITGAV	PTPRC (CD45)	Custom 5	Neg 2
CXCL9	AKT1	CD68	HAVCR2 (TIM3)	ITGAX (CD11C)	STAT2	Custom 6	Neg 3
CXCR6	ARG1	CD74	HIF1A	ITGB2	STAT3	Custom 7	Neg 4
HLA-DQA1/2	B2M	CD86	ICAM1	ITGB8	TBX21	Custom 8	Neg 5
HLA-DRB	BATF3	CSF1R	ICOSLG	LY6E	TNF	Custom 9	Neg 6
HLA-E	BCL2	CTLA4	IFNAR1	MKI67	TNFRSF9 (41-BB)	Custom 10	Neg 7
IDO1	CCND1	CTNNB1	IFNG	MS4A1 (CD20)	VEGFA	UBB	Neg 8

3 Examples in Immuno-Oncology

nature
medicine

Neoadjuvant nivoluma

Christian U. Blank¹,
Pia Kvistborg², Oscar
Johannes V. van Thielen³,
Lindsay G. Griepink⁴,
Harm van Tinteren⁵,
and Ton N. Schumacher⁶



TERS
018-0197-1

ia⁴,

Wells⁷,
d³,

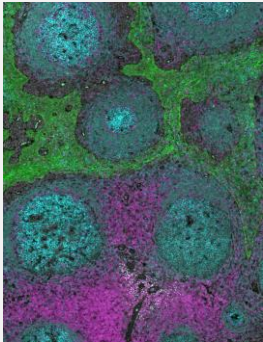
ian³,
rma^{5,10},

James Allison⁵, Michael T. Tetzlaff^{9,11,13} and Jennifer A. Wargo^{3,8,13*}



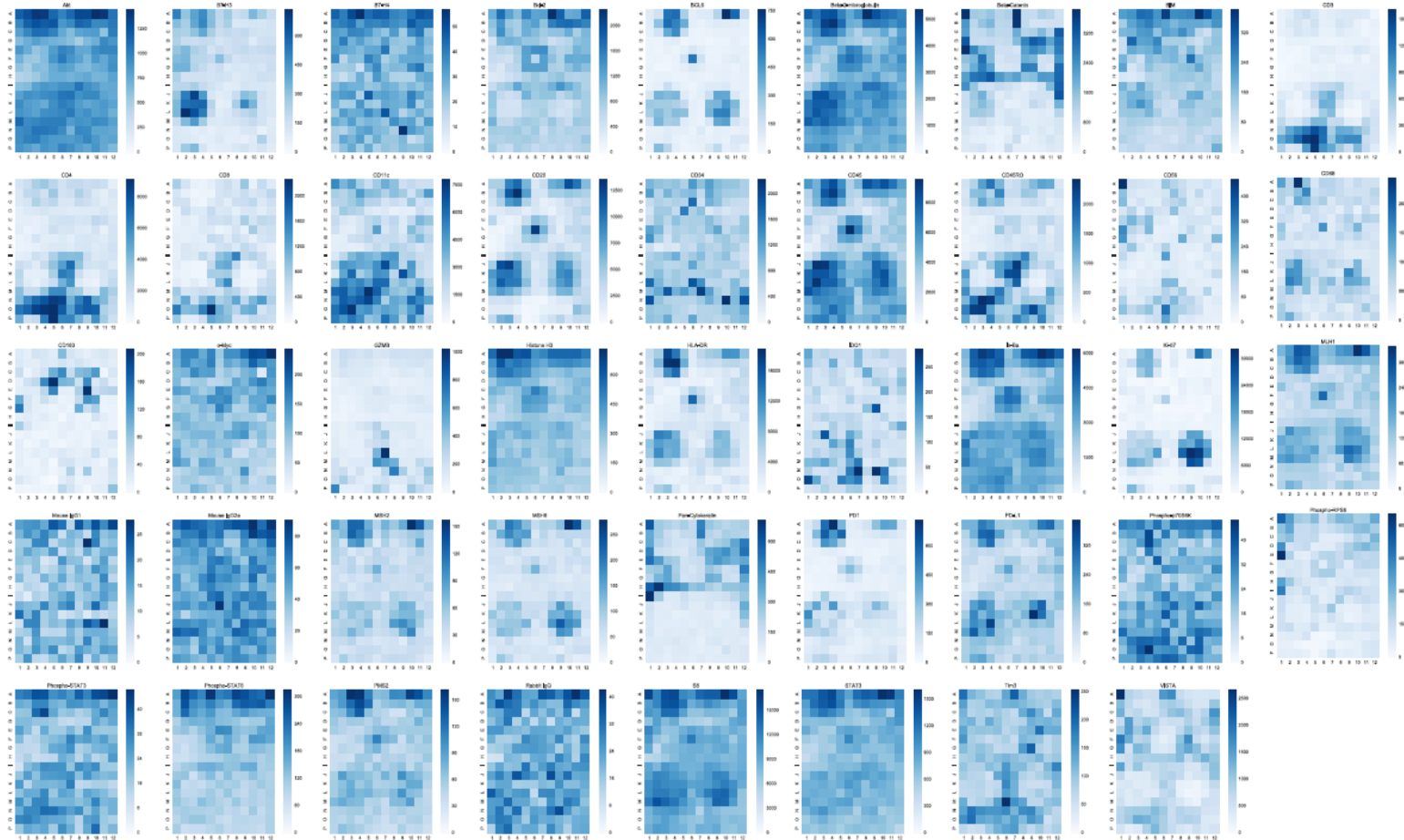
3 Examples in Immuno-Oncology

Rastered Region



5 um FFPE
Lymphoid Tissue
PanCK = green
CD20 = blue
CD3 = purple

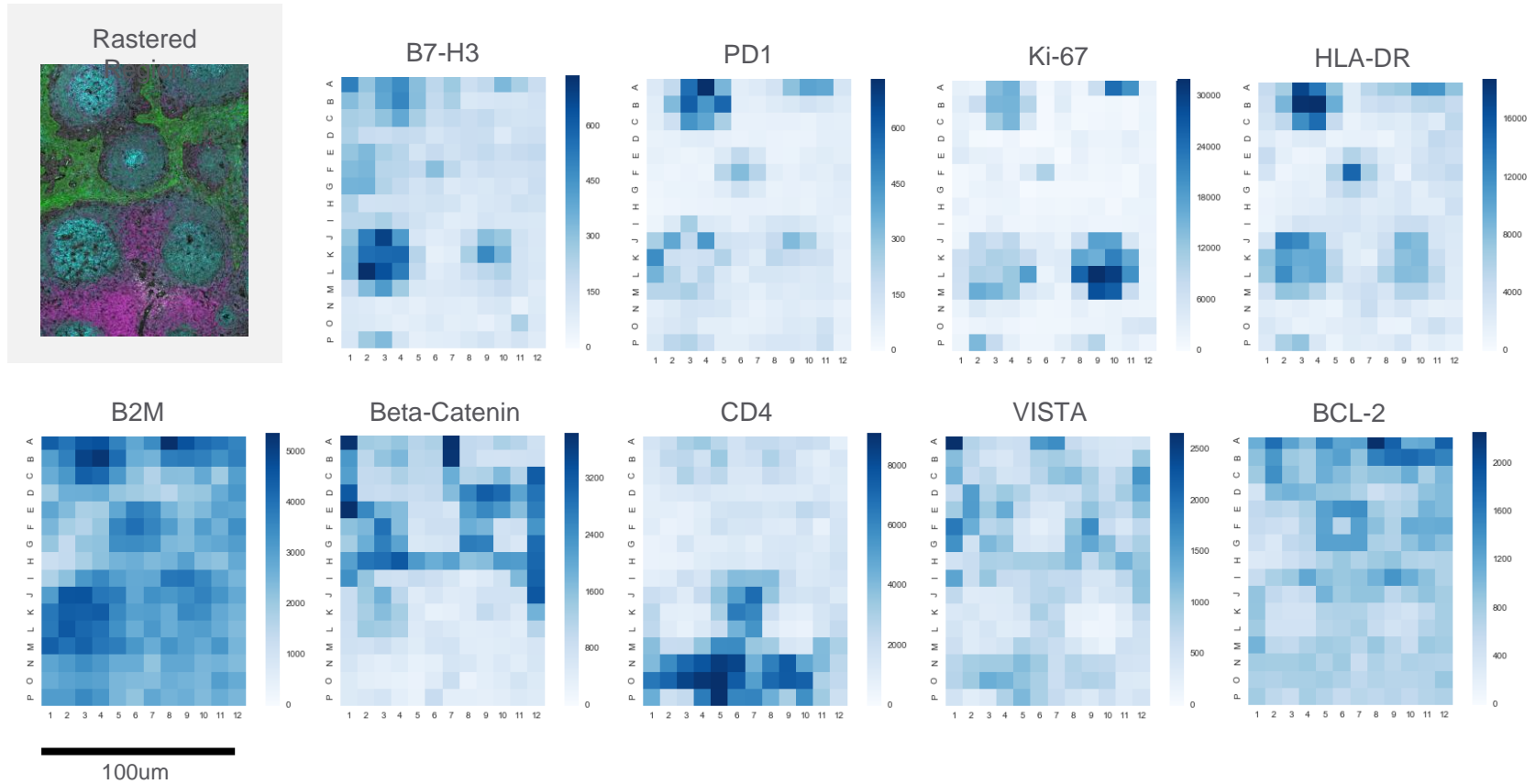
Cell numbers per
172 cells on aver
(113 min, 259 m)



Spatial & functional organization of a lymphoid tissue



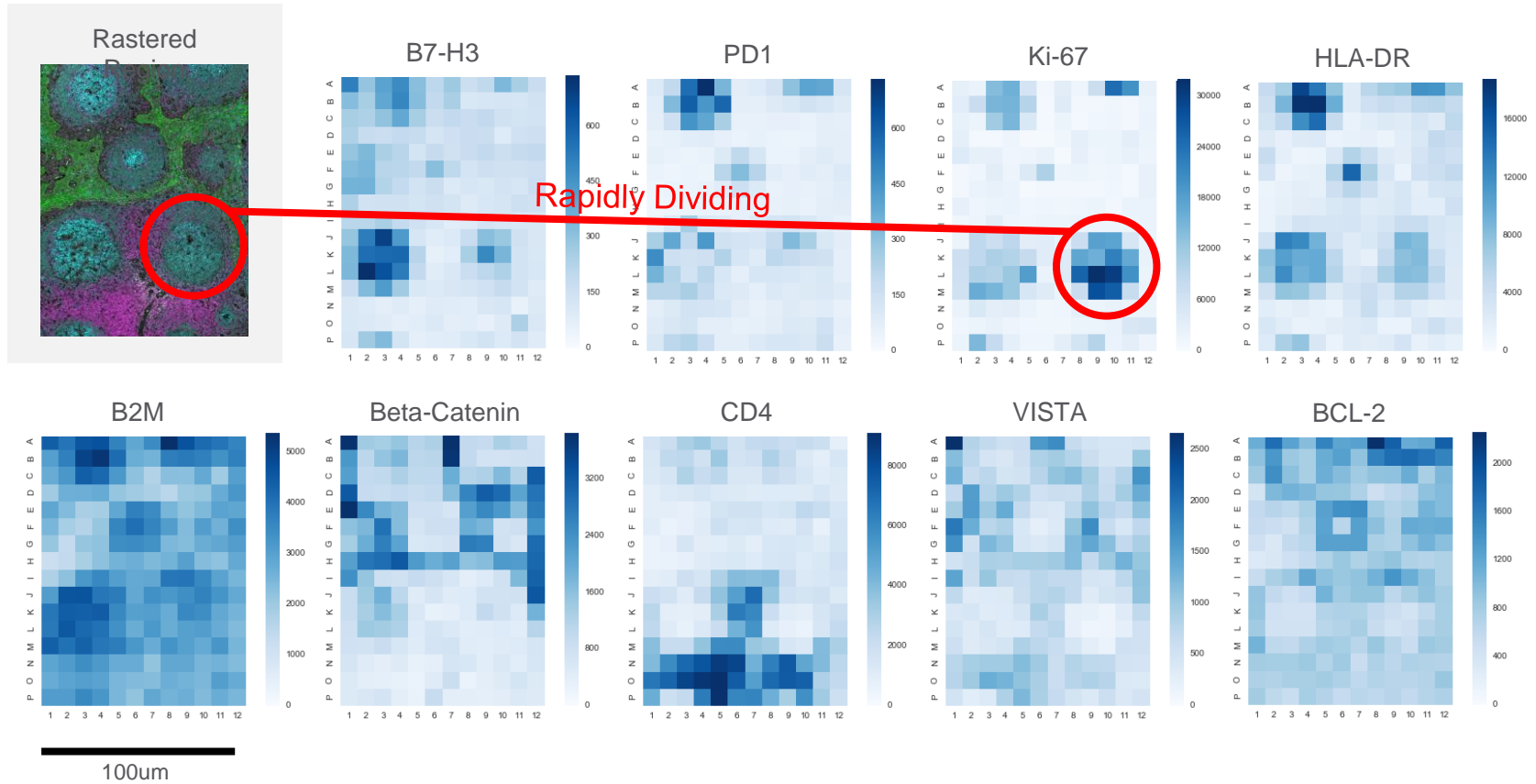
3 Examples in Immuno-Oncology



Spatial & functional organization of a lymphoid tissue



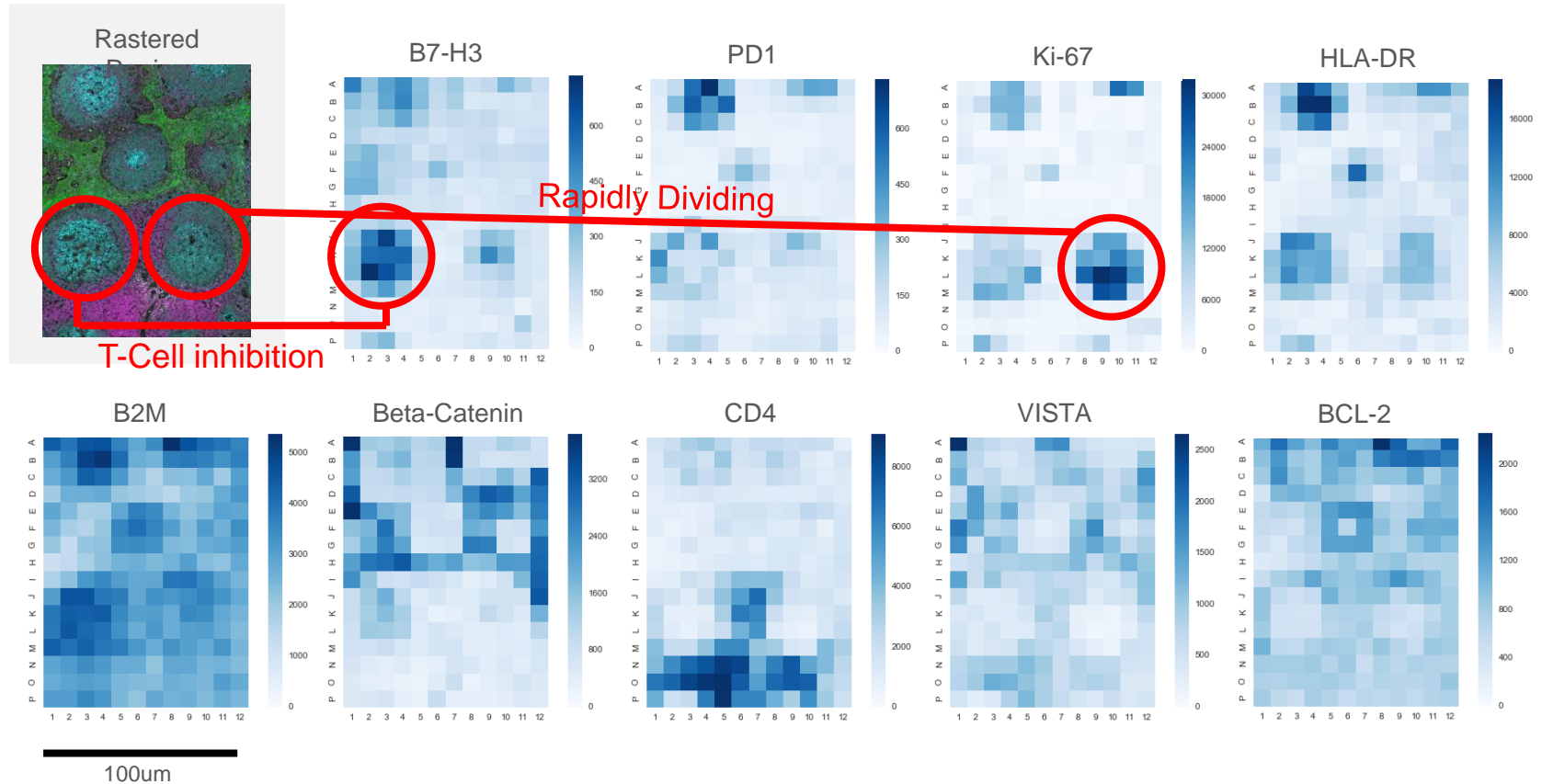
3 Examples in Immuno-Oncology



Spatial & functional organization of a lymphoid tissue



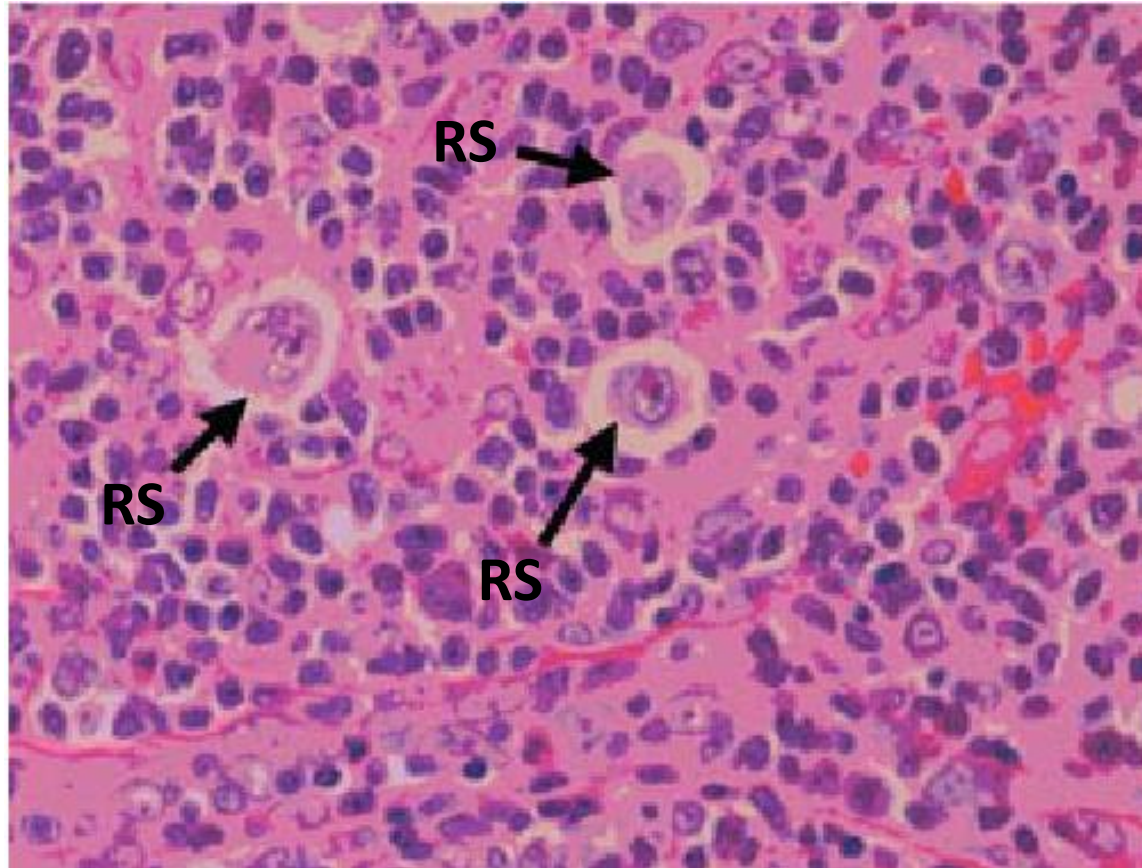
3 Examples in Immuno-Oncology



Spatial & functional organization of a lymphoid tissue



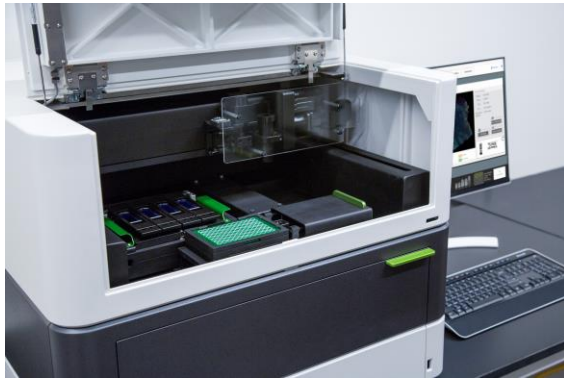
③ Examples in Immuno-Oncology



Hodgkin lymphoma > immunological gradient around CD30+ cells

Antibodies to select ROI :
CD30 (Reed Sternberg cell), **CD3** & **CD68**

4 GeoMx DSP outline



Multiplex

Many analytes on one tissue slice in a single pass

Multi-Analyte

High plex spatial analysis of both protein and RNA

Quantitation

Based on linear single-molecule counting: up to 6 logs

Single-cell

Limit of detection

Non-destructive

Sample is only touched by light

Throughput

Up to 20 sections per day

**This is working, this is easy to implement
BUT how do we integrate this level of complexity
into a clinical report that benefit to the patient?**





Hospices Civils de Lyon



votre santé,
notre engagement

jonathan.lopez@chu-lyon.fr