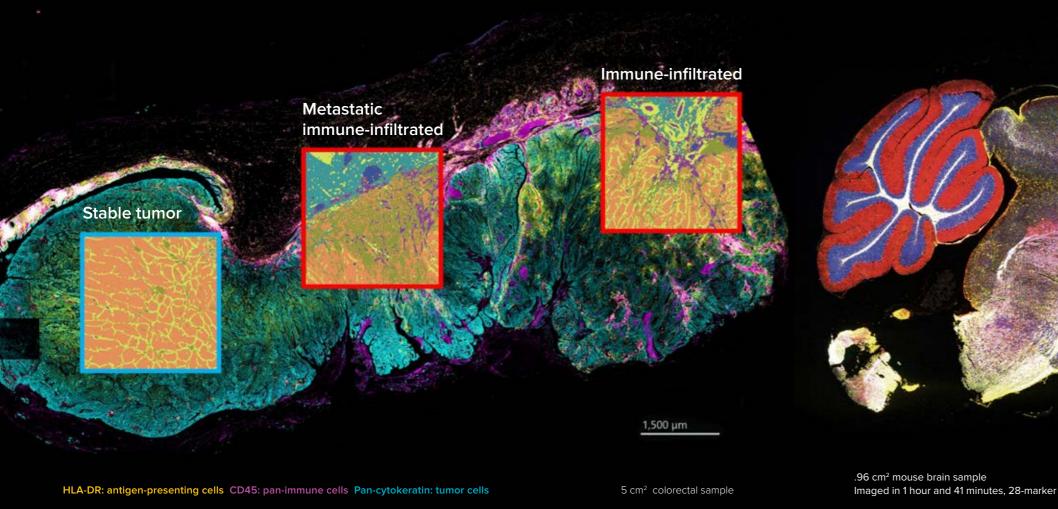
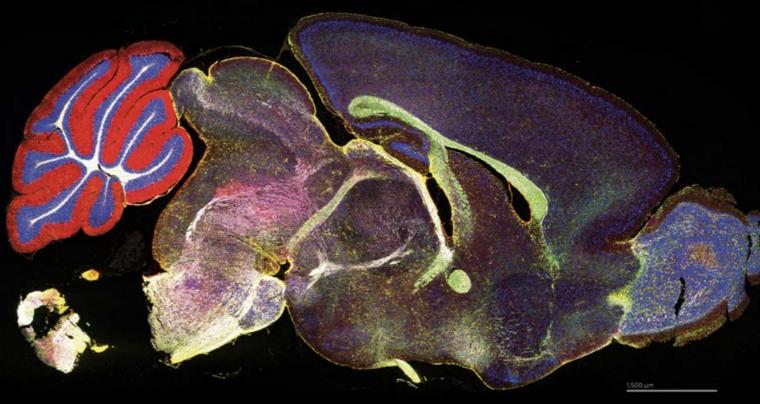


# 

# See the See th

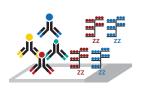




Visualize 40-plus markers in real time.

Understanding the complex tissue microenvironment is essential to the timely evaluation of disease progression and the response to the rapeutics. Imaging Mass Cytometry (IMC™) uniquely enables 40-plus protein and RNA markers to be simultaneously acquired and visualized, without time-consuming acquisition cycles.

#### A ONE-STEP MULTIPLEXED IMAGING APPROACH



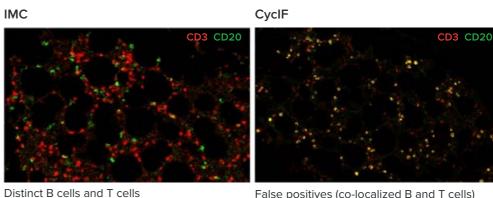




Real-time data

### Interpret data with confidence.

Quality and clarity of data are critical when deriving biological insights from limited, precious tissue samples. Clearly define areas in any tissue type – including lung, bone marrow, colon and brain – without autofluorescence interference. Imaging Mass Cytometry utilizes metal-tagged antibodies, instead of fluorophores, eliminating background autofluorescence and spectral overlap.



False positives (co-localized B and T cells)

In bone marrow, cyclic immunofluorescence (cycIF) data (right) shows false positives, highlighted by co-localization of B cells (CD20+) and T cells (CD3+) (yellow). Conversely, distinct B cells (green) and T cells (red) can be seen with IMC (left).

GFAP: astrocyte marker Neurofilament: neuronal cytoskeleton

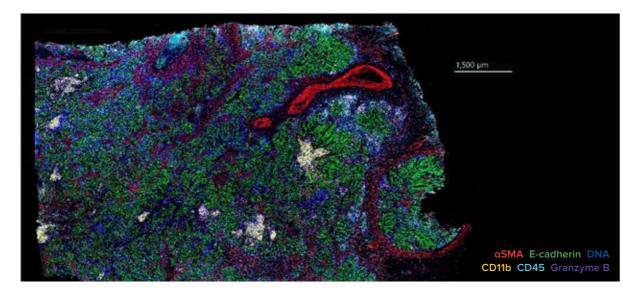
al cell bodies S100B: activated glia MBP: myelinated axon

# **PREVIEW**

# Three imaging modes to accommodate any research

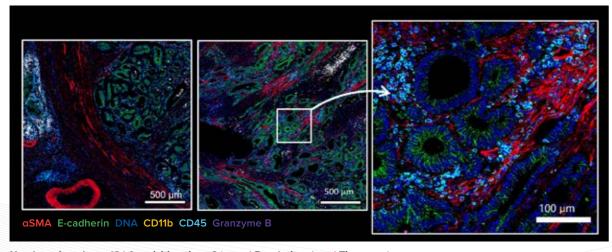
Quickly scan your tissue to preview the whole slide in 20 minutes.

Make quick decisions to identify regions of interest and subsequent acquisition mode(s).



Number of markers: 42 | Acquisition time: 20 minutes | Resolution: 1 µm subsampling (28-pixel spaces) | Tissue: colon cancel

Dig deeper within regions of interest at single-cell resolution.



Number of markers: 42 | Acquisition time: 2 hours | Resolution: 1 µm | Tissue: colon

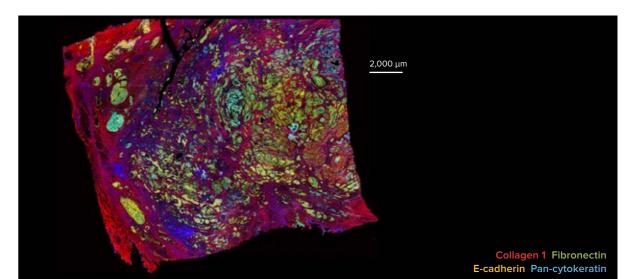
# Fast-forward SPATIAL BIOLOGY

Multiplexed spatial mapping – without compromising speed.

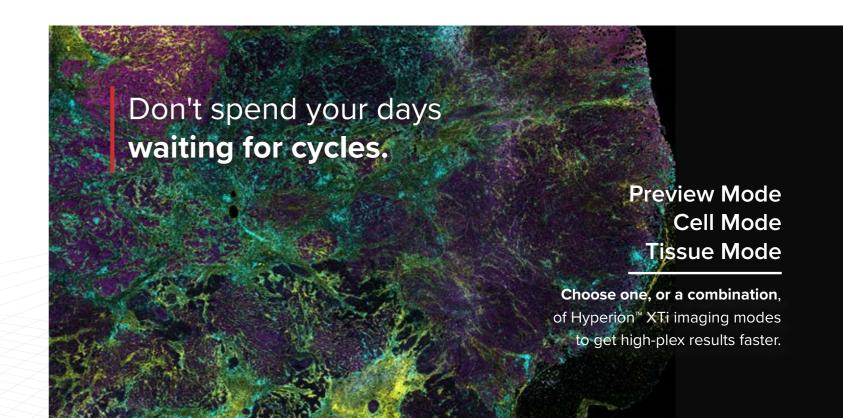
3

**TISSUE** 

Visualize heterogeneity with whole slide tissue imaging.



Number of markers: 41 | Acquisition time: 5 hours and 7 minutes | Resolution: 5 µm | Tissue: prostate cancel



# Ready-to-go panels that simplify spatial biology

### START WITH APPLICATION-SPECIFIC PANELS

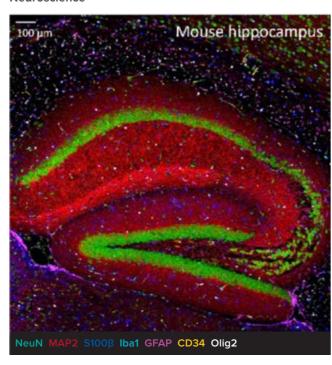
Choose from over 15 panels to easily target 40-plus markers.

### Immuno-oncology



The Human Immuno-Oncology IMC Panel, 31 Antibodies and the Human Immune Cell Expansion IMC Panel, 7 Antibodies were combined to interrogate the tumor microenvironment in breast cancer. This identified numerous cell types such as cancer-associated fibroblasts (FAP) and lymphoid (CD45) and myeloid (CD68) cells as well as epithelial-to-mesenchymal markers, cell functional states and tissue architecture identification.

#### Neuroscience

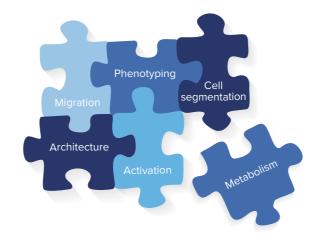


The Maxpar Neuro Phenotyping IMC Panel Kit identifies distinct spatial positioning of major brain cell lineages in a normal mouse hippocampus FFPE sample. Scale bar size = 100  $\mu$ m. Image size = 1,600  $\times$  1,600  $\mu$ m

# Easily combine panels or customize targets of interest.

Review the full list of imaging panels here.

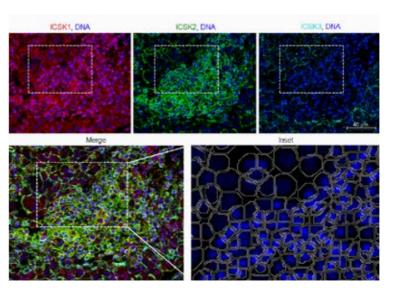




### **ADD CELL SEGMENTATION**

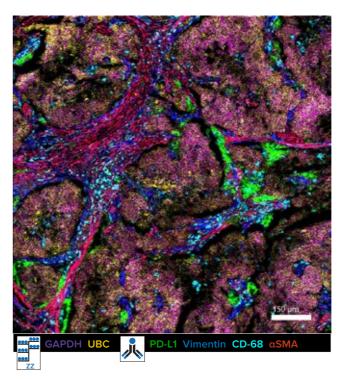
### Solving the most important step in spatial imaging.

The Maxpar® IMC Cell Segmentation Kit simplifies quantitative single-cell analysis in which cell types, cellular functions and intra- and intercellular processes can easily be defined.



The Maxpar IMC Cell Segmentation Kit contains three markers that can easily be added to existing panels.

Human formalin-fixed, paraffin-embedded non-small-cell lung cancer tissue stained with the Maxpar IMC Cell Segmentation Kit. Scale bar is 40 μm. Red, ICSK1; green, ICSK2; teal, ICSK3; blue, DNA stain. Cell segmentation was generated using Visiopharm® Phenoplex™ software.



### This image highlights RNA and protein co-detection in lung squamous cell carcinoma using the Hyperion XTi Imaging System and a 33-marker panel combined from three ready-to-go reagent kits.

### GET DEEPER INSIGHTS WITH RNA CO-DETECTION

### Combine spatial phenotyping with knowledge of the cell's transcriptome.

Detect protein and RNA on the same tissue sample to correlate transcriptional signatures and spatial context of pathogens, host cells or protein sources. Quantify mRNA, proteins and post-translational modifications to expand knowledge of cellular networks and cell type-specific gene expression.



### **WALK-AWAY AUTOMATION**

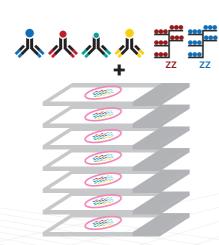
Incorporating a new level of throughput and efficiency, an integrated slide loader enables researchers to load up to 40 slides and walk away.



### **ALL-AT-ONCE STAINING**

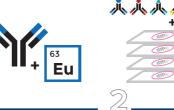
IMC uniquely enables a stain all-at-once approach to streamline experiment workflows.

Large batches of slides can be stained simultaneously to eliminate batch effects and technical variation, then stored until you are ready for analysis.



### A workflow to get results faster

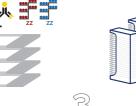
### Automated slide loader Image 40 slides per day



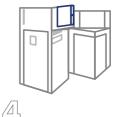
### Modularized panels Swap markers without panel revalidation.

Batch staining

Stable antibodies enable stained slides to be acquired at any time.



Bulk slide loading Load up to 40 slides for automated imaging.



Walk-away acquisition

Automatically process slides, imaging 40-plus markers per sample simultaneously.



Real-time data

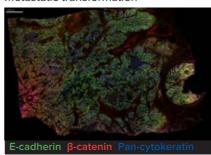
Visualize 40-plus markers in minutes.

### Analysis templates so you can work smarter.

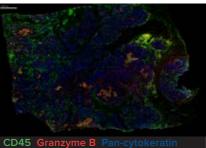
Accelerate data analysis using quick-view templates from MCD™ SmartViewer. These templates allow you to swiftly interpret high-plex data, getting you results faster.

### Quick views

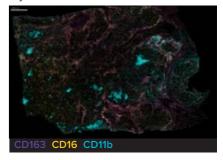
Metastatic transformation



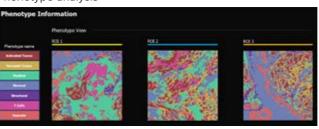
Immune cell infiltration and activation



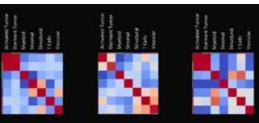
Myeloid immune cell infiltration



Phenotype analysis



Neighborhood



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