

From Complexity to Clarity



ZEISS Axioscan 7 spatial biology

Enabling Easy, Reproducible Multiplex Spatial Profiling at Scale



From Complexity...

...To Clarity.

The challenges to transition spatial biology from Basic to Clinical research

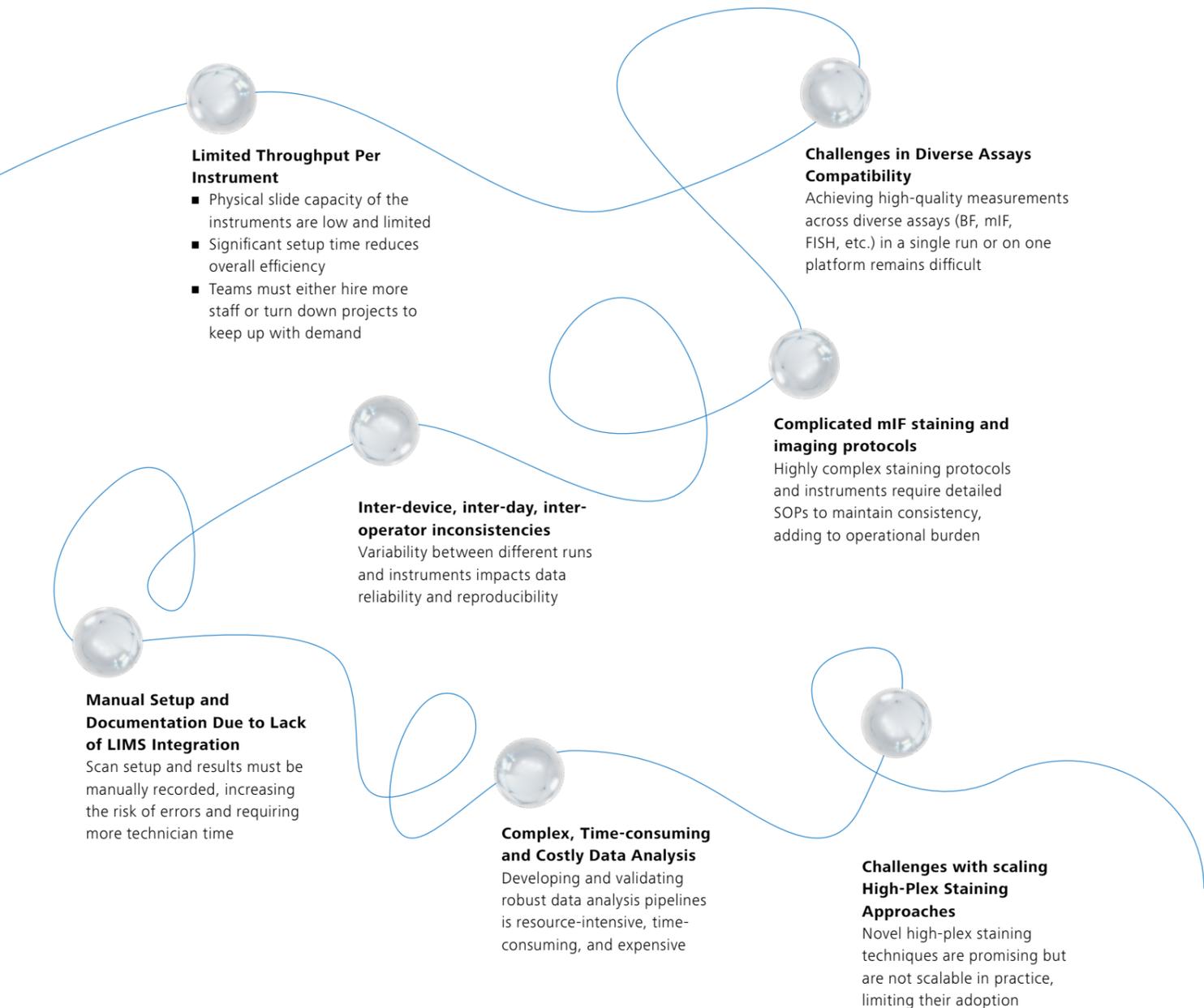
Spatial biology allows for the comprehensive analysis of cell presence, abundance, and spatial organization within their native tissue microenvironment. As the field rapidly transitions from discovery to translational and clinical applications, challenges with scalability such as low throughput, efficiency and reproducibility issues continue to hinder widespread adoption of spatial biology as a routine mainstream application. Despite its immense potential, many labs struggle to integrate spatial biology into their standard routine workflows.

Fast and high-throughput Imaging & Analysis

1-Click to effortlessly scan and analyze 100+ slides per day

With our ZEISS spatial biology offering, we simplify high-throughput multiplex IF imaging and analysis, making it robust, scalable, and accessible. Our seamless **tissue multiplexing workflow** aims to empower routine histopathology labs, including those with no prior spatial biology expertise, to generate comprehensive, reproducible biomarker data across large sample cohorts.

By combining automation, optimized imaging, and AI-powered analysis, we pave the way for the translational adoption of spatial proteomics and the transition to clinical practice.



One Streamlined Workflow



1-Click to Start and walk away

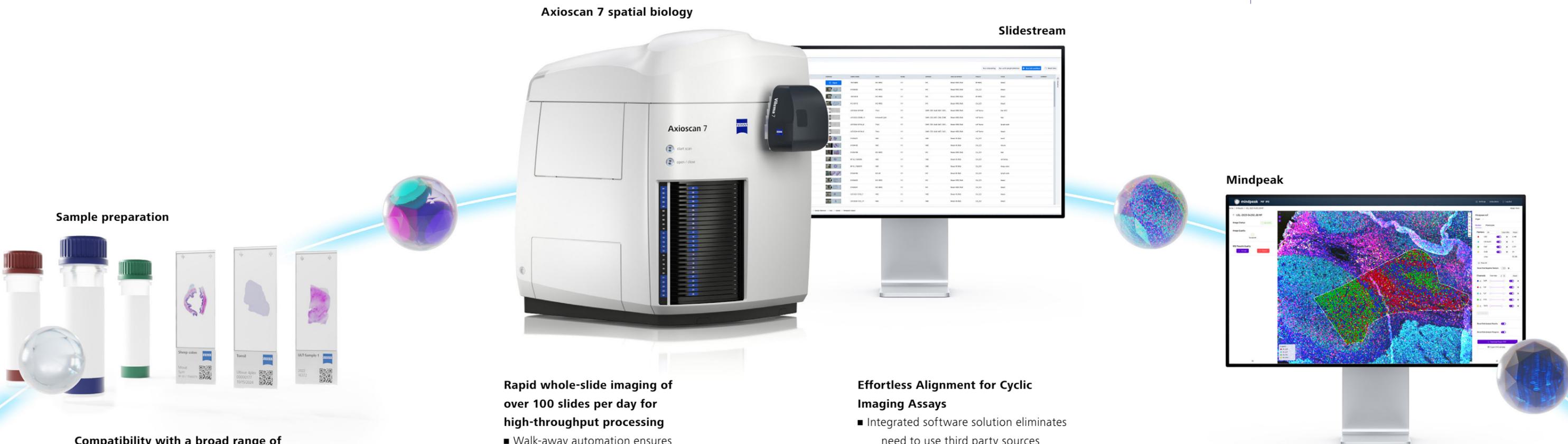
SlideStream Streamlines your workflow

Broad compatibility. High throughput.

Our workflow automation allows reading a vast variety of barcode formats used in autostainers and is compatible with a broad range of multiplex immunofluorescence (mIF) and chromogenic reagents and assays. Scan over 100 mIF and brightfield slides per day without manual intervention.

Automated Workflow

-  **Sample preparation**
Compatible assays and reagents
-  **Axioscan 7 spatial biology**
Automated HDR imaging
-  **SlideStream**
Workflow Manager Software
-  **Mindpeak**
AI-powered image analysis



Sample preparation

Compatibility with a broad range of reagents chemistries and assays

- One instrument to scan mIF and IHC stained samples
- Streamlined End-to-End workflow, from sample to report, eliminates the need for specialized technical expertise

Axioscan 7 spatial biology

Rapid whole-slide imaging of over 100 slides per day for high-throughput processing

- Walk-away automation ensures seamless operation, with the scanner automatically initiating slide analysis after scanning
- Physical capacity of 100 slides per run

Seamlessly scan and analyze 8-plex images on a single slide

- Examine cell-cell spatial relationships within the tumor microenvironment and evaluate expression across multiple ADC targets

SlideStream

Effortless Alignment for Cyclic Imaging Assays

- Integrated software solution eliminates need to use third party sources

Advanced automation

- Accelerate your work with AI-powered software features like never before
- Custom services for integration into LIMS

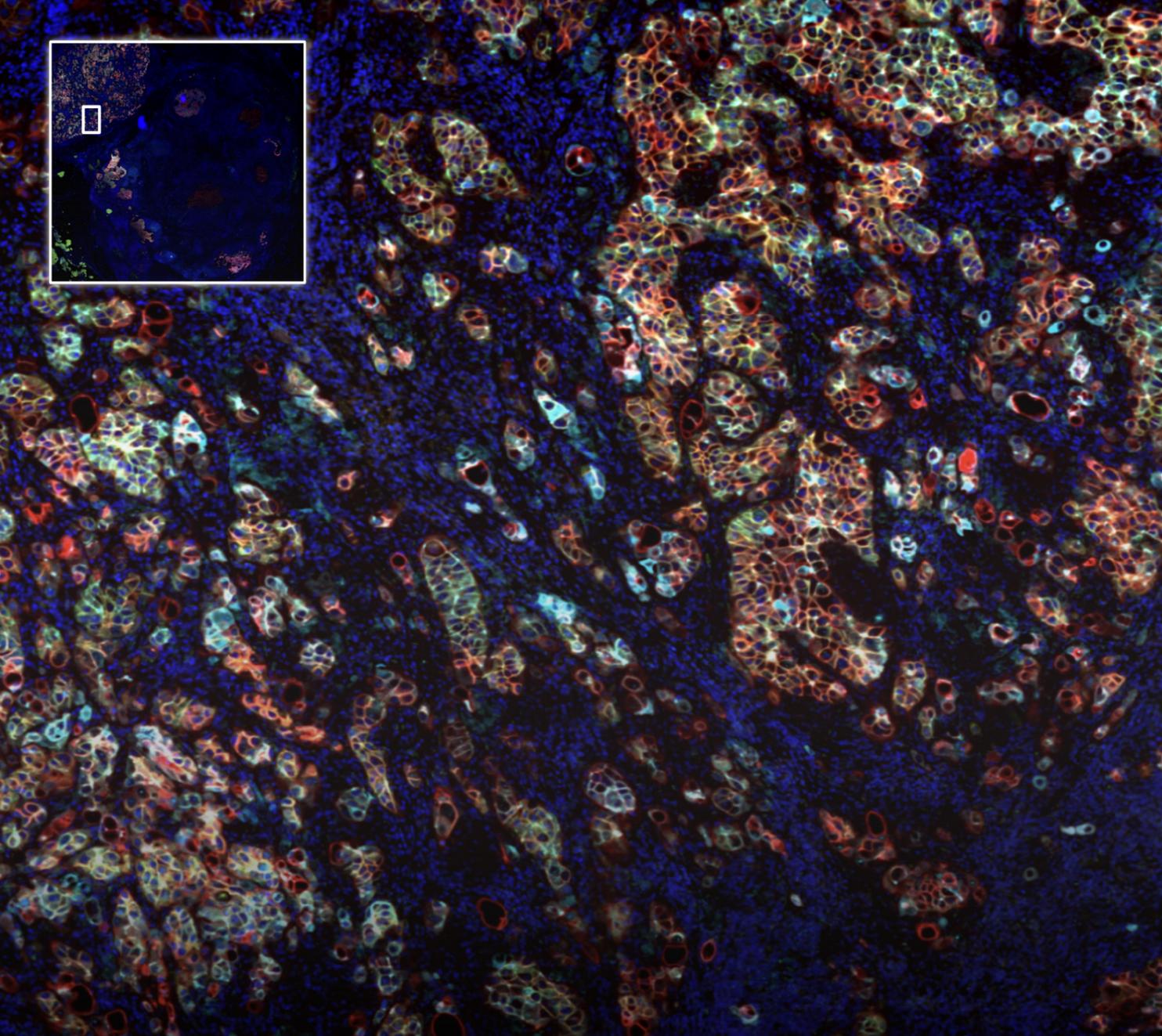
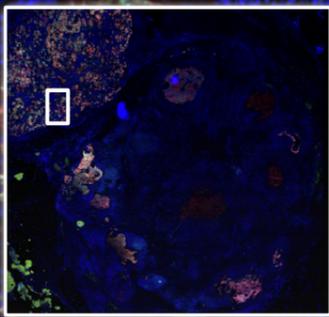
High reproducibility

- Achieved with smart combination of highly robust hardware and software components

Mindpeak

Automated spatial mIF Results

- Robust, pre-trained AI analysis provides reproducible spatial mIF data independent of the operator



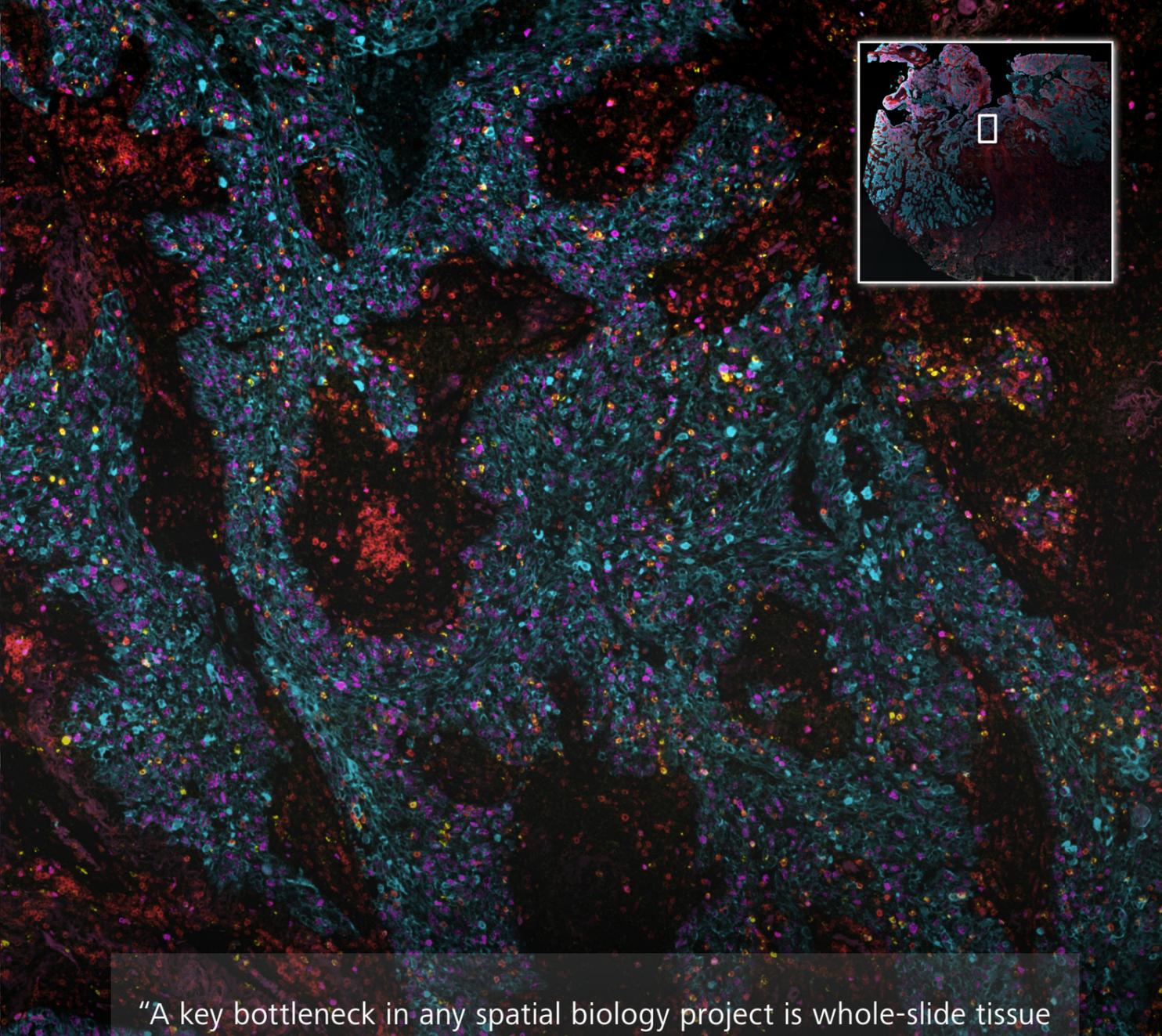
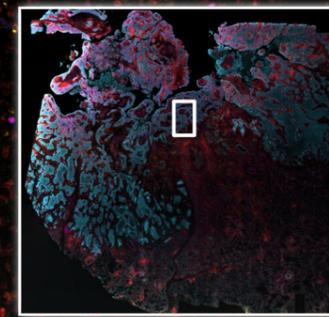
“The seamless integration of reagents, Axioscan 7, SlideStream, and Mindpeak analytics offers an incredibly user-friendly and fully automated workflow. It has transformed the way we generate spatial phenotypic data; fast, reproducible, and scalable for our clinical research.”

**Prof. Christoph Kuppe, MD, PhD,
University Clinic RWTH Aachen**

A composite image of bladder cancer tissue

TROP2 HER2 NECT4 HER3 DAPI

The image shown on this page represents research content, courtesy of University Clinic Aachen. ZEISS explicitly excludes the possibility of making a diagnosis or recommending treatment for possibly affected patients on the basis of the information generated with an Axioscan 7 slide scanner.



“A key bottleneck in any spatial biology project is whole-slide tissue scanning, which can lead to unplanned reviews and repeat imaging, making it difficult to predict delivery timelines and associated costs. ZEISS is addressing this challenge with an efficient and reproducible automated workflow that truly benefits CROs offering this service. I’m thrilled to be collaborating with ZEISS in advancing scanning technology and delivering superior outcomes for our clients.”

**Christopher Mills, Senior Scientist
Concept Life Sciences (CRO in United Kingdom)**

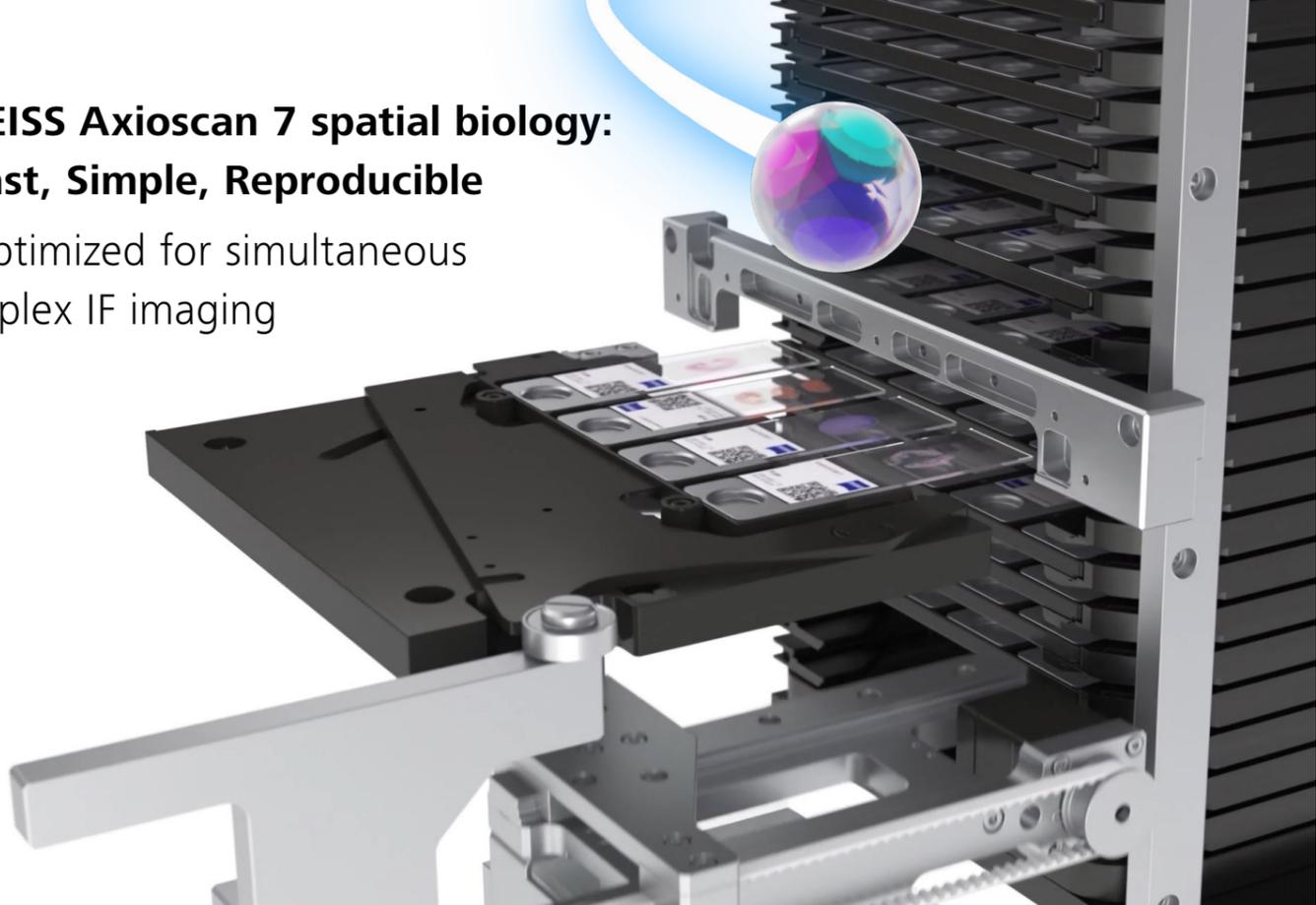
A composite image of Non-Small Cell Lung Cancer tissue

Ki67 GranzymeB CD3 CK/SOX10

The image shown on this page represents research content, courtesy of Concept Life Sciences. ZEISS explicitly excludes the possibility of making a diagnosis or recommending treatment for possibly affected patients on the basis of the information generated with an Axioscan 7 slide scanner.

ZEISS Axioscan 7 spatial biology: Fast, Simple, Reproducible

Optimized for simultaneous
8-plex IF imaging



In today's fast-paced research environments, CROs, pharmaceutical industry, and high-throughput imaging labs demand faster, more reliable and reproducible, and automated solutions for spatial biology.

The Axioscan 7 spatial biology configuration is designed to address these needs, delivering exceptional speed, sensitivity, and workflow integration optimized for 8-plex multiplex immunofluorescence (mIF) imaging.

Fast multiplexing imaging

Seamlessly scan up to 8-plex images with superior speed: 10 mm² tissue area can be scanned in 3 to ~10 minutes for 5- or 8-Plex respectively.

Superior Sensitivity

Reveal fine subcellular details with 170 nm pixel size (high-resolution mode) or 345 nm (high-sensitivity mode). Advanced optics and a high-performance camera deliver exceptional clarity in multiplexed samples.

Advanced HDR Imaging

Capture the full dynamic range from the dimmest to the brightest signals and stop wasting time to find multiple optimized exposure time settings to address staining intensity variability in your samples.

Advanced automation

Accelerate your work with enhanced barcode recognition, AI-driven mIF tissue detection, and pre-defined scan profiles optimized for robust mIF and H&E imaging.

Best-in-class reproducibility

The combination of highly robust light output of the new Viluma LED light source and an improved shading correction provides optimal image quality for tiled imaging and highly reproducible image intensities across devices.

Efficiency and workflow integration

Optional software bundle (SlideStream and Mindpeak) support routine users with a highly automated and streamlined pipeline for image acquisition and image analysis and the possibility of custom integration into LIMS systems.

Effortless Alignment for Cyclic Imaging Assays

Integrated co-registration feature ensures seamless, precise alignment of images across multiple imaging rounds, enhancing reproducibility and data integrity.

Custom Services

System upgrades and customization
Complimentary LIMS integration and lab automation
Custom AI analysis development and deployment

Introducing SlideStream Workflow Manager

Fast, Scalable, and Effortless Multiplex IF Imaging and Data Analysis

3 Main Steps to get from Sample to Report

Efficiency, Reproducibility, Ease-of-Use



POSITION	STATUS	OVERVIEW	SAMPLE NAME
(1, 1)	Analyzing		70016809
(1, 2)	Analyzing		01694283
(1, 3)	Analyzing		10016018
(1, 4)	Analyzing		01218713
(2, 1)	Analyzing		ULT-2024-101949
(2, 2)	Analyzing		ULT-2022-23098_11

Autoselect scan and analysis

Barcodes on slides are used to retrieve slide information from the integrated sample data base and automatically select the appropriate pre-optimized scan and analysis routine.

POSITION	STATUS	OVERVIEW	SAMPLE NAME
(1, 1)	Done		70016809
(1, 2)	Done		01694283
(1, 3)	Uploading		10016018
(1, 4)	Uploading		01218713
(2, 1)	Uploading		ULT-2024-101949
(2, 2)	Uploading		ULT-2022-23098_11

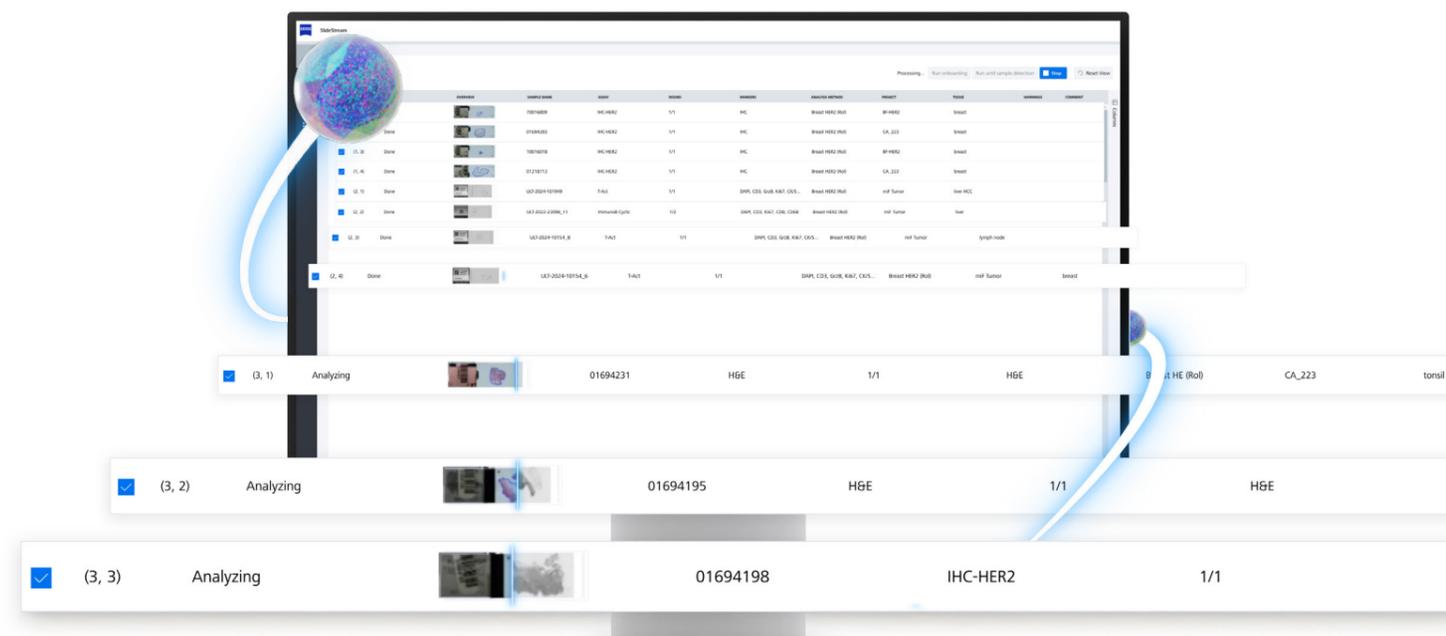
Automated image upload

Fully automated image upload and analysis trigger without any manual intervention

POSITION	STATUS	OVERVIEW	SAMPLE NAME
(1, 1)	Done		70016809
(1, 2)	Done		01694283
(1, 3)	Uploading		10016018
(1, 4)	Uploading		01218713
(2, 1)	Uploading		ULT-2024-101949
(2, 2)	Uploading		ULT-2022-23098_11

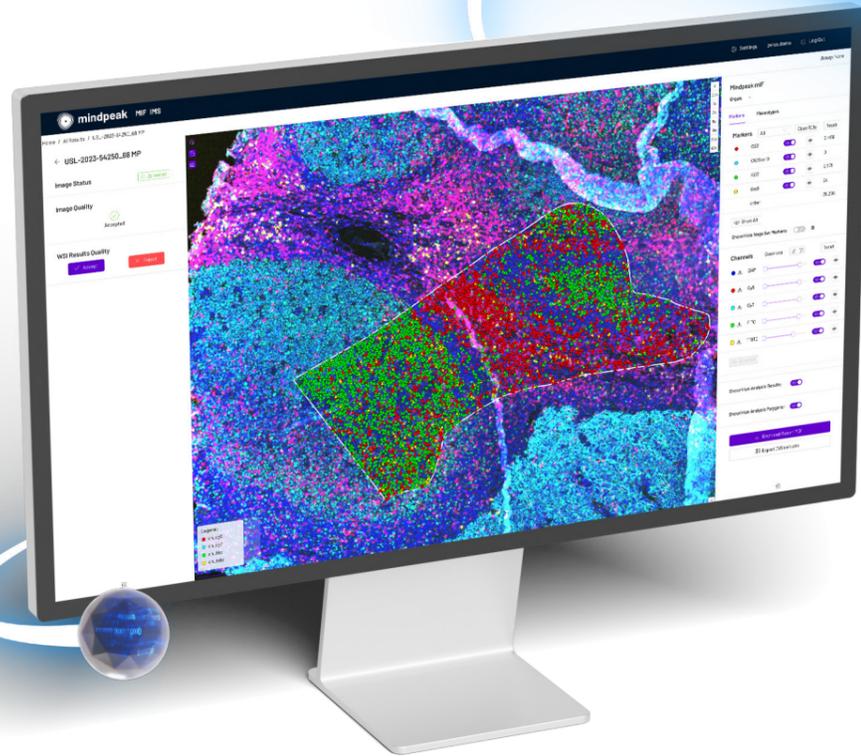
Integration services

Services for custom workflow automation, such as integration of Laboratory Information Management System (LIMS) and Image Management System (IMS) for smooth workflow optimization and data management, and many more adaptations.



AI-powered 1-click automated mIF image analysis

Image In, Report Out

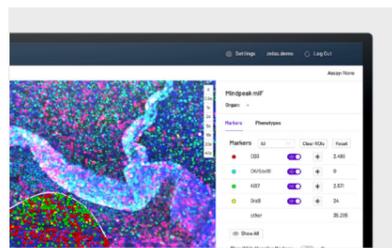
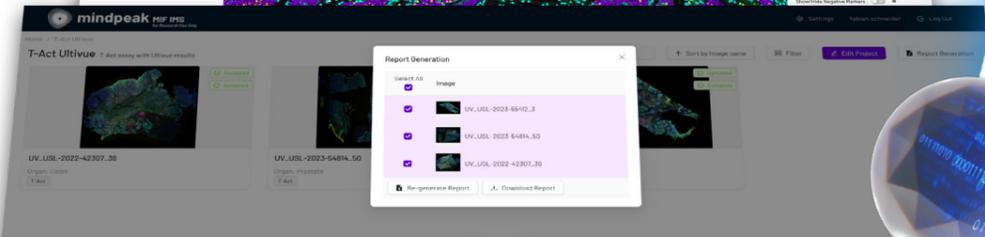
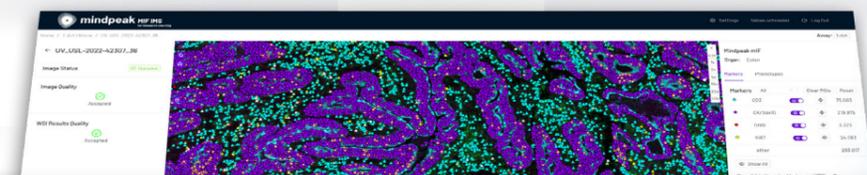
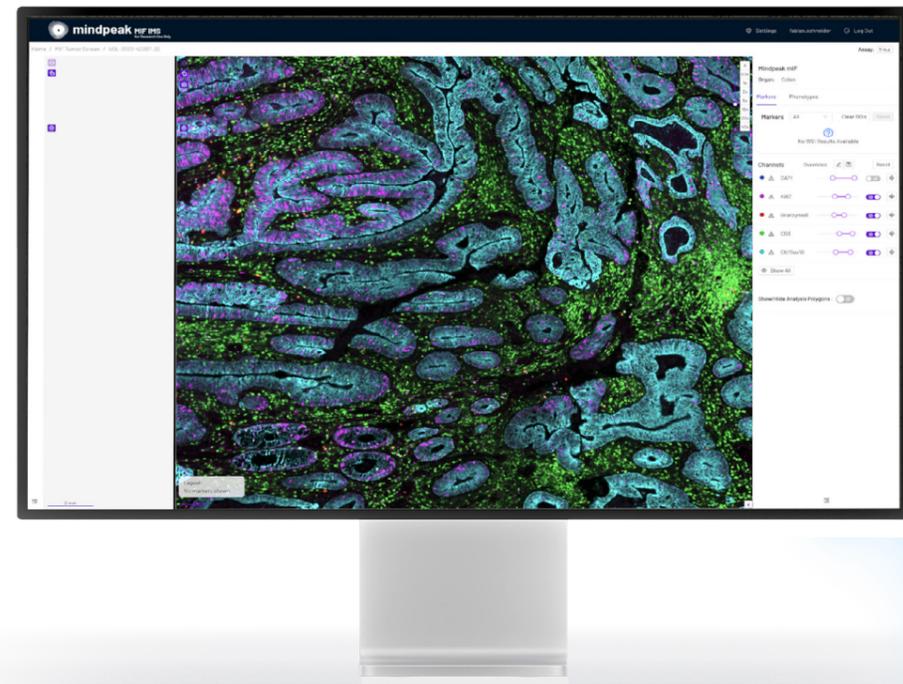


Robust, pre-developed AI-based analysis for supported assays.

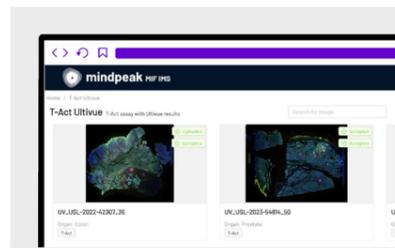
No image analysis skills needed.

Mindpeak AI-based analysis

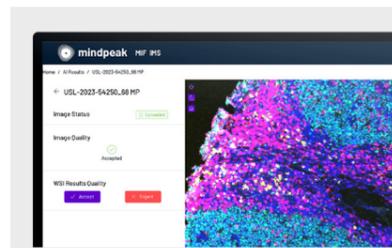
Automated image annotation and report generation



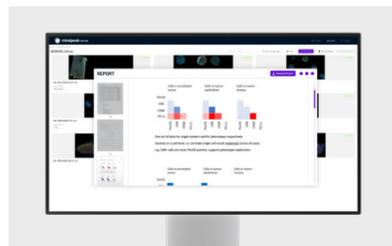
Easy and Intuitive to Use



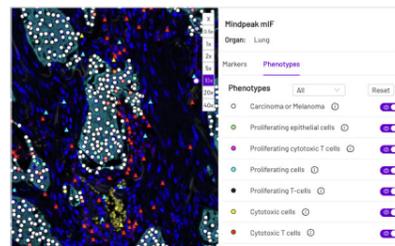
Browser Access to images and analysis - anywhere, anytime



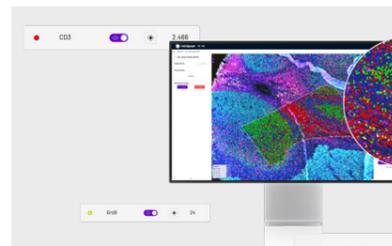
Review image and analysis quality and document QC results



Ready to download report



Standardized reports with User defined Phenotypes



Custom AI algorithm development services up to IVD certified use cases possible

mindpeak **ZEISS**

Immuno-Oncology Assay Report

1. Executive Summary
This report summarizes the multiplex immunofluorescence (mIF) analysis performed on the patient's lung tissue sample. The spatial distribution and density of key immune and tumor markers was quantified to characterize the tumor microenvironment (TME) and assess the status of T cell infiltration. Our findings indicate a predominantly inflamed TME with high CD8+ T cell infiltration into the tumor core. These results may suggest a positive response to immunotherapy.

2. Patient & Sample Information

- Patient Age: 62
- Patient Gender: Male
- Primary Diagnosis: Adenocarcinoma
- Sample Site: Tumor core

3. Marker Expression & Quantification

Marker	Density in Stroma (#/mm ²)	Typical range	Remarks	Density in Epithelium (#/mm ²)	Typical range	Remarks
CD3	2560	30-250	High	560	0-20	High
CD4	1204	12-250	High	40	0-5	High
CD8	1356	65-645	High	523	0-18	High
FOXP3	30	0-52		2	0-2	
Ki67	326	26-316	Elevated	1189	500-2500	
CD68	220	10-250		13	1-10	Elevated
CD206	350	15-500		15	10-30	
pan-CK	1	0-2		4700	3000-5000	

mindpeak **ZEISS**

Key Interactions: CD8+ T cells are frequently observed in direct contact with PanCK+ tumor cells. FoxP3+ Treg cells show some clustering near CD68+ macrophages.

Spatial Distribution: Immune cells are predominantly located in the stroma, but a significant population of CD8+ T cells has infiltrated the tumor core.

6. T Cell Infiltration Score

- Infiltration Status: Inflamed
- Score: 8/10 (Example Score)
- Description: The tumor exhibits an "inflamed" phenotype, characterized by a high density of CD8+ T cells within the tumor core and close proximity to tumor cells. This suggests an active immune response against the tumor.

7. Pathologist's Comments
The high density of CD8+ T cells within the tumor suggests a favorable microenvironment for immune checkpoint inhibitors. Further investigation is warranted.



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