

Development of a Multi-Modal Imaging Platform for the Analysis of Patient Biopsies In Translational Studies

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School of
Medicine

What is a biopsy for?

“Medical procedure that involves taking a small sample of body tissue so it can be examined under a microscope”

But what if we want to look at what's happening at the molecular level – what you can't see under a microscope?

Drug discovery and drug metabolism

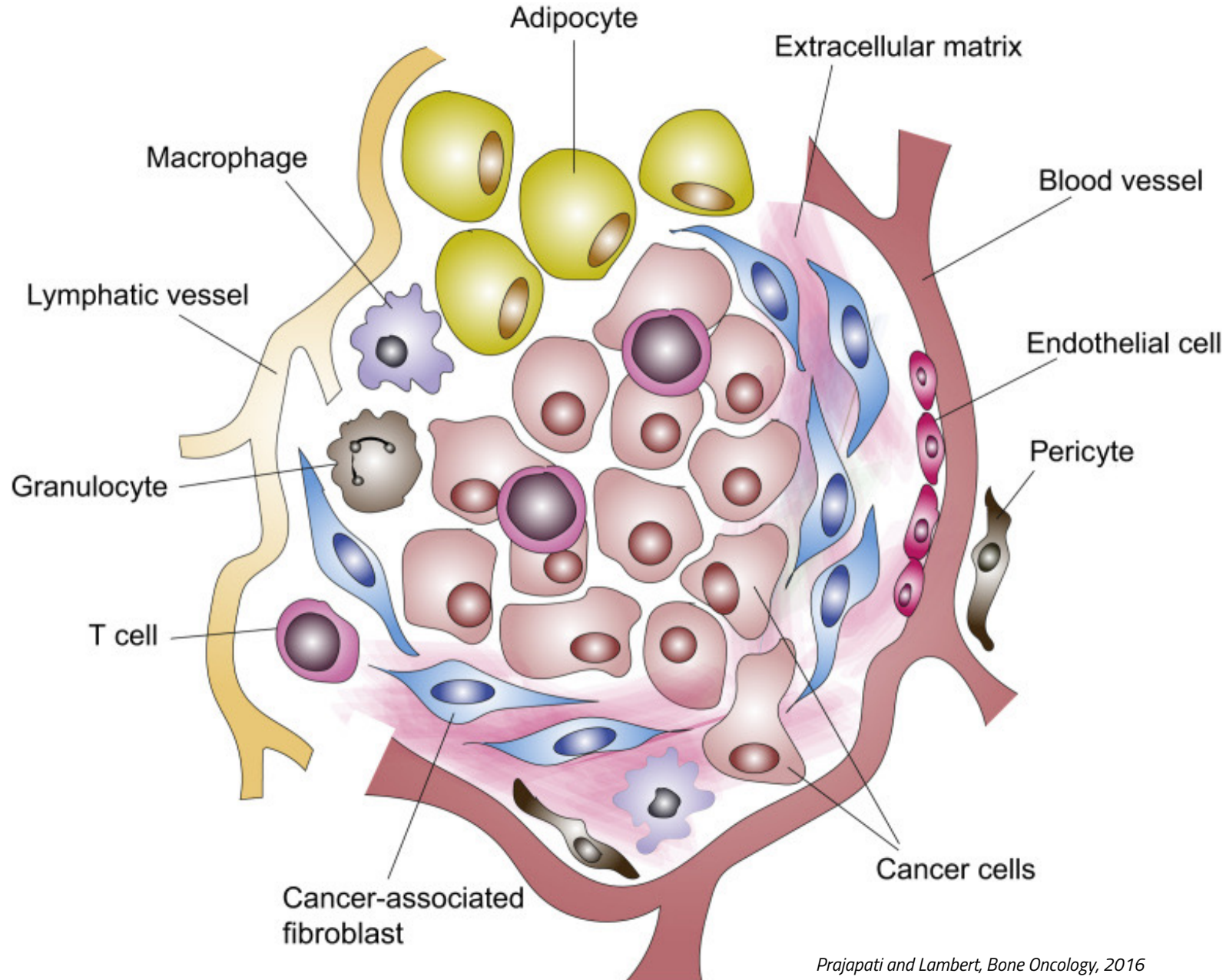
- Pharmacokinetic (PK)
- Pharmacodynamic (PD)

Two approaches to process patient biopsy

- Mash it up “grind and find” (homogenise)
- *in situ* analysis



Tissue is a lot more complex...

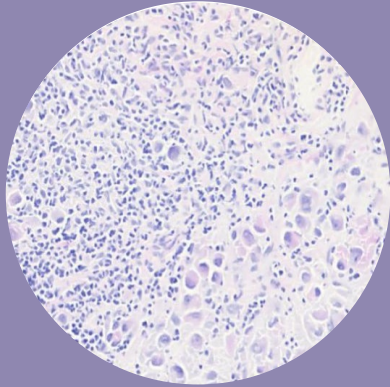


- Understanding of the cellular microenvironment
- Tissue architecture
- Immune microenvironment
- Phenotyping

Imaging Toolkit

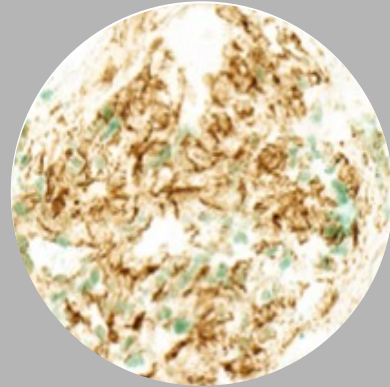
High Throughput

Large Data Volumes



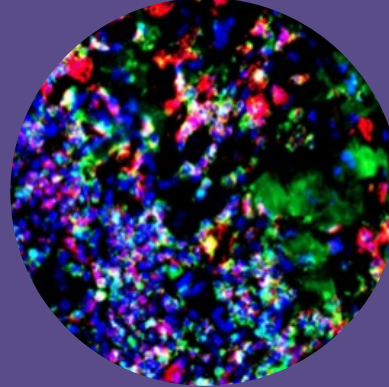
Histology

Pathology
Morphology



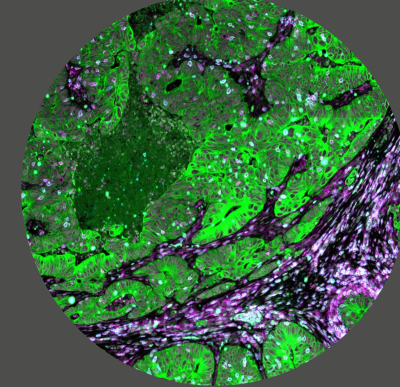
mIHC

Immuno-
Histochemistry



mIF

Immuno-
Fluorescence



ST

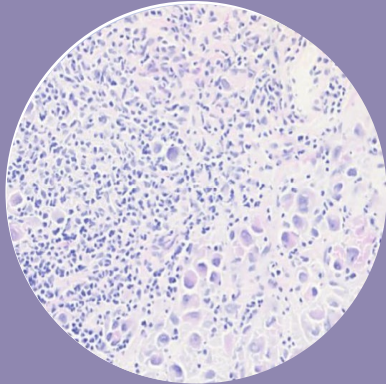
Spatial
Transcriptomics

Data Integration for AI and Machine Learning

Imaging Toolkit

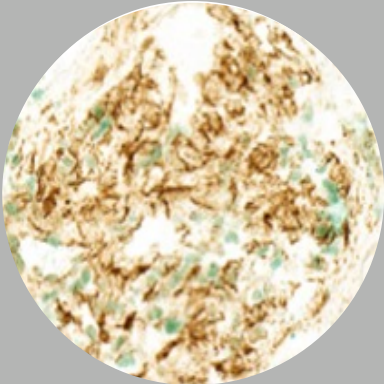
High Throughput

Large Data Volumes



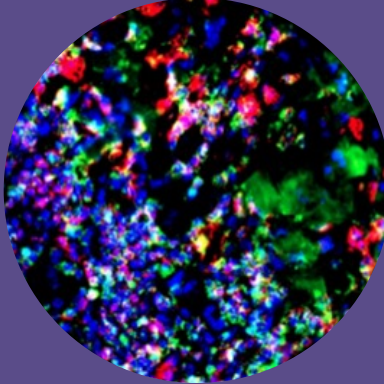
Histology

Pathology
Morphology



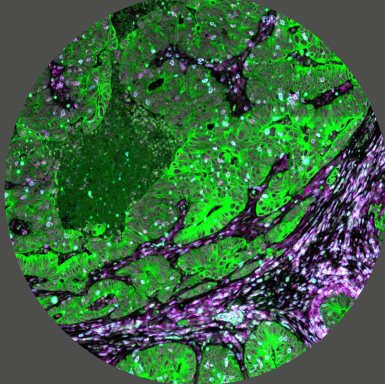
mIHC

Immuno-
Histochemistry



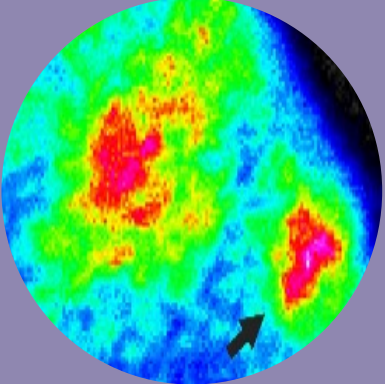
mIF

Immuno-
Fluorescence



ST

Spatial
Transcriptomics

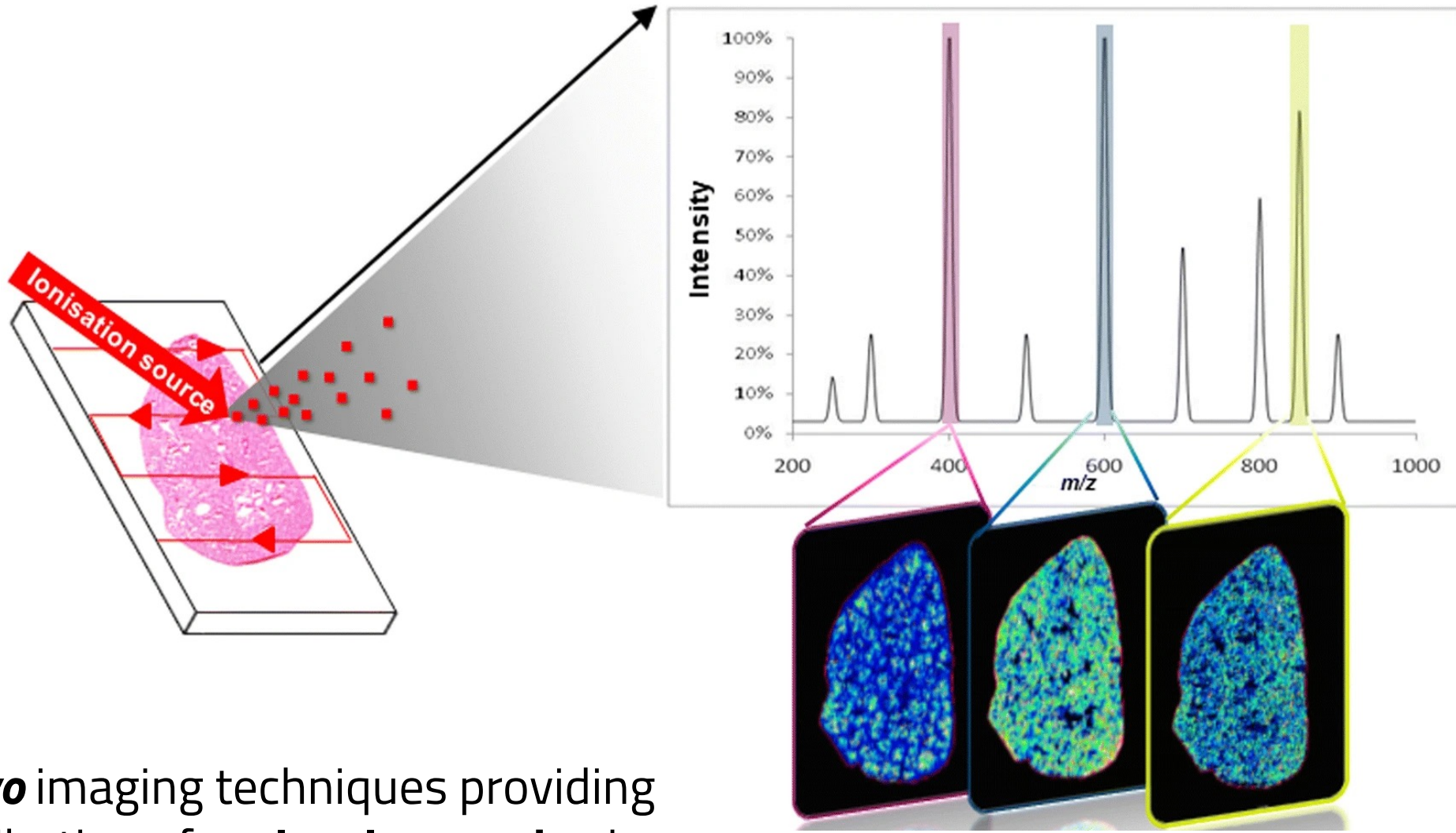


MSI

Mass
Spectrometry
Imaging

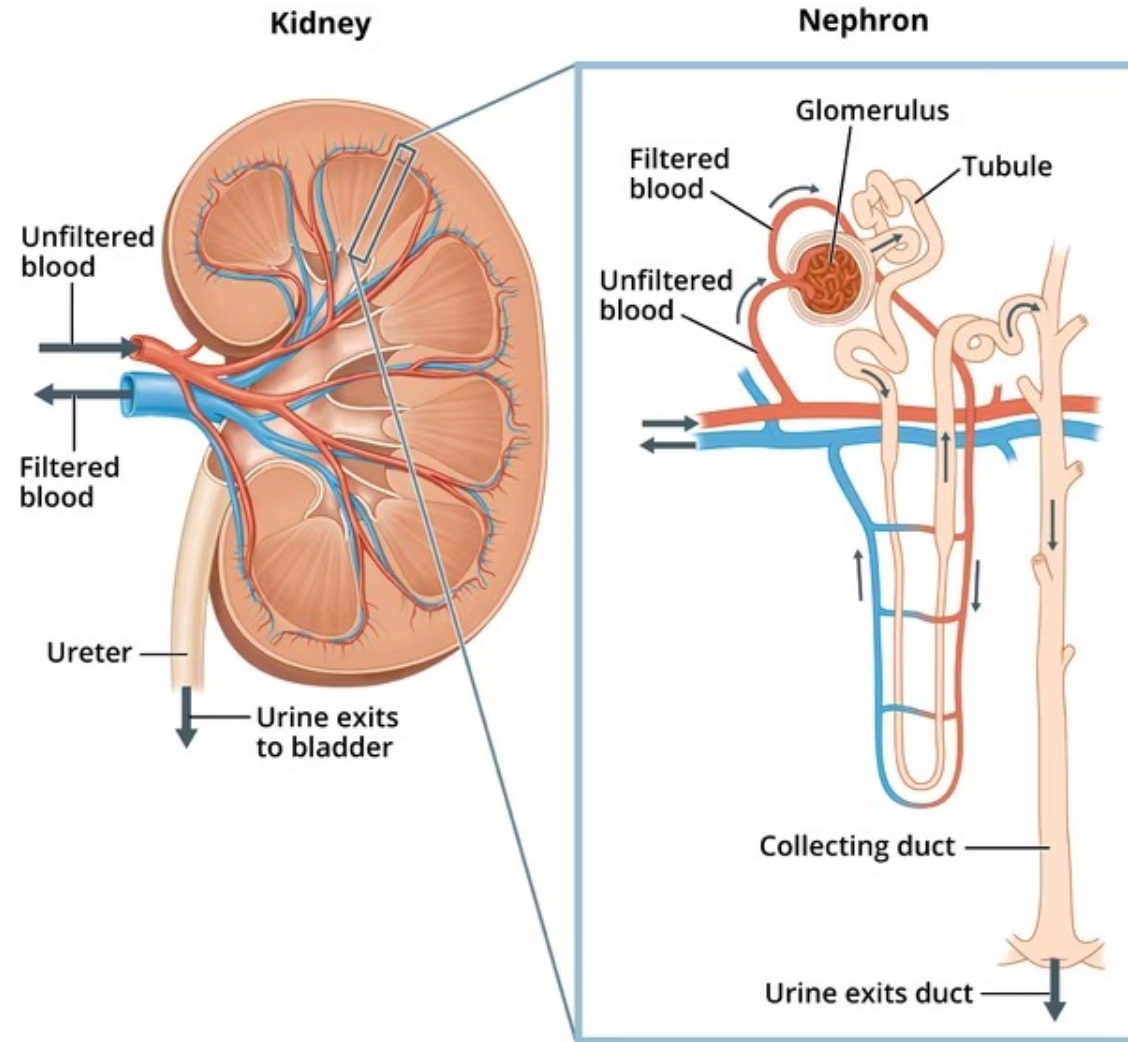


Mass Spectrometry Imaging (MSI)

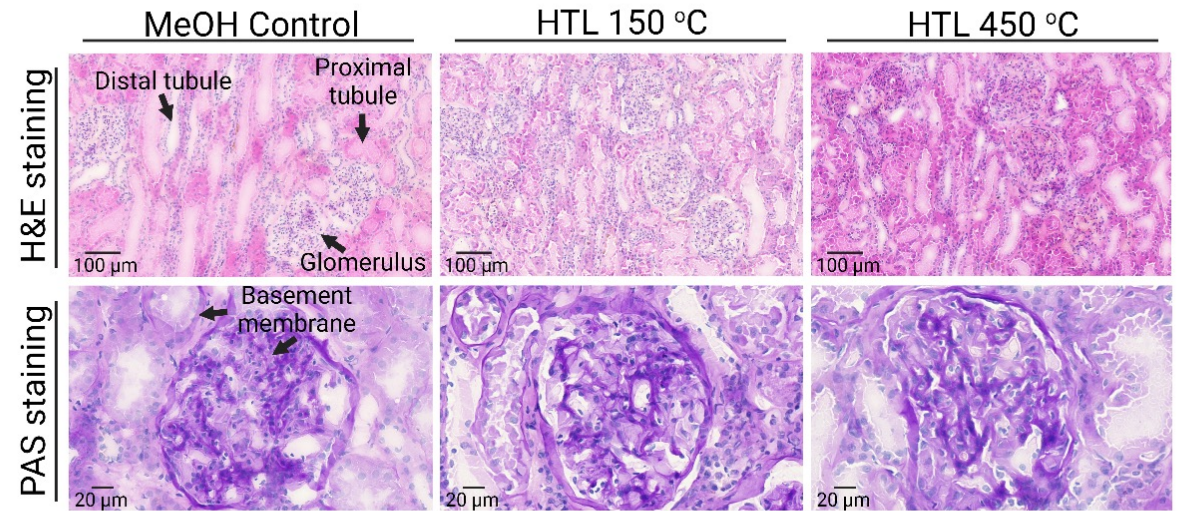
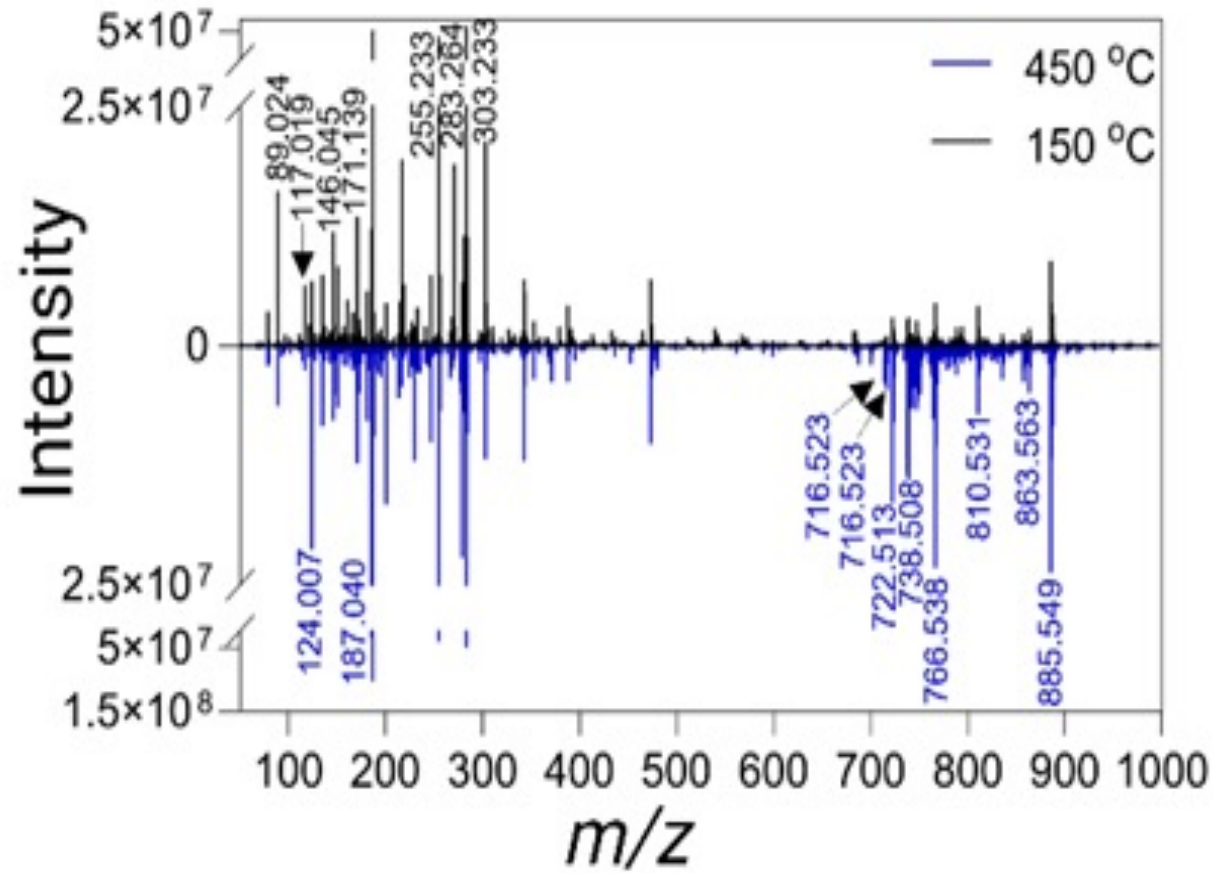


Set of ***ex-vivo*** imaging techniques providing spatial distribution of **molecular species** in **biological tissue**

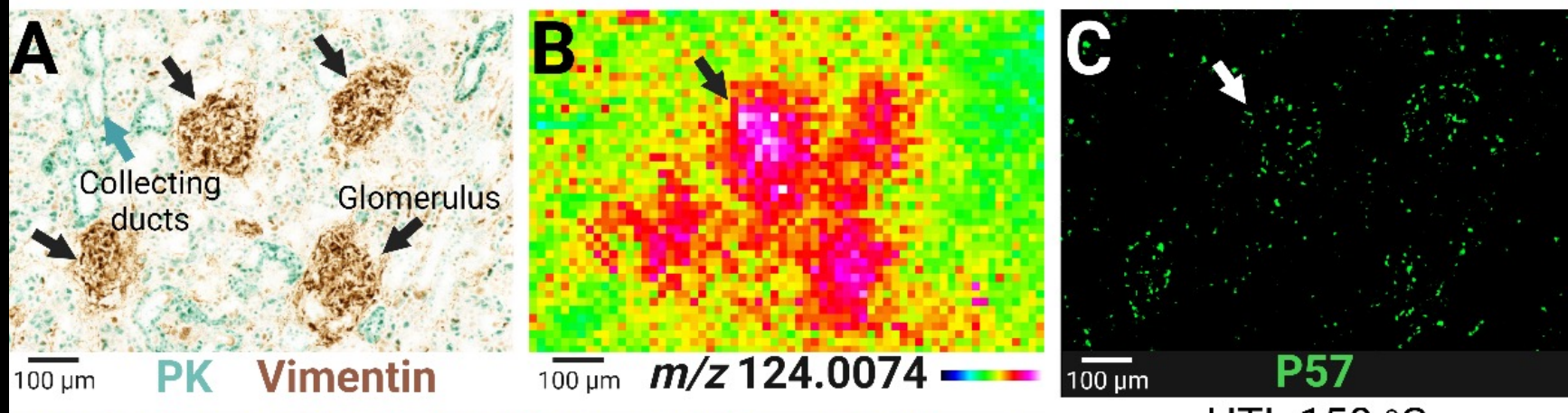
Take the kidney for example...



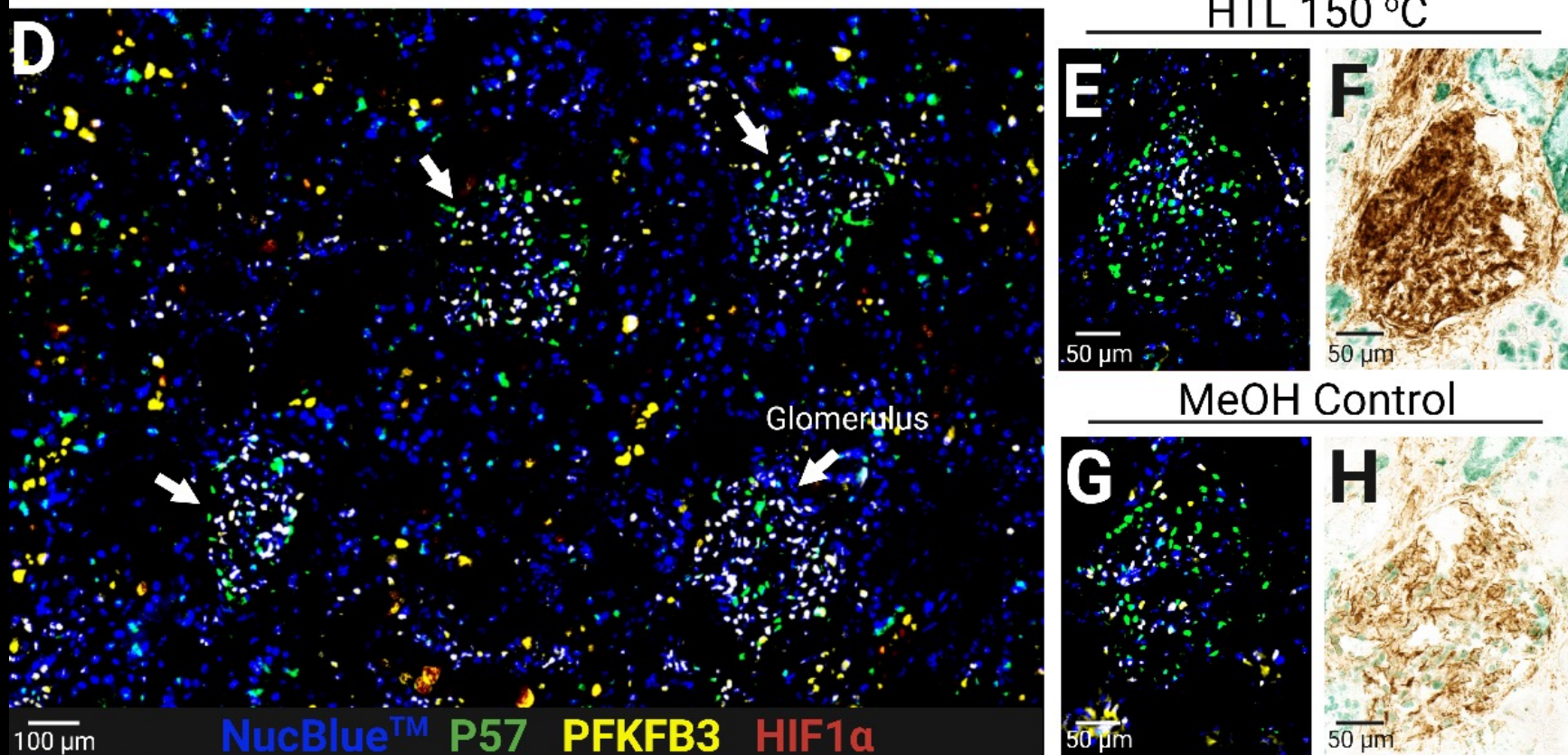
What's cooking with DESI-MSI?



HTL 450 °C

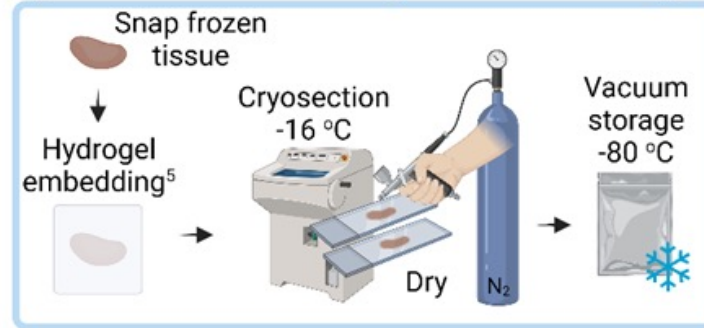


HTL 150 °C

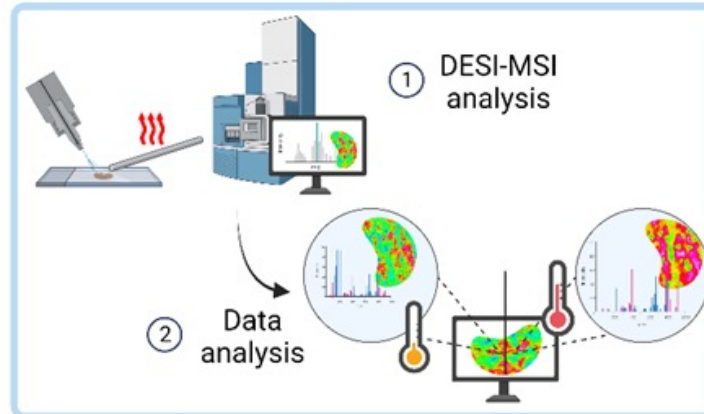


Vast amounts of data from a single tissue section

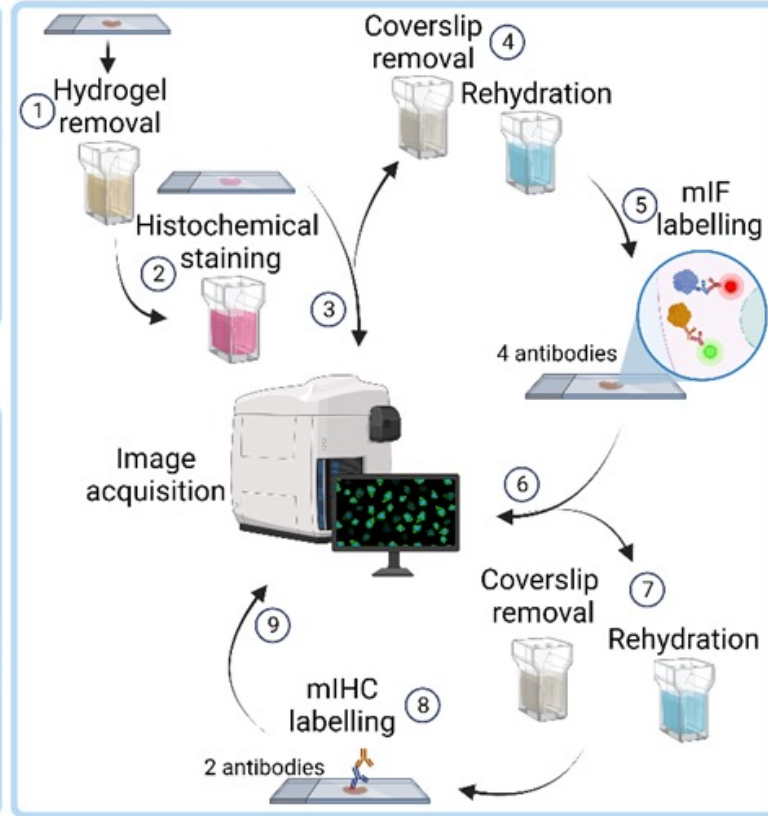
① Tissue embedding and sectioning



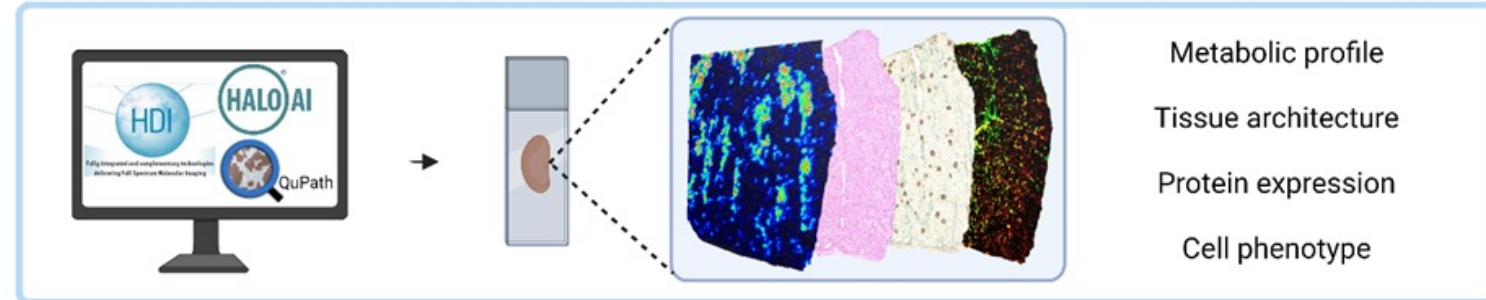
② DESI-MSI analysis



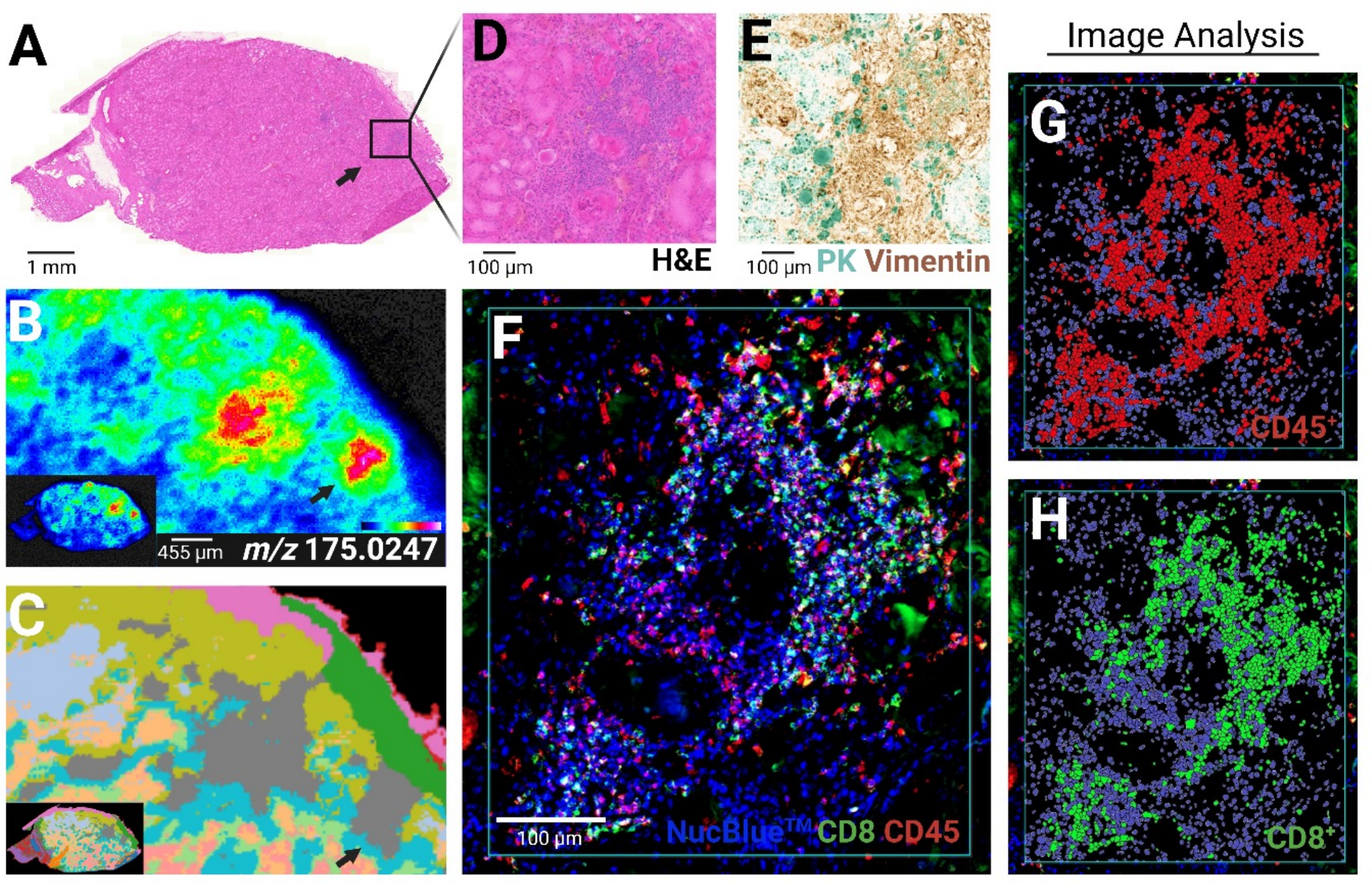
③ Histology and protein labelling



④ Data integration, analysis and interpretation



Application of the workflow – scarring and immune cell infiltration

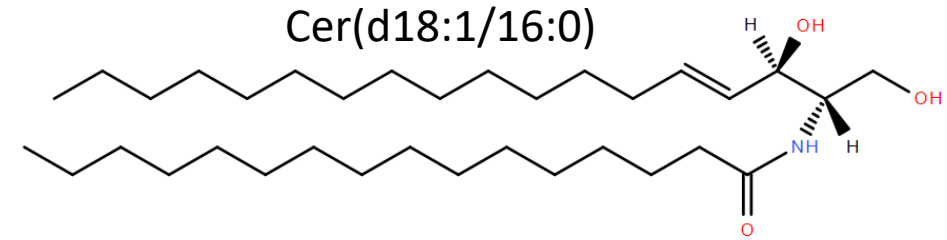
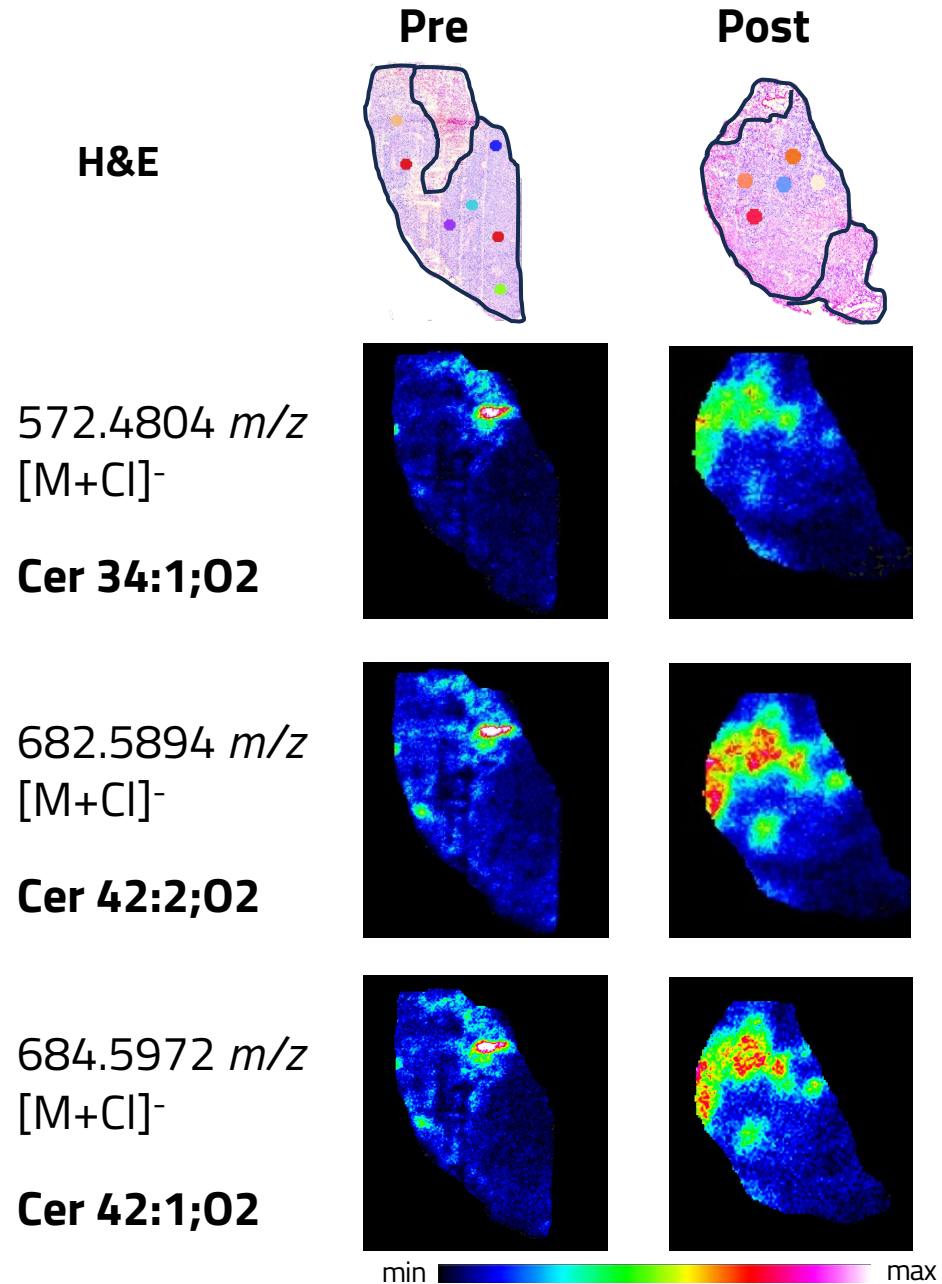


Translational Studies – biomarker discovery

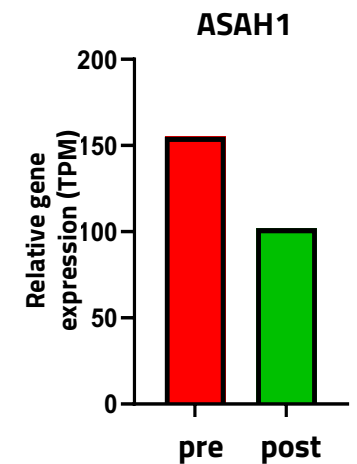
- **Translational medicine** “from bench to bedside” - requires a multidisciplinary approach
- Benefits of a well-designed clinical study allow **Paired** pre and on-treatment biopsies from patients in on-going Phase I/II clinical studies
- Challenges in getting tissue from patients with cancer
- Crucial that we obtain as much data as possible from a small sample



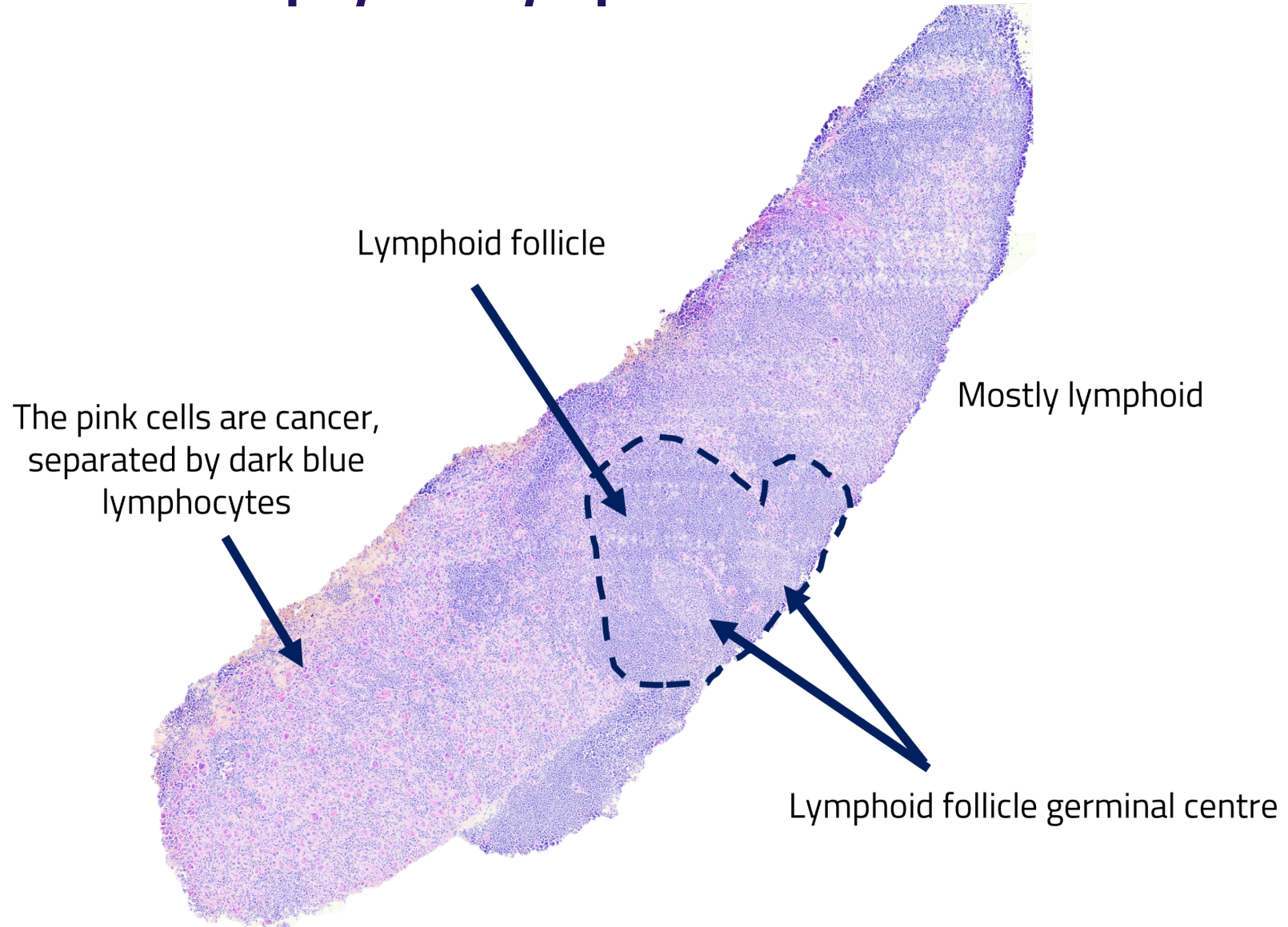
Case 1: necrotic tissue and dense areas of cancer cells



- 3 Ceramide (Cer) species significantly increased ($|d| > 5$) following treatment
- Bulk RNAseq of same biopsy shows a reduction in the expression of *ASAH1*, the gene that encodes the enzyme acid ceramidase (AC)
- AC is involved in metabolism of ceramides and often overexpressed in certain cancers
- Ceramides are pro-apoptotic

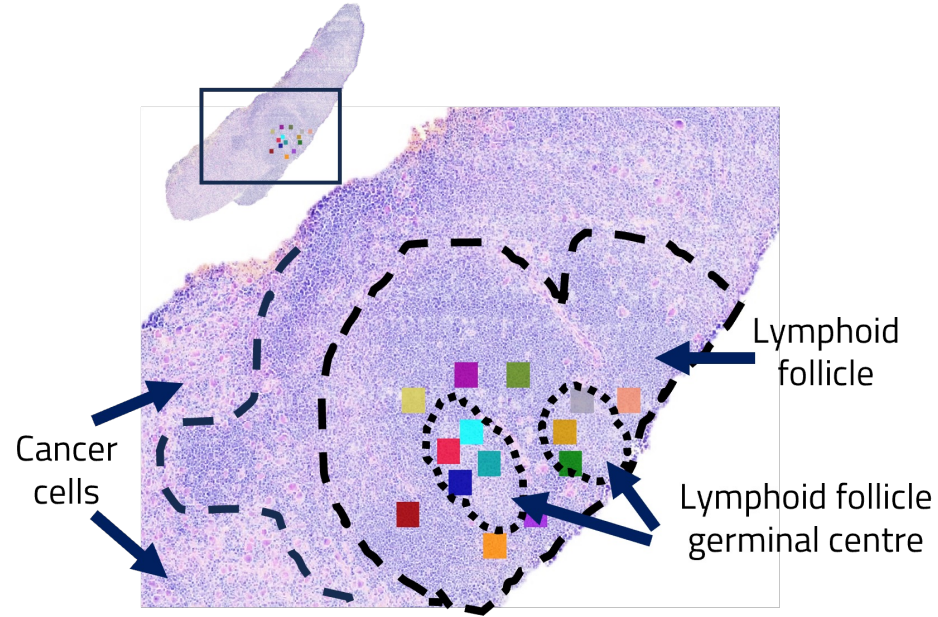


Case 2: Pre-Treatment biopsy of a lymph node

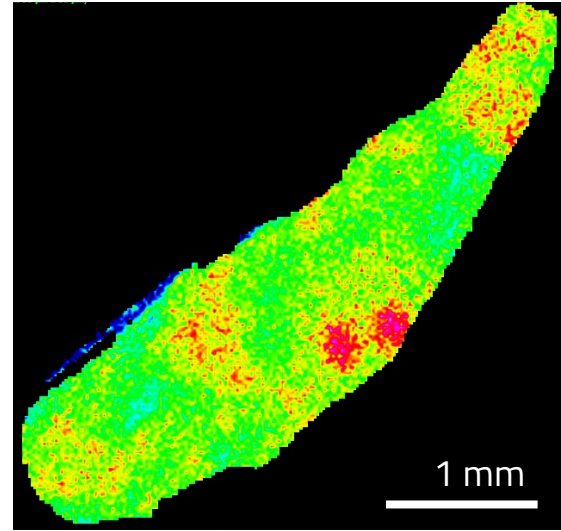
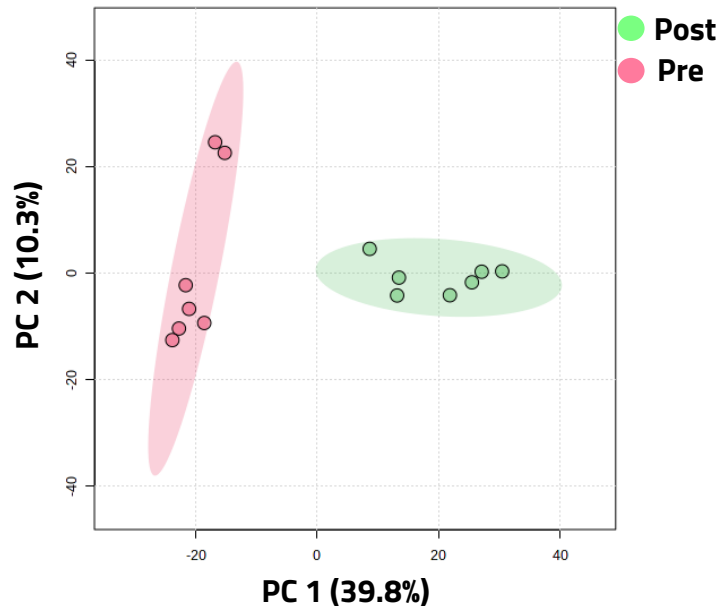


- Germinal centres, contain few T cells as these are full of maturing B cells

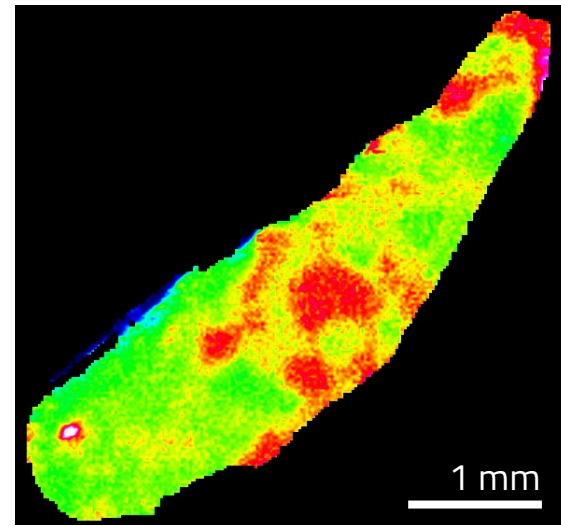
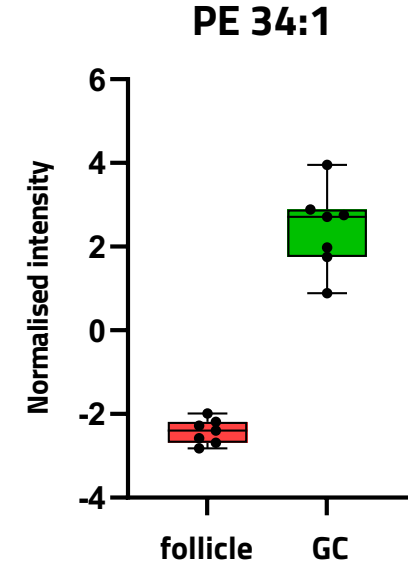
Metabolic changes revealed by MSI show an immune response in action



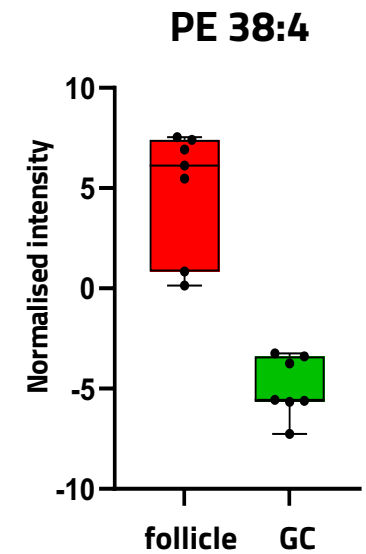
PCA 2D Scores Plot



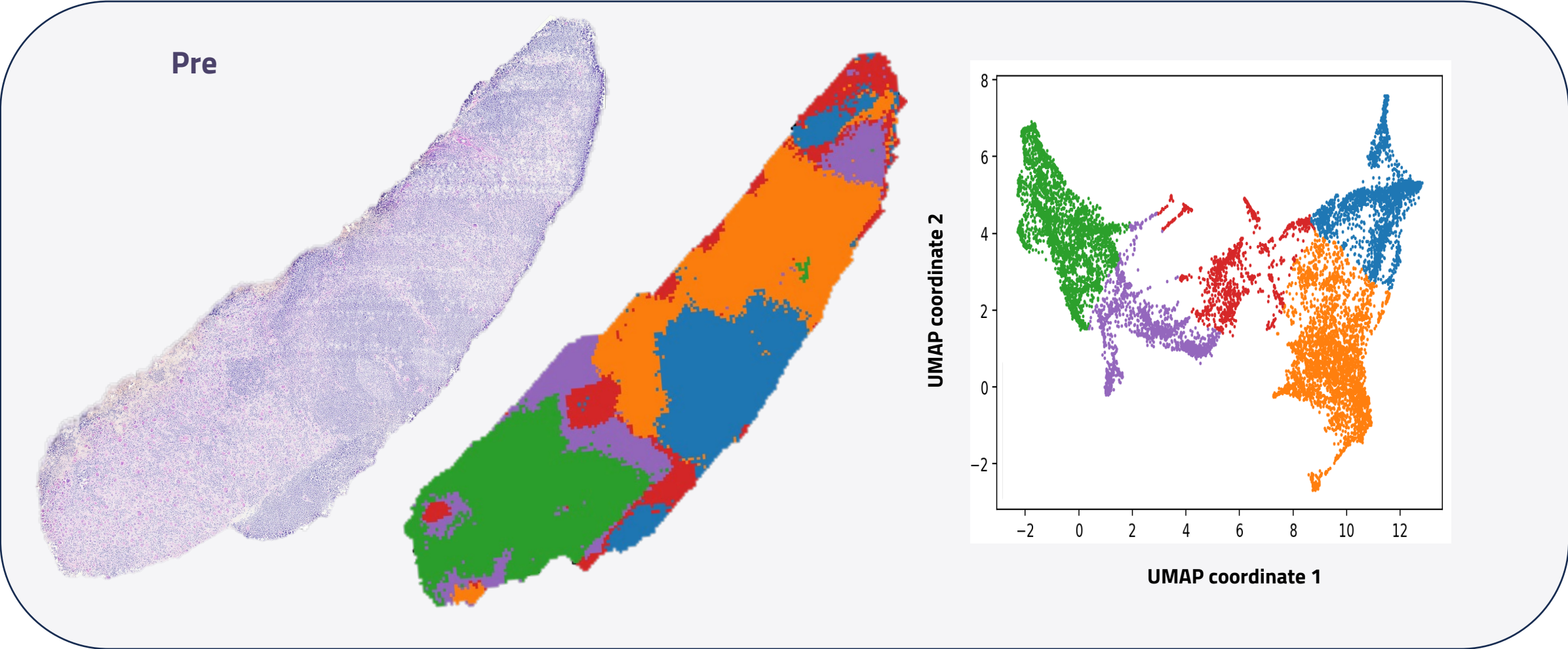
min max



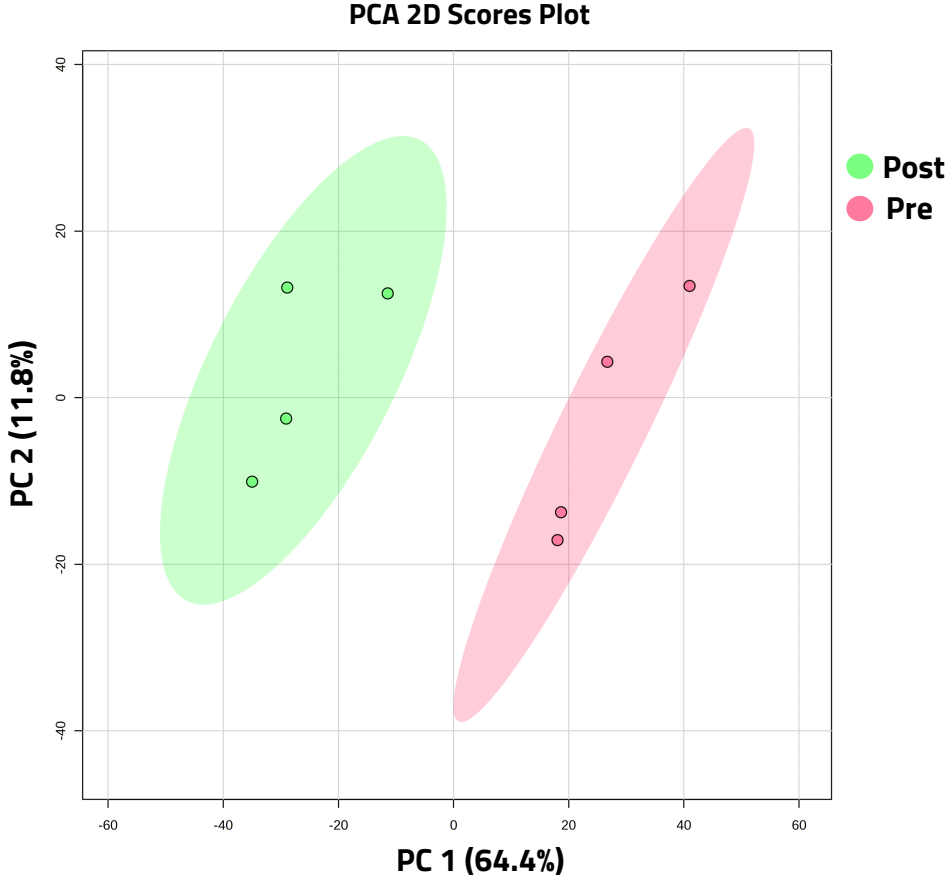
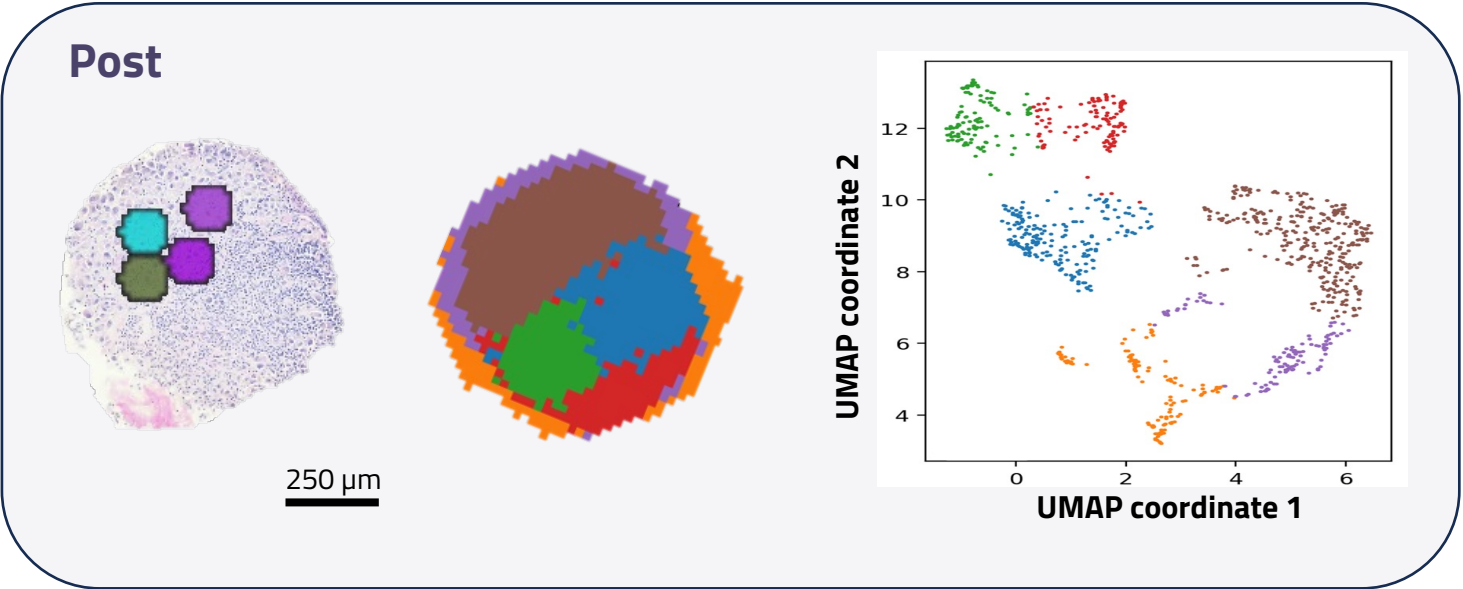
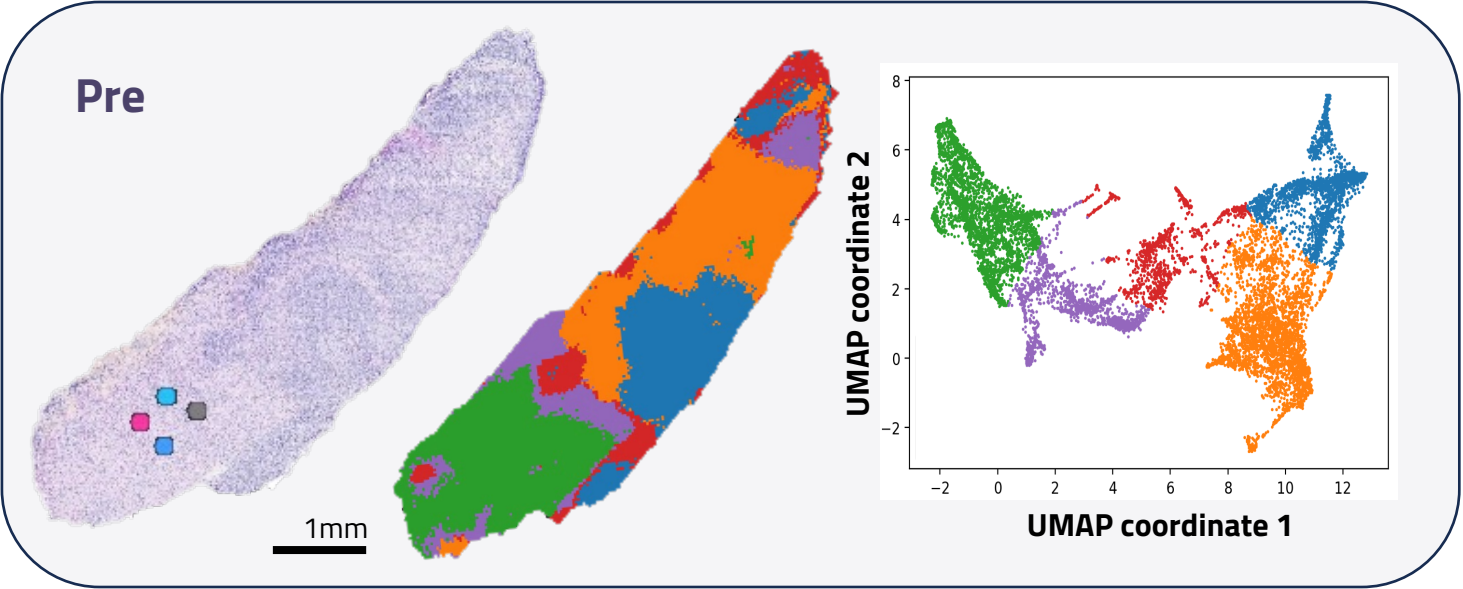
min max



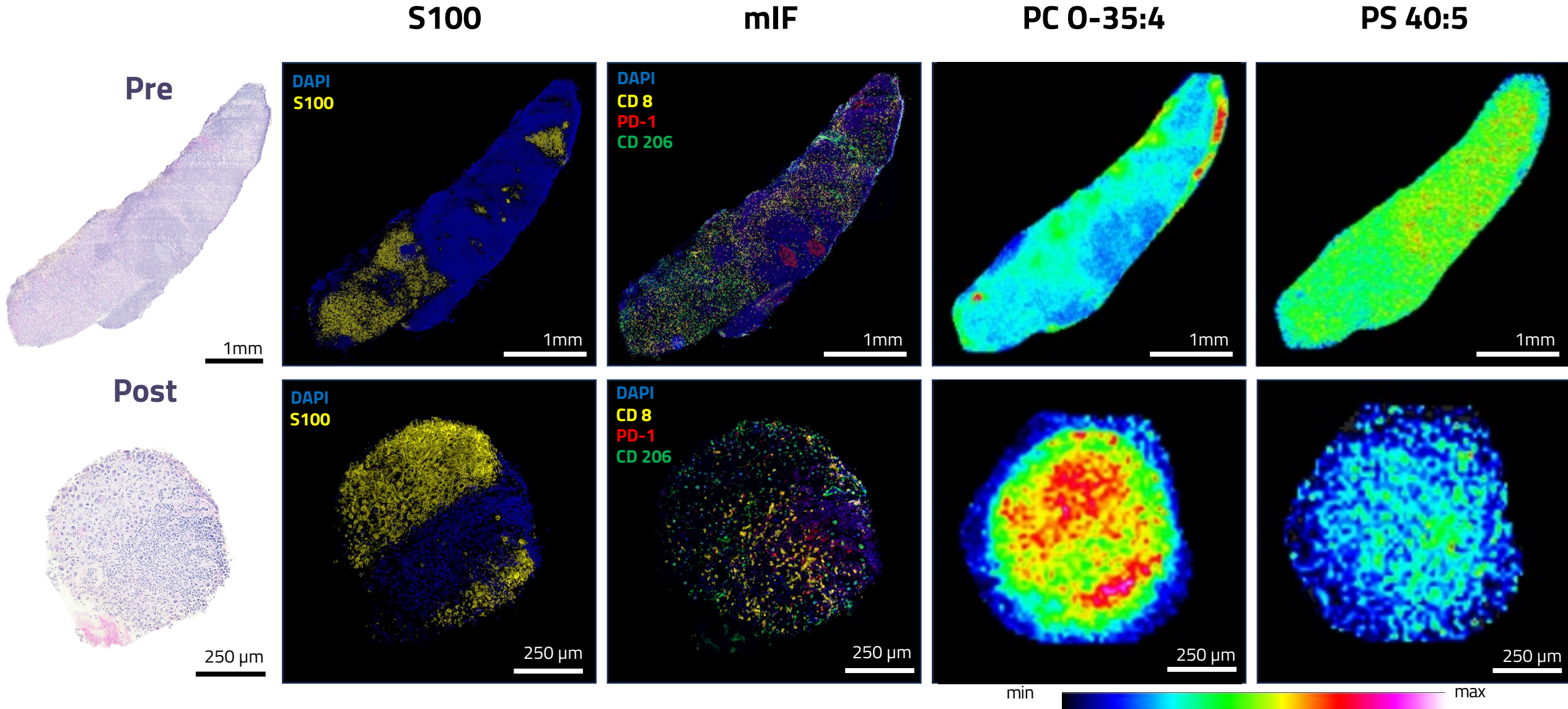
Pre and Post treatment biopsies – heterogeneity



Pre and Post treatment biopsies – heterogeneity

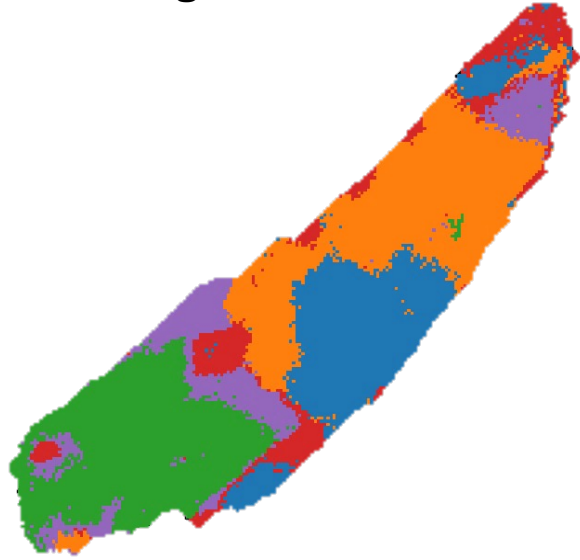


Pre and Post treatment biopsies – metabolome and protein changes



MSI data integrated into AI enabled image analysis software

UMAP Clustering of MSI data



Metabolite annotation layer

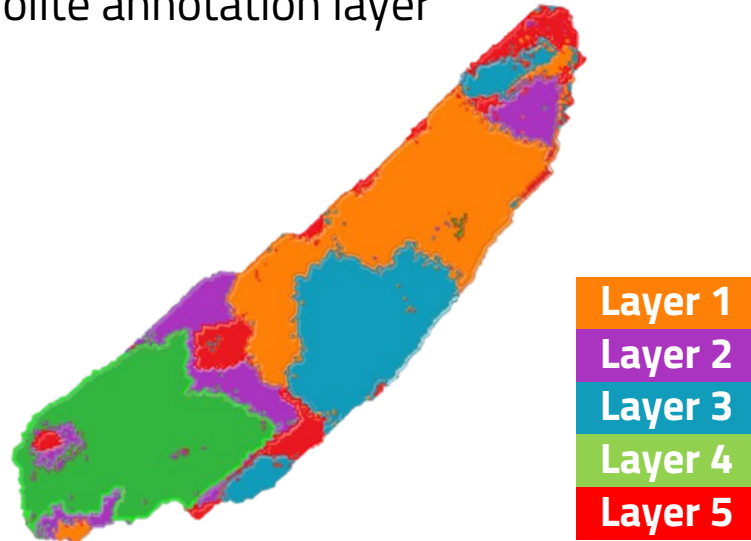
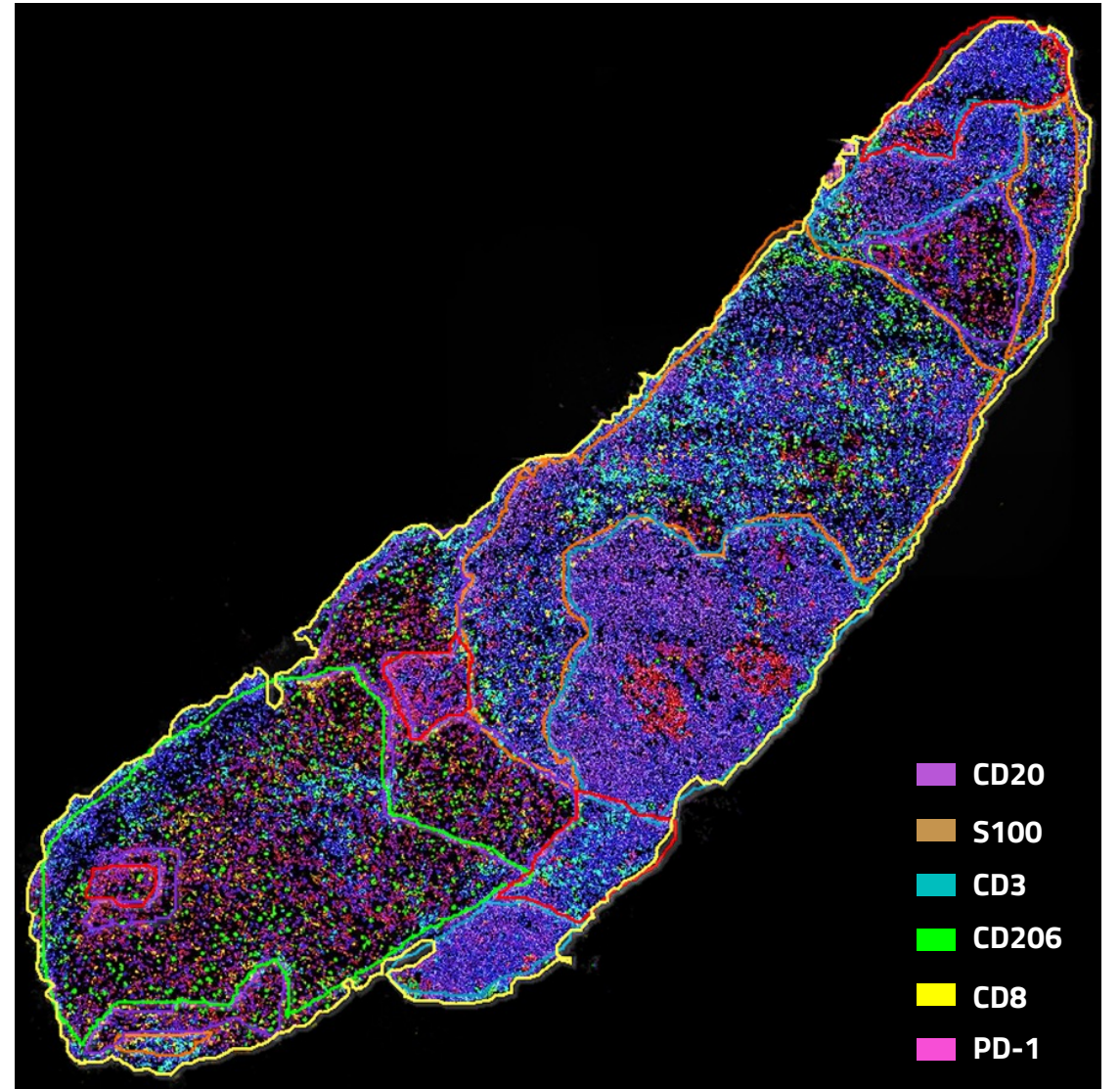
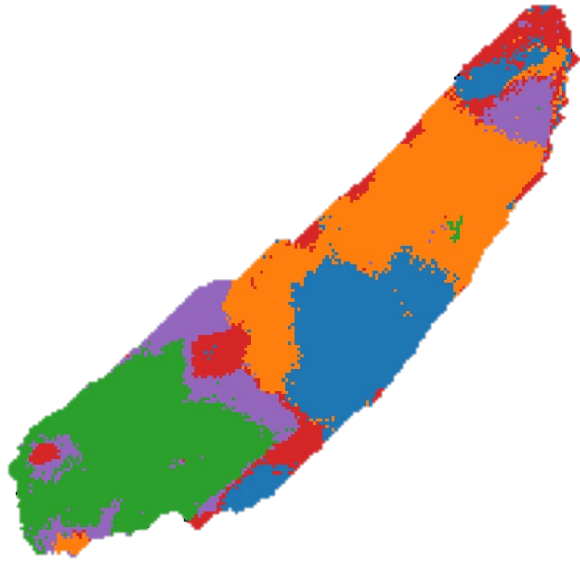


Image Analysis



MSI data integrated into AI enabled image analysis software

UMAP Clustering of MSI data



Metabolite annotation layer

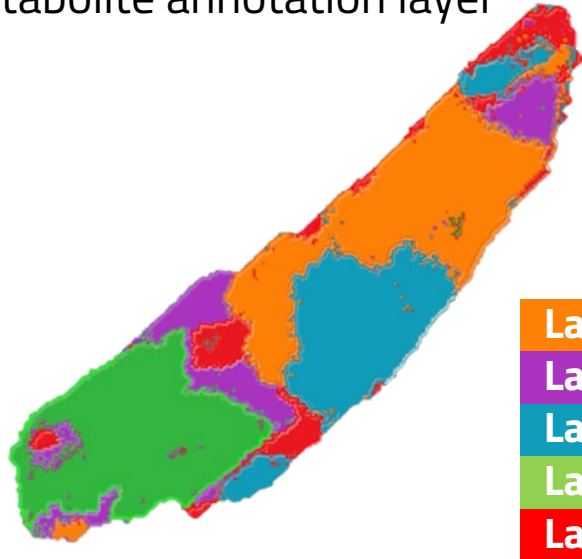
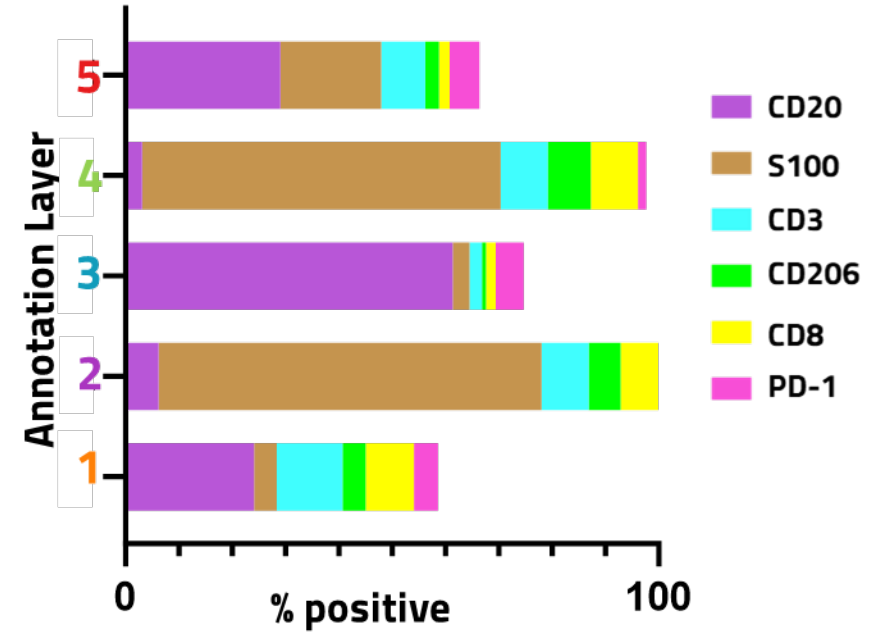
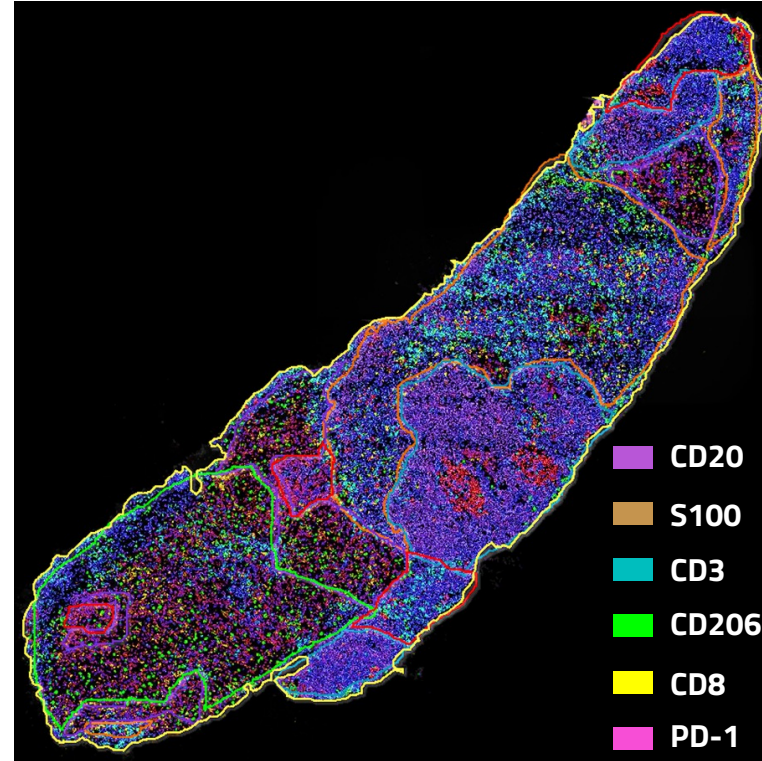
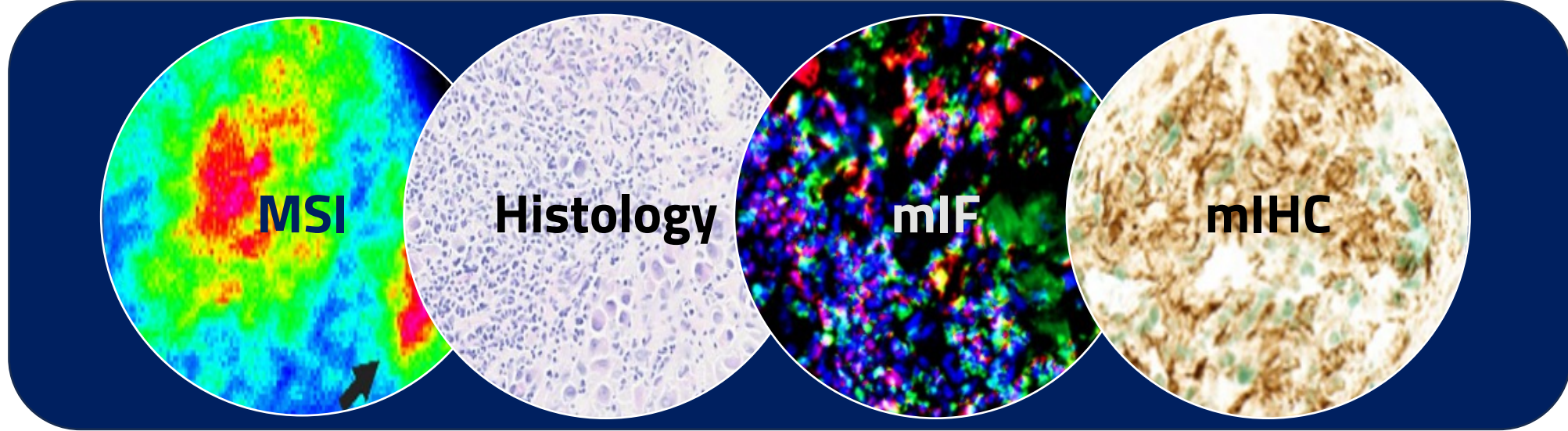


Image Analysis



Summary: Using a single 10 μm tissue section...



- DESI MSI – successfully incorporated into image analysis workflows using a single tissue section
- AI enabled pipeline that incorporates histology, mIHC, mIF following DESI-MSI
- Well-designed clinical studies provide opportunities to investigate a patient cancer and PD response to therapy
- Metabolic changes from MSI coupled to histology reveal immune responses
- Metabolomic changes are occurring between regions of interest of high cancer cell density pre and post treatment
- Metabolic signatures can be integrated into AI analysis software for molecular phenotyping

Collaborators/Network

University of St Andrews Imaging and AI Team

Greice Zickuhr
In Hwa Um
David Harrison

NuCana plc Translational Team

Mustafa Elshani
Ying Zhang

NuCana plc Clinical Collaborators

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Aglaia Skolariki
Noor Md Haris
Zhuang Boh
Sarah P Blagden
Natalie Cook
TR Jeffry Evans
E. Ruth Plummer

The logo for NuCana, featuring the word "NUCANA" in a bold, dark blue, sans-serif font. A stylized, curved blue line arches over the letters "A" and "N".