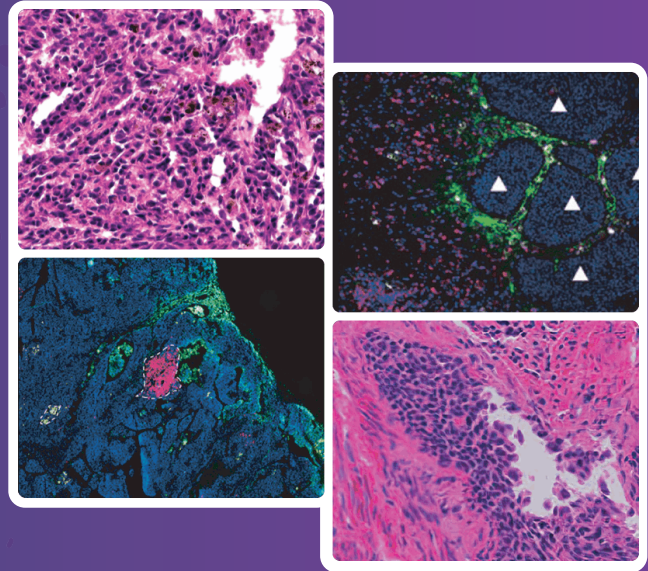


# STOmics Complete GENOMICS™

Tumor cell heterogeneity and microenvironment are key to research on tumor mechanism, prevention, and treatment. Spatial transcriptome technology -- Stereo-seq, can map tumor cell heterogeneity and microenvironment structure at subcellular resolution, bringing significant breakthroughs in tumor research. Similarly, spatial multi-omics technology can also be applied to other diseases that are tied to spatial structures, providing a powerful tool for further exploration.



## Technology highlights

- True single-cell resolution allowing visualization on cellular heterogeneities and efficient cell type annotations

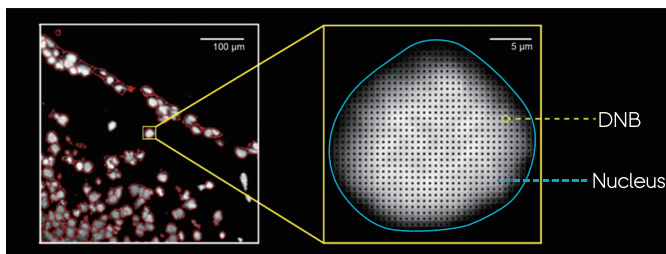


Figure 1. Stereo-seq capturing spots (DNBs) arranged down to single cell resolution, allowing mRNA within one cell to be captured with hundreds of coordinate ID containing DNBs

- Multiple chip sizes allowing various tissue types and scalable capture areas

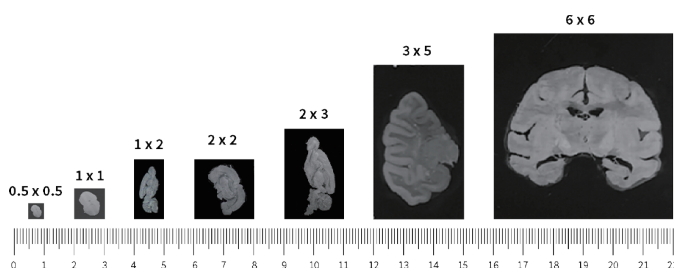
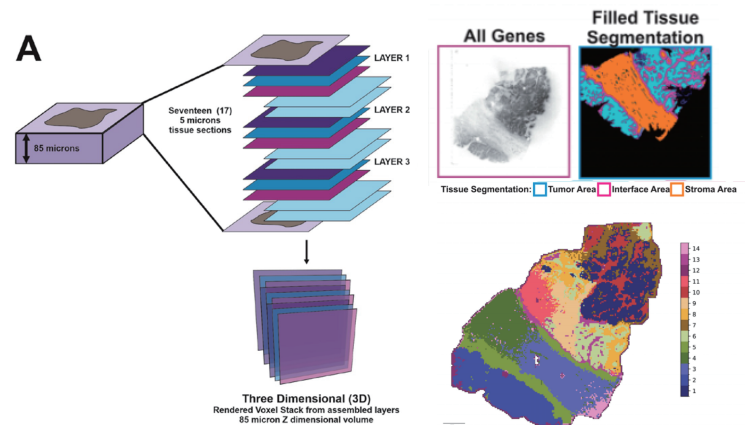


Figure 2. Demonstration of Stereo-seq chip at different sizes

## Featured publication

Molecular, Metabolic, and Subcellular Mapping of the Tumor Immune Microenvironment via 3D Targeted and Non-Targeted Multiplex Multi-Omics Analyses

Ferri-Borgogno & Burks et al., *Cancers*, 2024



### Highlights

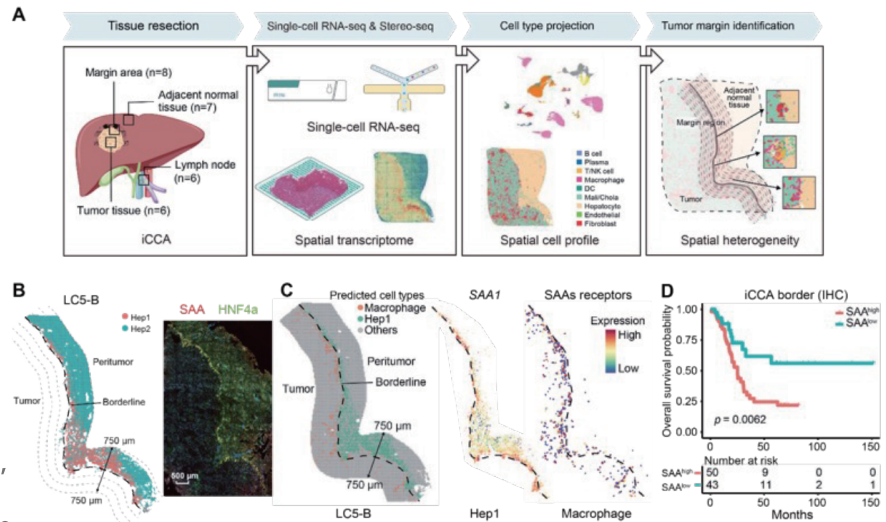
- This approach aims to unveil intricate signaling networks between different cell types and the extracellular matrix (ECM) in the 3D tumor microenvironment (TIME) using two FFPE gynecologic tumor samples (HGSOE & AEH)
- Stereo-seq for FFPE enabled the capture of not only human coding and non-coding RNA but also microbiome RNAs with subcellular resolution.

STOmics product:  
Early Access Stereo-seq Transcriptomics Solution for FFPE

## Featured publication

An invasive zone in human liver cancer identified by Stereo-seq promotes hepatocyte-tumor cell crosstalk, local immunosuppression and tumor progression.

Wu et al., *Cell Research*, 2023



### Highlights

- Spatial transcriptomics reveals the heterogeneity in tumor margin area.
- Characterization of distinctive immune cells, the suppressive immune microenvironment, and metabolic reprogramming of tumor cells in invasive zone.
- In the invasive zone, hepatocyte injury upregulates SAAs, further recruiting and polarizing macrophages to assist tumor progression.

STOmics product: Stereo-seq Transcriptomics Solution

### Publications

Molecular, Metabolic, and Subcellular Mapping of the Tumor Immune Microenvironment via 3D Targeted and Non-Targeted Multiplex Multi-Omics Analyses  
*Cancers* 2024, 16(5), 846

### Experiment snapshot

STOmics product: Early Access Stereo-seq Transcriptomics Solution for FFPE  
Sample type: Human, FFPE

### Link



A cellular hierarchy in melanoma uncouples growth and metastasis.  
*Nature* 610: 190–198 (2022)

STOmics product: Stereo-seq Transcriptomics Solution  
Sample type: Mouse, Fresh frozen



Presence of onco-fetal neighborhoods in hepatocellular carcinoma is associated with relapse and response to immunotherapy  
*Nature Cancer* 5:167–186 (2024)

STOmics product: Stereo-seq Transcriptomics Solution  
Sample type: Human, Fresh frozen



Single-cell landscape of idiopathic Multicentric Castleman Disease in identical twins  
*Blood* 2023021992.

STOmics product: Stereo-seq Transcriptomics Solution  
Sample type: Human, Fresh frozen



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