Explore spatial protein and subcellular gene expression profile with Stereo-seq for advancement in fundamental and translational research

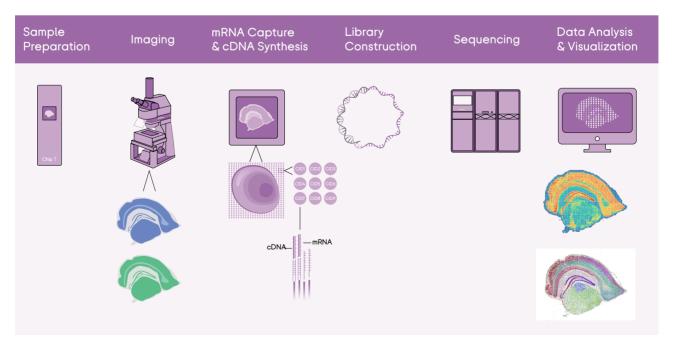
01 Stereo-seq Transcriptome and Multiplex Immunofluorescence (mIF) Co-detection

STOmics Stereo-seq Transcriptomics Set for Chip-on-a-slide is intended for generating a spatiallyresolved 3' mRNA library from biological tissue sections. Built upon DNA Nanoball (DNB) technology, STOmics Stereo-seq Transcriptomics Set for Chip-on-a-slide enables a "tissue-to-data" solution through in situ capture of the whole transcriptome, at nanoscale resolution and centimeter-sized field of view. Stereo-seq Transcriptomics Solution utilizes DNB patterned array chips loaded with spatiallybarcoded probes that capture and prime poly-adenylated mRNA from tissue sections in situ. Each cDNA synthesized from mRNA captured on a particular spot is linked to its spatially-barcoded probe, allowing subsequent gene expression mapping of a tissue section following sequencing and visualization analysis using the StereoMap visualization platform.

By integrating mIF staining method into the process of Stereo-seq Transriptomics standard workflow, Stereo-seq transcriptome and multiplex immunofluorescence (mIF) co-detection technology enables spatial visualization of multiple proteins on top of the unbiased whole transcriptome information on the same tissue slice. Without affecting mRNA capturing, the additional detected protein information can be integrated with gene expression data to in-depth evaluate valuable samples, and to parse complex pathological and physiological processes. The amount of protein that can be detected depends on user's antibody selection and imaging configuration. In this user manual, we are showing DAPI with stainings of three antibodies as an example.

02 Application Highlights

- Study protein, unbiased high-resolution whole transcriptome information and tissue morphology on the same tissue section.
- Discover new tissue biomarkers and molecular targets at a spatial scale with cellular specificity.
- Interrogate gene expression profiles and protein markers in different tumor regions to further explore tumor progressing mechanisms.
- Characterize immune cells and immune repertoires within healthy tissue microenvironment vs TME.
- · Identify the spatial location of secreted proteins within the cell under different conditions.



03 Gene Expression & Protein Cluster Demo Display

Mouse Testis

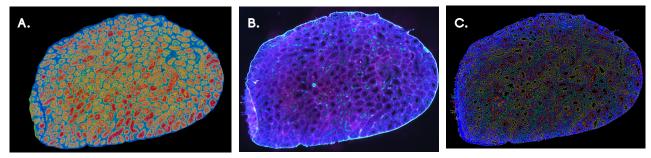
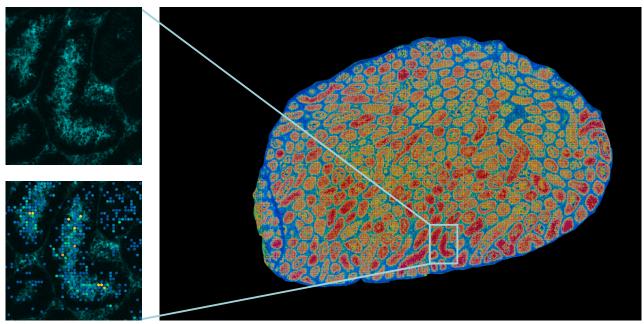


Figure 1. A. Spatial gene expression heatmap of mouse testis at bin20 (~10 µm). B. Merged immunofluorescence images of mouse testis stained for DAPI (nuclei staining), AKAP3 protein and TESK2 protein by immunofluorescence. C. Mouse testis single-cell clustering results using cellbinning and overlayed with merged immunofluorescence images.



Bin 20

Bin 20

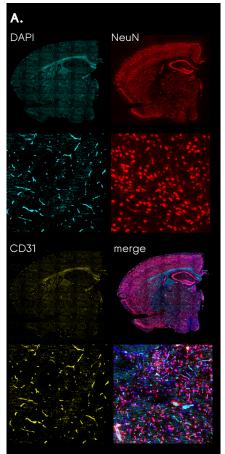
Figure 2. A mouse testis section was stained with anti-AKAP3 antibody and anti-TESK2 antibody and went through Stereoseq Transcriptomics mIF workflow. Here, demonstrating a zoomed in region within the tissue with AKAP3 staining and colocalization of AKAP3 gene.

AKAP3 only AKAP3 protein + AKAP3 gene

STOmics

Mouse Brain

Mouse Spleen



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Figure 3. A. Individual fluorescence images of DAPI stained nuclei, anti-NeuN antibody staining and anti-CD31 antibody staining of mouse brain and merged immuno-fluorescence images. B. Spatial gene expression heatmap of mouse brain at bin20 resolution (~10 μ m) after immuno-fluorescence assay in the same section.

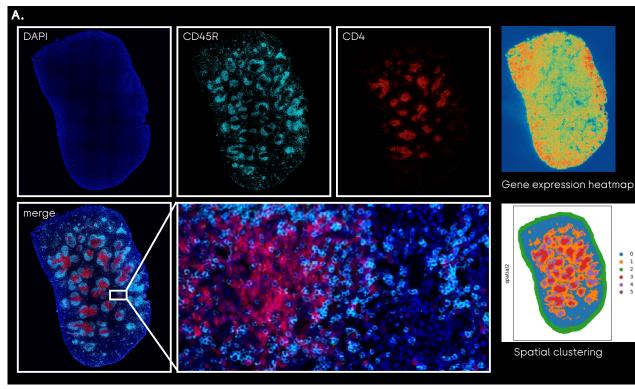


Figure 4. A. Individual fluorescence images of DAPI stained nuclei, anti-CD45R antibody staining and anti-CD4 antibody staining of mouse spleen and merged immuno-fluorescence images. B. Spatial gene expression heatmap of mouse spleen at bin50 resolution (~25 µm) after immuno-fluorescence assay in the same section. C. Spatial clustering of unbiased transcriptome with Leiden algorithm using Scanpy.

04 Product Reagents

Perform Stereo-seq in your own laboratory with Stereo-seq Kits.

Stereo-seq Permed	ibilization Set for Chip-on-a-sli	ide		
Part Number	Product		Specification	Description
211SP118	Stereo-seq Permeabilization Kit	111KP118	8 RXN	For determining permeabilization parameters to optimize mRNA capture
	Stereo-seq Chip P Slide (1cm*1cm)	210CP118	8 EA	
	STOmics Accessory Kit	1000033700	5 PCS	
Stereo-seq Transci	iptomics Set for Chip-on-a-slid	e		
Part Number	Product		Specification	Description
211ST114	Stereo-seq Transcriptomics T Kit	111KT114	4 RXN	For generating a spatially-resolved 3' mRNA library from biological tissue sections
	Stereo-seq Chip T Slide (1cm*1cm)	210CT114	4 EA	
	STOmics Accessory Kit	1000033700	5 PCS	
Stereo-seq Library	Preparation Kit			
Part Number	Product		Specification	Description
111KL114	Stereo-seq Library Preparation Kit		4 RXN	For constructing STOmics Library
Stereo-seq PCR Ac	laptor			
Part Number	Product		Specification	Description
301AUX001	Stereo-seq PCR Adaptor		2 EA	Compatible with PCR thermal cycler as a heating unit

05 Key Features

- High fidelity protein staining imaging for enhanced spatial protein identification.
- O Chip-on-a-slide for efficient experiment operation and compatible with existing Stereo-seq solutions.
- O Detect multiple proteins with unbiased whole transcriptome simultaneously in fresh frozen and PFA fixed samples.
- I Flexibility in choosing antibodies based on user's research needs and project design.
- 6 Co-detection of protein and RNA in a single experiment for extensible applications in tissue and cellular type study.

06 Intuitive Software Pipelines and Suits

ImageStudio image processing software, **SAW** (Stereo-seq Analysis Workflow) and **StereoMap** visualizaiton platform are offered free of charge to users and enables them to discover spatial biology knowledge with multiplexed tissue images.



Learn more: https://en.stomics.tech/BioinfoTools