# Resource Summary Report

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## Mayo Clinic Spatial Biology Core Facility

RRID:SCR_026548
Type: Tool

### Proper Citation

Mayo Clinic Spatial Biology Core Facility (RRID:SCR_026548)

### Resource Information

**URL:** https://www.mayo.edu/research/core-resources/spatial-biology-core/overview

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**Description:** Core provides broad range of spatial transcriptomic and targeted proteomics services including: Conventional nCounter analysis to determine RNA abundance, GeoMx Digital Spatial Profiling platform to determine protein and RNA abundance, CosMx Spatial Molecular Imaging platform for single cell spatial analysis of RNA and protein.

**Resource Type:** access service resource, core facility, service resource

**Keywords:** spatial transcriptomic, targeted proteomics, determine RNA abundance, determine protein abundance, single cell spatial analysis of RNA and protein

**Availability:** Open

**Resource Name:** Mayo Clinic Spatial Biology Core Facility

**Resource ID:** SCR_026548

**Record Creation Time:** 20250308T060322+0000

**Record Last Update:** 20260919T200052+0000

### Ratings and Alerts

No rating or validation information has been found for Mayo Clinic Spatial Biology Core Facility.

No alerts have been found for Mayo Clinic Spatial Biology Core Facility.

### Data and Source Information

**Source:** SciCrunch Registry

## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at nidm-terms .

Liu Y, et al. (2026) Spatially resolved single-cell landscape of tumor immunotypes reveals the central role of interferon signaling and plasmacytoid dendritic cells in triple-negative breast cancer. Journal for immunotherapy of cancer, 14(6).