

# Resource Summary Report

Generated by ASWG on Aug 4, 2026

## Harvard Medical School Single Cell Core Facility

RRID:SCR\_025837

Type: Tool

### Proper Citation

Harvard Medical School Single Cell Core Facility (RRID:SCR\_025837)

### Resource Information

**URL:** <https://singlecellcore.hms.harvard.edu>

**Proper Citation:** Harvard Medical School Single Cell Core Facility (RRID:SCR\_025837)

**Description:** Core specializes in single cell and spatial transcriptomics, and multiomics. Provides custom services, education, training and expertise in multiple high-throughput single cell technologies.

**Abbreviations:** SCC

**Synonyms:** Harvard Medical School Single Cell Core, Single Cell Core

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

**Keywords:** single cell, spatial transcriptomics, multiomics, single cell technologies,

**Availability:** Open

**Resource Name:** Harvard Medical School Single Cell Core Facility

**Resource ID:** SCR\_025837

**Alternate IDs:** ABRF\_2939

**Alternate URLs:** <https://coremarketplace.org/?FacilityID=2939&citation=1>

**Record Creation Time:** 20241003T053303+0000

**Record Last Update:** 20260804T043851+0000

### Ratings and Alerts

No rating or validation information has been found for Harvard Medical School Single Cell Core Facility.

No alerts have been found for Harvard Medical School Single Cell Core Facility.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 4 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [ASWG](#).

Yeo AT, et al. (2022) Single-cell RNA sequencing reveals evolution of immune landscape during glioblastoma progression. Nature immunology, 23(6), 971.

Avagyan S, et al. (2021) Single-cell ATAC-seq reveals GATA2-dependent priming defect in myeloid and a maturation bottleneck in lymphoid lineages. Blood advances, 5(13), 2673.

Tattikota SG, et al. (2020) A single-cell survey of Drosophila blood. eLife, 9.

Southard-Smith AN, et al. (2020) Dual indexed library design enables compatibility of in-Drop single-cell RNA-sequencing with exAMP chemistry sequencing platforms. BMC genomics, 21(1), 456.