

Resource Summary Report

Generated by T1D on Sep 5, 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: access service resource, core facility, service resource

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: 20260904T001019+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.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Weinberg MS, et al. (2026) Bacillus Calmette-Guérin (BCG) immunotherapy reprograms CNS immunity and alters Alzheimer's biomarkers: results from two open-label clinical trials. Communications medicine, 6(1).

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.

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.

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