Resource Summary Report

Generated by RRID on Aug 5, 2026

Stanford University Shared FACS Core Facility

RRID:SCR_017788
Type: Tool

Proper Citation

Stanford University Shared FACS Core Facility (RRID:SCR_017788)

Resource Information

URL: http://facs.stanford.edu/

Proper Citation: Stanford University Shared FACS Core Facility (RRID:SCR_017788)

Description: Provides flow cytometry instrumentation and expertise. Provides operator assisted analyzer and sorter use, as well as training and support for user instrument operation.

Synonyms: Stanford Shared FACS Facility

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

Keywords: Flow, cytometry, instrumentation, analysis, cell, sorting, training, service, core

Funding: NCRR S10 RR025518;
NCRR S10 RR027431;
NIH Office of the Director S10 OD016318;
Parker Institute for Cancer Immunotherapy

Availability: Open

Resource Name: Stanford University Shared FACS Core Facility

Resource ID: SCR_017788

Alternate IDs: ABRF_398

Record Creation Time: 20220129T080337+0000

Record Last Update: 20260805T044412+0000

Ratings and Alerts

No rating or validation information has been found for Stanford University Shared FACS Core Facility.

No alerts have been found for Stanford University Shared FACS Core Facility.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 16 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Jiang H, et al. (2026) IL-9 as a naturally orthogonal cytokine with optimal JAK/STAT signaling for engineered T cell therapy. Immunity, 59(1), 177.

Schwartz CI, et al. (2025) Toward optimizing diversifying base editors for high-throughput mutational scanning studies. Nucleic acids research, 53(12).

Chen D, et al. (2025) Connectome-seq: High-throughput Mapping of Neuronal Connectivity at Single-Synapse Resolution via Barcode Sequencing. bioRxiv : the preprint server for biology.

Jiang H, et al. (2025) IL-9 as a naturally orthogonal cytokine with optimal JAK/STAT signaling for engineered T cell therapy. bioRxiv : the preprint server for biology.

Butler DSC, et al. (2025) Eosinophils enhance granuloma-mediated control of persistent Salmonella infection in mice. Nature microbiology, 10(12), 3176.

Lee S, et al. (2025) Pharmacologic stimulation of insulin granule acidification increases ?-cell zinc content and augments ?-cell-targeted drug delivery. The Journal of biological chemistry, 301(10), 110645.

Raposo CJ, et al. (2025) Functional memory T cells are derived from exhausted clones and expanded by checkpoint blockade. bioRxiv : the preprint server for biology.

Zhao H, et al. (2025) Elucidating the role of N-myristoylation in the excessive membrane localization of PD-L1 in hypoxic cancers and developing a novel NMT1 inhibitor for combination with immune checkpoint blockade therapy. Journal of experimental & clinical cancer research : CR, 44(1), 181.

Monack D, et al. (2025) Eosinophils Enhance Granuloma-Mediated Control of Persistent Salmonella Infection. Research square.

Yamamoto Y, et al. (2025) Scaled multidimensional assays of variant effect identify sequence-function relationships in hypertrophic cardiomyopathy. bioRxiv : the preprint server for biology.

Wang T, et al. (2025) Sensitive, direct detection of non-coding off-target base editor unwinding and editing in primary cells. bioRxiv : the preprint server for biology.

Naktang C, et al. (2025) Chromosome-level assemblies of two hexaploid bamboos, Thyrsostachys oliveri and Thyrsostachys siamensis, provide a foundation for functional and comparative genomics studies. GigaScience, 14.

Yang Y, et al. (2025) A spatiotemporal and machine-learning platform facilitates the manufacturing of hPSC-derived esophageal mucosa. Developmental cell, 60(9), 1359.

Park JH, et al. (2024) p53 engagement is a hallmark of an unfolded protein response in the nucleus of mammalian cells. bioRxiv : the preprint server for biology.

Li NY, et al. (2024) Basal-to-inflammatory transition and tumor resistance via crosstalk with a pro-inflammatory stromal niche. Nature communications, 15(1), 8134.

Yang Y, et al. (2023) A Spatiotemporal and Machine-Learning Platform Accelerates the Manufacturing of hPSC-derived Esophageal Mucosa. bioRxiv : the preprint server for biology.