Resource Summary Report

Generated by RRID on Sep 10, 2026

Rockefeller University Flow Cytometry Resource Center Core Facility

RRID:SCR_017694
Type: Tool

Proper Citation

Rockefeller University Flow Cytometry Resource Center Core Facility (RRID:SCR_017694)

Resource Information

URL: https://www.rockefeller.edu/fcrc/

Proper Citation: Rockefeller University Flow Cytometry Resource Center Core Facility (RRID:SCR_017694)

Description: Core offers researchers multi laser and multi color flow cytometry sorters and analyzers, including spectral and imaging flow cytometers.

Abbreviations: FCRC

Synonyms: Flow Cytometry Resource Center

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

Keywords: Flow, cytometry, sorter, analyzer, imaging, core, Rockefeller, service

Availability: Restricted

Resource Name: Rockefeller University Flow Cytometry Resource Center Core Facility

Resource ID: SCR_017694

Alternate IDs: SCR_017710, ABRF_102

Alternate URLs: https://coremarketplace.org/?FacilityID=102&citation=1

Record Creation Time: 20220129T080336+0000

Record Last Update: 20260905T133410+0000

Ratings and Alerts

No rating or validation information has been found for Rockefeller University Flow Cytometry Resource Center Core Facility.

No alerts have been found for Rockefeller University Flow Cytometry Resource Center Core Facility.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 25 mentions in open access literature.

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

Yatim A, et al. (2026) Human LFA-1 governs T cell immune surveillance of the skin. Science immunology, 11(116), eadz8360.

Liu S, et al. (2026) Radial-glia-to-astrocyte trans-differentiation and astrocyte transcriptional convergence are coordinated by CEH-43/DLX in C. elegans. bioRxiv : the preprint server for biology.

Patriotis AL, et al. (2026) The conserved N-terminal SANT1-binding domain (SBD) of EZH2 regulates PRC2 activity. Genes & development, 40(9-10), 650.

Kojima H, et al. (2026) A post-translational regulatory map of chronic antigen-driven human T cell dysfunction. bioRxiv : the preprint server for biology.

Kim C, et al. (2026) A flow cytometry-based assay to quantify the binding of transmembrane ligands to their cognate receptors using fluorescent virus-like particles. bioRxiv : the preprint server for biology.

Varandas KC, et al. (2025) Glia detect and transiently protect against dendrite substructure disruption in C. elegans. Nature communications, 16(1), 79.

Saidoune F, et al. (2025) Enhanced TLR7-dependent production of type I interferon by pDCs underlies pandemic chilblains. The Journal of experimental medicine, 222(7).

Chapa CM, et al. (2025) RNA-Seq dataset of wild type and PRC2 mutant mouse embryonic stem cells undergoing ground state conversion. Data in brief, 62, 111957.

Zhang Z, et al. (2024) A Panoramic View of Cell Population Dynamics in Mammalian Aging. bioRxiv : the preprint server for biology.

Fang Y, et al. (2024) ATRX guards against aberrant differentiation in mesenchymal progenitor cells. Nucleic acids research, 52(9), 4950.

Arias AA, et al. (2024) Tuberculosis in otherwise healthy adults with inherited TNF deficiency. Nature, 633(8029), 417.

Xu Z, et al. (2024) Dissecting key regulators of transcriptome kinetics through scalable single-cell RNA profiling of pooled CRISPR screens. Nature biotechnology, 42(8), 1218.

Lee J, et al. (2024) Principles of peptide selection by the transporter associated with antigen processing. Proceedings of the National Academy of Sciences of the United States of America, 121(23), e2320879121.

Stewart KS, et al. (2024) Stem cells tightly regulate dead cell clearance to maintain tissue fitness. Nature, 633(8029), 407.

Khan A, et al. (2024) Metabolic gene function discovery platform GeneMAP identifies SLC25A48 as necessary for mitochondrial choline import. Nature genetics, 56(8), 1614.

Stefanakis N, et al. (2024) LET-381/FoxF and its target UNC-30/Pitx2 specify and maintain the molecular identity of C. elegans mesodermal glia that regulate motor behavior. The EMBO journal, 43(6), 956.

Kelley ME, et al. (2024) Spastin regulates anaphase chromosome separation distance and microtubule-containing nuclear tunnels. Molecular biology of the cell, 35(4), ar48.

Zhang S, et al. (2024) Resistance of estrogen receptor function to BET bromodomain inhibition is mediated by transcriptional coactivator cooperativity. bioRxiv : the preprint server for biology.

Liu Y, et al. (2023) Autoregulatory control of mitochondrial glutathione homeostasis. Science (New York, N.Y.), 382(6672), 820.

Brahma A, et al. (2023) Transcriptional and post-transcriptional control of odorant receptor choice in ants. Current biology : CB, 33(24), 5456.