Resource Summary Report

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New York University Grossman School of Medicine Cytometry and Cell Sorting Laboratory Core Facility

RRID:SCR_019179
Type: Tool

Proper Citation

New York University Grossman School of Medicine Cytometry and Cell Sorting Laboratory Core Facility (RRID:SCR_019179)

Resource Information

URL: https://med.nyu.edu/research/scientific-cores-shared-resources/cytometry-cell-sorting-laboratory

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Description: Provides access to flow cytometry and cell sorting technologies and instruments. If your research requires cytometric analysis, instruments acquire optical measurements using different lasers to detect fluorophores with high level of precision.

Synonyms: Cytometry and Cell Sorting Laboratory

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

Keywords: USEDit, flow cytometry, cell sorting, cytometric analysis, ABRF, ABRF

Funding: NCI P30CA016087

Resource Name: New York University Grossman School of Medicine Cytometry and Cell Sorting Laboratory Core Facility

Resource ID: SCR_019179

Alternate IDs: ABRF_821

Alternate URLs: https://coremarketplace.org/?FacilityID=821

Record Creation Time: 20220129T080343+0000

Record Last Update: 20250412T060255+0000

Ratings and Alerts

No rating or validation information has been found for New York University Grossman School of Medicine Cytometry and Cell Sorting Laboratory Core Facility.

No alerts have been found for New York University Grossman School of Medicine Cytometry and Cell Sorting Laboratory Core Facility.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 8 mentions in open access literature.

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

Samper N, et al. (2024) Kir6.1, a component of an ATP-sensitive potassium channel, regulates natural killer cell development. Frontiers in immunology, 15, 1490250.

Calderon A, et al. (2024) Chromatin accessibility and cell cycle progression are controlled by the HDAC-associated Sin3B protein in murine hematopoietic stem cells. Epigenetics & chromatin, 17(1), 2.

Cattle MA, et al. (2024) An enhanced Eco1 retron editor enables precision genome engineering in human cells from a single-copy integrated lentivirus. bioRxiv : the preprint server for biology.

Nakamura R, et al. (2024) High-dose methylprednisolone mediates YAP/TAZ-TEAD in vocal fold fibroblasts with macrophages. Research square.

Barcia Durán JG, et al. (2024) Immune checkpoint landscape of human atherosclerosis and influence of cardiometabolic factors. Nature cardiovascular research, 3(12), 1482.

Samper N, et al. (2024) Kir6.1, a component of an ATP-sensitive potassium channel, regulates natural killer cell development. bioRxiv : the preprint server for biology.

Glytsou C, et al. (2023) Mitophagy Promotes Resistance to BH3 Mimetics in Acute Myeloid Leukemia. Cancer discovery, 13(7), 1656.

Calderon A, et al. (2023) The Sin3B chromatin modifier restricts cell cycle progression to dictate hematopoietic stem cell differentiation. bioRxiv : the preprint server for biology.