Resource Summary Report

Generated by PRECISE-TBI on Aug 2, 2026

Pennsylvania State University Huck Institutes Flow Cytometry Core Facility

RRID:SCR_024460
Type: Tool

Proper Citation

Pennsylvania State University Huck Institutes Flow Cytometry Core Facility (RRID:SCR_024460)

Resource Information

URL: https://www.huck.psu.edu/core-facilities/flow-cytometry-facility

Proper Citation: Pennsylvania State University Huck Institutes Flow Cytometry Core Facility (RRID:SCR_024460)

Description: Facility is equipped with flow cytometers and cell sorters to examine cell samples within range of micron. Staff engage in experimentation, training, project collaboration, and consultation.

Synonyms: Huck Institutes' Flow Cytometry Facility

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

Keywords: ABRF, flow cytometry, cell sorting, experimentation, training, project collaboration, consultation,

Resource Name: Pennsylvania State University Huck Institutes Flow Cytometry Core Facility

Resource ID: SCR_024460

Alternate IDs: ABRF_2452

Alternate URLs: https://coremarketplace.org/RRID:SCR_024460?citation=1, https://coremarketplace.org/?FacilityID=2452&citation=1

Record Creation Time: 20230922T050237+0000

Record Last Update: 20260801T191249+0000

Ratings and Alerts

No rating or validation information has been found for Pennsylvania State University Huck Institutes Flow Cytometry Core Facility.

No alerts have been found for Pennsylvania State University Huck Institutes Flow Cytometry Core Facility.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at PRECISE-TBI .

Ruan B, et al. (2025) Itaconate promotes the differentiation of murine stress erythroid progenitors by increasing Nrf2 activity. Blood. Red cells & iron, 1(1).

Harris JE, et al. (2025) The activity of soil microbial taxa in the rhizosphere predicts the success of root colonization. mSystems, 10(9), e0045825.

Kulkarni S, et al. (2025) Auxin-induced depletion of human CCR4-NOT subunits reveals opposing functions of CNOT1 and CNOT4 in mRNA metabolism. The Journal of biological chemistry, 301(12), 110862.

Kulkarni S, et al. (2024) Human CCR4-NOT globally regulates gene expression and is a novel silencer of retrotransposon activation. bioRxiv : the preprint server for biology.