

# Resource Summary Report

Generated by [nidm-terms](#) on Aug 9, 2026

## University Health Network Princess Margaret Flow Cytometry Core Facility

RRID:SCR\_027537

Type: Tool

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### Proper Citation

University Health Network Princess Margaret Flow Cytometry Core Facility  
(RRID:SCR\_027537)

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### Resource Information

**URL:** <https://pmflow.ca/>

**Proper Citation:** University Health Network Princess Margaret Flow Cytometry Core Facility (RRID:SCR\_027537)

**Description:** Core offers services in Analytical Flow Cytometry, Fluorescence Activated Cell Sorting, consultation and data analysis. Staff provides training, assisted acquisition, consultation, and data analysis upon request.

**Synonyms:** Princess Margaret Flow Cytometry Facility

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

**Keywords:** ABRF, Analytical Flow Cytometry, Fluorescence Activated Cell Sorting, consultation, data analysis,

**Availability:** Open

**Resource Name:** University Health Network Princess Margaret Flow Cytometry Core Facility

**Resource ID:** SCR\_027537

**Alternate IDs:** ABRF\_5439

**Alternate URLs:** [https://coremarketplace.org/RRID:SCR\\_027537/?citation=1](https://coremarketplace.org/RRID:SCR_027537/?citation=1)

**Record Creation Time:** 20251011T055736+0000

**Record Last Update:** 20260808T190803+0000

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### Ratings and Alerts

No rating or validation information has been found for University Health Network Princess Margaret Flow Cytometry Core Facility.

No alerts have been found for University Health Network Princess Margaret Flow Cytometry Core Facility.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [nidm-terms](#) .

Ihara F, et al. (2026) CD28-driven ex vivo generation of stem-like memory CD8+ T cells bypassing CD3/TCR signaling. Proceedings of the National Academy of Sciences of the United States of America, 123(17), e2524626123.

Chen HTT, et al. (2026) Aberrant splicing of MBD1 reshapes the epigenome to drive convergent myeloerythroid defects in MDS. Blood.