

Resource Summary Report

Generated by [ASWG](#) on Sep 18, 2026

Dartmouth Genomics Shared Resource

RRID:SCR_009763

Type: Tool

Proper Citation

Dartmouth Genomics Shared Resource (RRID:SCR_009763)

Resource Information

URL: <http://dms.dartmouth.edu/dgml>

Proper Citation: Dartmouth Genomics Shared Resource (RRID:SCR_009763)

Description: Core facility that provides the following services: Illumina Bead Array service, Affymetrix Microarray service.

The Genomics Shared Resource provides technologies to the Norris Cotton Cancer Center and Dartmouth community investigators that enable profiling of gene expression, miRNA, GpC Island, and CGH on a whole-genome scale. The long-term goal of the Genomics Shared Resource is to provide an efficient and affordable fee-for-service operation that will provide high quality genomics and microarray data for the growing number of Cancer Center investigators who require this service.

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

Keywords: nucleic acid microarray assay

Resource Name: Dartmouth Genomics Shared Resource

Resource ID: SCR_009763

Alternate IDs: nlx_156226

Alternate URLs: <http://dartmouth.eagle-i.net/i/0000012e-0783-585d-4bd3-f7c180000000>, <http://cancer.dartmouth.edu/res/genomics.html>, <https://freedom7.dartmouth.edu/login>

Record Creation Time: 20220129T080254+0000

Record Last Update: 20260912T200321+0000

Ratings and Alerts

No rating or validation information has been found for Dartmouth Genomics Shared Resource.

No alerts have been found for Dartmouth Genomics Shared Resource.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [ASWG](#) .

Turnquist A, et al. (2026) PI3K?? inhibition alters CD8 T cell differentiation and reprograms the tumor microenvironment following adoptive immunotherapy. Journal of immunology (Baltimore, Md. : 1950), 215(5).