

Resource Summary Report

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Dartmouth Genomic Data Sciences Core Facility

RRID:SCR_025383

Type: Tool

Proper Citation

Dartmouth Genomic Data Sciences Core Facility (RRID:SCR_025383)

Resource Information

URL: <https://sites.dartmouth.edu/cqb/projects-and-cores/data-analytics-core/>

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Description: Core offers statistical analysis of genomics data. Provides suite of standardized analysis pipelines for common data types like RNA-seq, ATAC-seq, and genome assembly. For more complex datasets, such as single-cell genomics, provides tailored analysis solutions. Core team assists with all project stages, from experimental design to publication support and data management. GDSC hosts several multi-day bioinformatics workshops designed to introduce fundamental concepts for analysis of genomics data.

Abbreviations: GDSC

Synonyms: , Dartmouth Genomic Data Sciences Core, Genomic Data Sciences Core, Geisel School of Medicine at Dartmouth Genomic Data Sciences Core

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

Keywords: ABRF, statistical analysis, genomics data, RNA-seq data, ATAC-seq data, genome assembly data, data analysis, experimental design, publication support, data management,

Resource Name: Dartmouth Genomic Data Sciences Core Facility

Resource ID: SCR_025383

Alternate IDs: ABRF_2806

Alternate URLs: <https://coremarketplace.org/?FacilityID=2806&citation=1>

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Record Last Update: 20260731T043150+0000

Ratings and Alerts

No rating or validation information has been found for Dartmouth Genomic Data Sciences Core Facility.

No alerts have been found for Dartmouth Genomic Data Sciences Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Reinhold WC, et al. (2025) Acetalax and Bisacodyl for the Treatment of Triple-Negative Breast Cancer: A Combined Molecular and Preclinical Study. Cancer research communications, 5(2), 375.

Kolling FW, et al. (2025) Multi-site Assessment of Methods for Cell Preservation Upstream of Single Cell RNA Sequencing. bioRxiv : the preprint server for biology.