

Resource Summary Report

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Vermont University Genetics Network Bioinformatics Core Facility

RRID:SCR_017686

Type: Tool

Proper Citation

Vermont University Genetics Network Bioinformatics Core Facility (RRID:SCR_017686)

Resource Information

URL: <https://vgn.uvm.edu/bioinformatics/>

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Description: Core provides expertise in biostatistics, microarray data analysis, proteome informatics, next generation sequencing data analysis, functional analysis, database development and information technology, including data storage infrastructure and high performance computing. Working closely with VGN Proteomics Facility, offers investigators experimental design consultations, comprehensive data analysis, data management and publishing, and manuscript and grant support. Core personnel also engage in teaching and training activities for data analysis and compute resources necessary for VGN network investigators. Our goal is to provide network researchers with bioinformatics expertise.

Synonyms: Vermont Genetics Network Bioinformatics Core

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

Keywords: Vermont, genetic, network, bioinformatic, core, data, analysis, microarray, proteome, next, generation, sequencing, functional, database, core

Availability: Open

Resource Name: Vermont University Genetics Network Bioinformatics Core Facility

Resource ID: SCR_017686

Alternate IDs: SCR_017707, ABRF_8

Record Creation Time: 20220129T080336+0000

Record Last Update: 20260815T183132+0000

Ratings and Alerts

No rating or validation information has been found for Vermont University Genetics Network Bioinformatics Core Facility.

No alerts have been found for Vermont University Genetics Network Bioinformatics Core Facility.

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 [dkNET](#) .

Kennedy A, et al. (2024) Transcription factor VRT2 reinitiates vernalization when interrupted by warm temperatures in a temperate grass model. Plant physiology, 196(4), 2614.