

Resource Summary Report

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Duke University Molecular Genomics Core Facility

RRID:SCR_017860

Type: Tool

Proper Citation

Duke University Molecular Genomics Core Facility (RRID:SCR_017860)

Resource Information

URL: <http://dmpi.duke.edu/molecular-genomics-shared-resource>

Proper Citation: Duke University Molecular Genomics Core Facility (RRID:SCR_017860)

Description: Core offers variety of experimental platforms to facilitate genomics research. Accredited as Duke Shared Resource facility offers experience with genetic, genomic and epigenomic study design and technology, working closely with researchers to customize experiments to meet their needs. Applications include 10x Genomics NGS library generation for both single cell and gDNA experiments, DNA methylation microarrays, SNP genotyping and copy number microarrays, and Taqman targeted SNP genotyping.

Abbreviations: MGC

Synonyms: Molecular Genomics Core

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

Keywords: Genomic, epigenomic, design, technology, customize, experiment, library, single, cell, gDNA, DNA, methylation, microarray, SNP, genotyping, service, core, ABRF

Resource Name: Duke University Molecular Genomics Core Facility

Resource ID: SCR_017860

Alternate IDs: ABRF_682

Record Creation Time: 20220129T080337+0000

Record Last Update: 20260805T044418+0000

Ratings and Alerts

No rating or validation information has been found for Duke University Molecular Genomics Core Facility.

No alerts have been found for Duke University Molecular Genomics 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 [RRID](#) .

David-Bercholz J, et al. (2022) Conserved YKL-40 changes in mice and humans after postoperative delirium. Brain, behavior, & immunity - health, 26, 100555.