

Resource Summary Report

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University of Rochester Genomics Research Center Core Facility

RRID:SCR_012359

Type: Tool

Proper Citation

University of Rochester Genomics Research Center Core Facility (RRID:SCR_012359)

Resource Information

URL: <https://www.urmc.rochester.edu/research/rochester-genomics-center.aspx>

Proper Citation: University of Rochester Genomics Research Center Core Facility (RRID:SCR_012359)

Description: Provides core laboratory support, technical consultation, assistance with experimental design and data analysis for investigators using high throughput genomic sequencing, genotyping and gene expression in their research programs. Provides support for RNA-Seq, ATAC-seq, ChIP-Seq, Single Cell RNA-seq, Spatial Transcriptomics, Exome/Genome sequencing, Whole Genome and Reduced Representation Bisulfite Sequencing (WGBS, RRBS), 16S rRNA Sequencing, Small RNA-Seq, Metatranscriptomics, qRT-PCR, processing of RNA/DNA, as well as custom applications and approaches. Bioinformatic analysis support is also available.

Abbreviations: URGRC

Synonyms: URMG Genomics Research Center, UR Genomics Research Center, University of Rochester Medical Center Genomics Research Center

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

Keywords: Genomic sequencing, genotyping, gene expression, ABRF, USEDit

Availability: open

Resource Name: University of Rochester Genomics Research Center Core Facility

Resource ID: SCR_012359

Alternate IDs: ABRF_400, SciEx_12075

Alternate URLs: <http://www.scienceexchange.com/facilities/ur-genomics-research-center-rochester>, <https://coremarketplace.org/?FacilityID=400>

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Ratings and Alerts

No rating or validation information has been found for University of Rochester Genomics Research Center Core Facility.

No alerts have been found for University of Rochester Genomics Research Center Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

West JD, et al. (2025) The quantitative impact of 3'UTRs on gene expression. Nucleic acids research, 53(12).

Calcines-Rodríguez L, et al. (2025) Microglial interferon signaling and A β plaque pathology are enhanced in female 5xFAD Alzheimer's disease mice, independent of estrous cycle stage. Journal of neuroinflammation.

Cohen ED, et al. (2025) Whole genome transcriptomics reveal distinct atrial versus ventricular responses to neonatal hyperoxia. American journal of physiology. Heart and circulatory physiology, 328(4), H832.