

# Resource Summary Report

Generated by SPARC SAWG on Sep 12, 2026

## Case Western Reserve University Genomics Core Facility

RRID:SCR\_019021

Type: Tool

### Proper Citation

Case Western Reserve University Genomics Core Facility (RRID:SCR\_019021)

### Resource Information

**URL:** <https://case.edu/medicine/genomics/>

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**Description:** Core provides centralized resource working to take difficulty out of Next Generation Sequencing projects for both novice and experienced users alike. Provides assistance in all facets of experimentation from design to data analysis including grant and manuscript preparations. Offers NGS, Sanger, Informatics and Genotyping services based off of Illumina platform.

**Synonyms:** Case Western Reserve University CWRU Genomics Core, CWRU Genomics Core

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

**Keywords:** USEEdit, Next Generation Sequencing, data analysis, experiment design, grant preparation, manuscript preparation, genotyping services, Illumina, ABRF

**Availability:** open

**Resource Name:** Case Western Reserve University Genomics Core Facility

**Resource ID:** SCR\_019021

**Alternate IDs:** ABRF\_468

**Alternate URLs:** <https://coremarketplace.org/?FacilityID=468>

**Record Creation Time:** 20220129T080343+0000

**Record Last Update:** 20260912T200408+0000

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## Ratings and Alerts

No rating or validation information has been found for Case Western Reserve University Genomics Core Facility.

No alerts have been found for Case Western Reserve University Genomics Core Facility.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Sun S, et al. (2025) Combined effects of low temperature, hyperosmolarity and seawater-conditioned pathogens on open fracture healing in a rat model simulating circumpolar environments. Journal of orthopaedic surgery and research, 20(1), 875.

Vasu K, et al. (2021) Screening of CRISPR-Cas9-generated point mutant mice using MiSeq and locked nucleic acid probe PCR. STAR protocols, 2(4), 100785.