

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 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/>

Proper Citation: Case Western Reserve University Genomics Core Facility (RRID:SCR_019021)

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: 20260919T195937+0000

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.

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 [nidm-terms](#) .

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.