

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

University of Nebraska Medical Center UNMC Genomics Core Facility

RRID:SCR_023539

Type: Tool

Proper Citation

University of Nebraska Medical Center UNMC Genomics Core Facility
(RRID:SCR_023539)

Resource Information

URL: <https://www.unmc.edu/vcr/cores/vcr-cores/genomics/index.html>

Proper Citation: University of Nebraska Medical Center UNMC Genomics Core Facility
(RRID:SCR_023539)

Description: Provides Next Generation DNA and Sanger sequencing services; single cell genomics assays by 10x Genomics, and Nanostring technology targeted gene expression assays.

Synonyms: UNMC Genomics Core Facility

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

Keywords: USEEdit, ABRF, Next Generation DNA and Sanger sequencing services,

Resource Name: University of Nebraska Medical Center UNMC Genomics Core Facility

Resource ID: SCR_023539

Alternate IDs: ABRF_1757

Alternate URLs: <https://coremarketplace.org/?FacilityID=1757&citation=1>

Record Creation Time: 20230503T050210+0000

Record Last Update: 20260808T190659+0000

Ratings and Alerts

No rating or validation information has been found for University of Nebraska Medical Center UNMC Genomics Core Facility.

No alerts have been found for University of Nebraska Medical Center UNMC Genomics 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 [SPARC SAWG](#) .

Hernandez JB, et al. (2026) Mapping the metagenomic landscape: combined shotgun sequencing and quantitative PCR to profile gut metagenome-assembled genomes in marmosets following treatment with a broad-spectrum antibiotic cocktail. Gut microbes, 18(1), 2687925.

Tamanaeifar F, et al. (2026) Luteinizing hormone activates the Hippo pathway to promote progesterone synthesis in bovine luteal cells. Cell communication and signaling : CCS, 24(1).

Hernandez C, et al. (2026) An Exploratory Analysis of Transcriptional Responses to Peanut Exposure in Drosophila melanogaster. International journal of molecular sciences, 27(10).