

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## Marshall University School of Medicine Genomics Core Facility

RRID:SCR\_018885

Type: Tool

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### Proper Citation

Marshall University School of Medicine Genomics Core Facility (RRID:SCR\_018885)

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### Resource Information

**URL:** <https://jcesom.marshall.edu/research/office-of-research-and-graduate-education/core-facilities/genomics/>

**Proper Citation:** Marshall University School of Medicine Genomics Core Facility (RRID:SCR\_018885)

**Description:** Core provides services including high throughput next generation sequencing (NGS) to support whole genome, whole exome, RNA-Seq, single cell RNA-Seq, microbiome and global chromatin and methylation studies, biostatistical and bioinformatic support for NGS projects, access to DNA/RNA sequence analysis software, automated Sanger DNA sequencing, genotyping and RNA/DNA quality assessment, access to shared instrumentation such as plate readers, real time thermal cyclers, Agilent Bioanalyzers, fluorimeters, and spectrophotometers.

**Synonyms:** Marshall University Genomics Core Facility, MU Genomics Core Facility, Marshall University Genomics and Bioinformatics Core Facilities

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

**Keywords:** USEdit, genomics, next generation sequencing, biostatistical and bioinformatic support, RNA-Seq, global chromatin and methylation studies, ABRF

**Funding:** NIGMS 1P20GM121299;  
NIGMS 2U54GM104942;  
NIGMS P20GM103434

**Resource Name:** Marshall University School of Medicine Genomics Core Facility

**Resource ID:** SCR\_018885

**Alternate IDs:** ABRF\_163

**Alternate URLs:** <https://coremarketplace.org/?FacilityID=163>

**Record Creation Time:** 20220129T080342+0000

**Record Last Update:** 20260905T133417+0000

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## Ratings and Alerts

No rating or validation information has been found for Marshall University School of Medicine Genomics Core Facility.

No alerts have been found for Marshall University School of Medicine Genomics Core Facility.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Shultz C, et al. (2023) Association of genetic variants and survival in patients with acute myeloid leukemia in rural Appalachia. *Cancer reports* (Hoboken, N.J.), 6(3), e1746.

Barrett CF, et al. (2023) Mitochondrial genome sequencing and analysis of the invasive *Microstegium vimineum* : a resource for systematics, invasion history, and management. *bioRxiv* : the preprint server for biology.

Lin J, et al. (2023) RNA-Seq Dissects Incomplete Activation of Phytoalexin Biosynthesis by the Soybean Transcription Factors GmMYB29A2 and GmNAC42-1. *Plants* (Basel, Switzerland), 12(3).

Barrett CF, et al. (2022) Digitized collections elucidate invasion history and patterns of awn polymorphism in *Microstegium vimineum*. *American journal of botany*, 109(5), 689.

Barrett CF, et al. (2022) Lineage and role in integrative taxonomy of a heterotrophic orchid complex. *Molecular ecology*, 31(18), 4762.

Souder K, et al. (2021) Role of *dipA* and *pilD* in *Francisella tularensis* Susceptibility to Resazurin. *Antibiotics* (Basel, Switzerland), 10(8).