

Resource Summary Report

Generated by ASWG on Aug 23, 2026

Michigan State University Bioinformatics Core Facility

RRID:SCR_026706

Type: Tool

Proper Citation

Michigan State University Bioinformatics Core Facility (RRID:SCR_026706)

Resource Information

URL: <https://bioinformatics.msu.edu>

Proper Citation: Michigan State University Bioinformatics Core Facility (RRID:SCR_026706)

Description: Core brings together expertise from across MSU to support research through advanced omics data analysis, grant proposal assistance, and tailored training for researchers at all career stages.

Synonyms: , MSU Bioinformatics Core, Michigan State University MSU Bioinformatics Core

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

Keywords: ABRF, omics data analysis, grant proposal assistance, expertise, training,

Resource Name: Michigan State University Bioinformatics Core Facility

Resource ID: SCR_026706

Alternate IDs: ABRF_3127

Alternate URLs: <https://coremarketplace.org/?FacilityID=3127&citation=1>

Record Creation Time: 20250404T060341+0000

Record Last Update: 20260821T194344+0000

Ratings and Alerts

No rating or validation information has been found for Michigan State University Bioinformatics Core Facility.

No alerts have been found for Michigan State University Bioinformatics Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Mugume Y, et al. (2026) A Lipoxygenase 3 mutation reverses growth phenotypes in an Arabidopsis Plastid Lipase 3 overexpression line. PloS one, 21(6), e0350738.

Bahmani K, et al. (2026) -Identification of QTL for steviol glycoside biosynthesis using a SNP-based genetic linkage map for Stevia rebaudiana. G3 (Bethesda, Md.), 16(3).

Scarborough T, et al. (2026) Tangential flow filtration for isolating exomeres and other nanoscale extracellular particles. Nanoscale, 18(29), 15554.

Scarborough T, et al. (2025) A scalable filtration-based method for isolating exomeres and other nanoscale extracellular particles. bioRxiv : the preprint server for biology.