

# Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

## Michigan State University Bioinformatics Core Facility

RRID:SCR\_026706

Type: Tool

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### Proper Citation

Michigan State University Bioinformatics Core Facility (RRID:SCR\_026706)

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### 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:** 20260905T133529+0000

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### 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.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 4 mentions in open access literature.

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

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.