

# Resource Summary Report

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## MSU Proteomics Core Laboratory

RRID:SCR\_009944

Type: Tool

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### Proper Citation

MSU Proteomics Core Laboratory (RRID:SCR\_009944)

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### Resource Information

**URL:** <http://montana.eagle-i.net/i/0000012b-00be-96b8-df3b-3fdc80000000>

**Proper Citation:** MSU Proteomics Core Laboratory (RRID:SCR\_009944)

**Description:** Core facility that provides the following services: MicroTOF (access), GCMS (access), MALDI (access), XCT ion trap (access), MicroTOF high resolution (facility run), MicroTOF low res (facility run), MALDI (facility run), GCMS (facility run), HPLC-microTOF (facility run), 2-D gel (access), 2-D gel (facility run).

The mission of the Mass Spectrometry and Proteomics Facility is to seed mass spectrometry and proteomics technology to research labs at Montana State University and affiliated programs.

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

**Keywords:** mass spectrometry assay, high performance liquid chromatography, protein mass determination by mass spectrometry, protein identification, preparative chromatography, protein electrophoresis

**Resource Name:** MSU Proteomics Core Laboratory

**Resource ID:** SCR\_009944

**Alternate IDs:** nlx\_156414

**Alternate URLs:** <http://cores.montana.edu/index.php?page=proteomics>

**Record Creation Time:** 20220129T080255+0000

**Record Last Update:** 20260919T195849+0000

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### Ratings and Alerts

No rating or validation information has been found for MSU Proteomics Core Laboratory.

No alerts have been found for MSU Proteomics Core Laboratory.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.