

Resource Summary Report

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

MSU Proteomics Core Laboratory

RRID:SCR_009944

Type: Tool

Proper Citation

MSU Proteomics Core Laboratory (RRID:SCR_009944)

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: 20260905T133337+0000

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.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.