Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

University of Michigan Medical School BRCF Proteomics Resource Core Facility

RRID:SCR_026723
Type: Tool

Proper Citation

University of Michigan Medical School BRCF Proteomics Resource Core Facility (RRID:SCR_026723)

Resource Information

URL: https://cores.research.umich.edu/core/proteomics-resource-facility/

Proper Citation: University of Michigan Medical School BRCF Proteomics Resource Core Facility (RRID:SCR_026723)

Description: Core is dedicated to protein analysis and applying mass spectrometry-based proteomic approaches to varied biological questions.

Synonyms: , University of Michigan Medical School Pathology BRCF Proteomics Resource Core Facility, University of Michigan Medical School Pathology BRCF Proteomics Resource Facility, University of Michigan BRCF Proteomics Resource Facility

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

Keywords: ABRF, protein analysis, mass spectrometry

Availability: Restricted

Resource Name: University of Michigan Medical School BRCF Proteomics Resource Core Facility

Resource ID: SCR_026723

Alternate IDs: ABRF_3195

Alternate URLs: https://www.pathology.med.umich.edu/login, https://coremarketplace.org/RRID:SCR_026723/?citation=1

Record Creation Time: 20250409T060321+0000

Record Last Update: 20260905T133530+0000

Ratings and Alerts

No rating or validation information has been found for University of Michigan Medical School BRCF Proteomics Resource Core Facility.

No alerts have been found for University of Michigan Medical School BRCF Proteomics Resource Core Facility.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at PRECISE-TBI .

Upadhyay KK, et al. (2026) The inner nuclear membrane protein SUN1 regulates cullin-3 neddylation to maintain insulin signaling. bioRxiv : the preprint server for biology.

Tang VT, et al. (2026) Endothelial Cell Secretome Alterations Induced by Inflammatory Stress. bioRxiv : the preprint server for biology.

Popelka H, et al. (2026) Atg7-Atg12 conjugation and autophagy are negatively regulated by the disordered region of Atg12. Communications biology, 9(1).