

# Resource Summary Report

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

## Wayne State University Proteomics Core Facility

RRID:SCR\_022879

Type: Tool

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### Proper Citation

Wayne State University Proteomics Core Facility (RRID:SCR\_022879)

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### Resource Information

**URL:** <https://research.wayne.edu/proteomics>

**Proper Citation:** Wayne State University Proteomics Core Facility (RRID:SCR\_022879)

**Description:** Protein identification and proteome analysis using Orbitrao Eclipse,Fusion and Q Exactive LC-MS/MS platforms. Affinity selection for phosphopeptides with analysis using HCD, CID and ETD. Validation and quantitation of proteins using MRM with a TSQ Vantage triple quad.

**Synonyms:** WSU Proteomics Core, Wayne State University WSU Proteomics Core

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

**Keywords:** USEDit, ABRF, Protein identification and proteome analysis

**Resource Name:** Wayne State University Proteomics Core Facility

**Resource ID:** SCR\_022879

**Alternate IDs:** ABRF\_373

**Alternate URLs:** <https://coremarketplace.org/?FacilityID=373&citation=1>

**Record Creation Time:** 20221013T050144+0000

**Record Last Update:** 20260905T133430+0000

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### Ratings and Alerts

No rating or validation information has been found for Wayne State University Proteomics Core Facility.

No alerts have been found for Wayne State University Proteomics Core Facility.

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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.