

Resource Summary Report

Generated by PRECISE-TBI on Jul 31, 2026

University of Texas San Antonio Mass Spectrometry and Proteomics Core Facility

RRID:SCR_010151

Type: Tool

Proper Citation

University of Texas San Antonio Mass Spectrometry and Proteomics Core Facility
(RRID:SCR_010151)

Resource Information

URL: <https://research.utsa.edu/cores/mspc/>

Proper Citation: University of Texas San Antonio Mass Spectrometry and Proteomics Core Facility (RRID:SCR_010151)

Description: Core provides routine and custom mass spectrometry-based support. Services include High-resolution, high mass accuracy mass analysis (HRMS), Liquid chromatography-mass spectrometry (LC-MS), Proteomics-based analyses.

Abbreviations: UTSA MSPC

Synonyms: , USTA RCMI Proteomics & Protein Biomarkers Core, UTSA Mass Spectrometry & Proteomics Core

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

Keywords: ABRF, liquid chromatography-mass spectrometry, proteomics-based analyses, mass spectrometry,

Availability: <https://coremarketplace.org/?FacilityID=3095&citation=1>

Resource Name: University of Texas San Antonio Mass Spectrometry and Proteomics Core Facility

Resource ID: SCR_010151

Alternate IDs: SCR_011034, SciEx_9630, nlx_156630, ABRF_3095

Old URLs: <http://utsa.eagle-i.net/i/00000135-58f7-6c34-d7c8-cf3780000000>

Record Creation Time: 20220129T080257+0000

Record Last Update: 20260731T042830+0000

Ratings and Alerts

No rating or validation information has been found for University of Texas San Antonio Mass Spectrometry and Proteomics Core Facility.

No alerts have been found for University of Texas San Antonio Mass Spectrometry and Proteomics Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.