Resource Summary Report

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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/

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Description: Core provides routine and custom mass spectrometry-based support. Services include High-resolution, high mass accuracy mass analysis (HRMS), Liquid chromatographymass 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, access service resource, service resource

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

Funding:

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

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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 found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Mathew B, et al. (2021) Exosomes as Emerging Biomarker Tools in Neurodegenerative and Neuropsychiatric Disorders-A Proteomics Perspective. Brain sciences, 11(2).