

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

UTEP Biomolecule Analysis Core Facility

RRID:SCR_010139

Type: Tool

Proper Citation

UTEP Biomolecule Analysis Core Facility (RRID:SCR_010139)

Resource Information

URL: <http://utep.eagle-i.net/i/00000134-a9b2-8e15-bfc0-fe9780000000>

Proper Citation: UTEP Biomolecule Analysis Core Facility (RRID:SCR_010139)

Description: Core facility that provides the following services: Mass spectrometry analysis, Trypsin Digestion. The Biomolecule Analysis Core Facility (BACF) specializes in the separation and characterization of biomolecules. The facility possesses state-of-the-art equipment for proteomic analysis and the characterization of molecular binding interactions. These resources promote new insight into the mechanism of infectious agents, cancer, and other diseases that affect the population of the Paseo del Norte region. BACF services are available to external collaborators for a small fee.

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

Keywords: mass spectrometry assay, liquid chromatography-mass spectrometry

Resource Name: UTEP Biomolecule Analysis Core Facility

Resource ID: SCR_010139

Alternate IDs: nlx_156618

Record Creation Time: 20220129T080257+0000

Record Last Update: 20260731T042830+0000

Ratings and Alerts

No rating or validation information has been found for UTEP Biomolecule Analysis Core Facility.

No alerts have been found for UTEP Biomolecule Analysis Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.