

Resource Summary Report

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University of North Carolina at Chapel Hill Department of Chemistry Mass Spectrometry Core Facility

RRID:SCR_021937

Type: Tool

Proper Citation

University of North Carolina at Chapel Hill Department of Chemistry Mass Spectrometry Core Facility (RRID:SCR_021937)

Resource Information

URL: <http://mscore.web.unc.edu>

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Description: Core specializes in small molecule analysis. Services include, but are not limited to, quantitative mass spectrometry, liquid chromatography separations, fragmentation and structural elucidation mass spectrometry, complex mixture analysis, molecular formula confirmation, high resolution and accurate mass analysis. Provides individualized user training for undergraduate, graduate, and post-doctoral scholars of Department of Chemistry.

Synonyms: University of North Carolina at Chapel Hill Department of Chemistry Mass Spectrometry Core Laboratory, Department of Chemistry Mass Spectrometry Core Laboratory

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

Keywords: USEDit, ABRF, small molecule analysis, mass spectrometry, liquid chromatography

Funding:

Availability: open

Resource Name: University of North Carolina at Chapel Hill Department of Chemistry Mass

Spectrometry Core Facility

Resource ID: SCR_021937

Alternate IDs: ABRF_655

Alternate URLs: <https://coremarketplace.org/?FacilityID=655>

Record Creation Time: 20220421T050137+0000

Record Last Update: 20250409T061742+0000

Ratings and Alerts

No rating or validation information has been found for University of North Carolina at Chapel Hill Department of Chemistry Mass Spectrometry Core Facility.

No alerts have been found for University of North Carolina at Chapel Hill Department of Chemistry Mass Spectrometry 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](#).

Hunt EG, et al. (2024) Acetyl-CoA carboxylase obstructs CD8+ T cell lipid utilization in the tumor microenvironment. Cell metabolism.