

Resource Summary Report

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Emory University Emory Integrated Proteomics Core Facility

RRID:SCR_023530

Type: Tool

Proper Citation

Emory University Emory Integrated Proteomics Core Facility (RRID:SCR_023530)

Resource Information

URL: <https://www.cores.emory.edu/eipc/>

Proper Citation: Emory University Emory Integrated Proteomics Core Facility (RRID:SCR_023530)

Description: Core is full service proteomics facility offering Emory researchers ability to use the latest technologies in their research.

Abbreviations: EIPC

Synonyms: Emory University Emory Integrated Proteomics Core (EIPC), Emory Integrated Proteomics Core (EIPC)

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

Keywords: USEDit, ABRF, proteomics services,

Funding: Winship Cancer Institute ;
Georgia Clinical and Translational Science Alliance ;
Emory University School of Medicine

Resource Name: Emory University Emory Integrated Proteomics Core Facility

Resource ID: SCR_023530

Alternate IDs: ABRF_1745

Alternate URLs: <https://coremarketplace.org/?FacilityID=1745&citation=1>

Record Creation Time: 20230503T050210+0000

Record Last Update: 20260728T043659+0000

Ratings and Alerts

No rating or validation information has been found for Emory University Emory Integrated Proteomics Core Facility.

No alerts have been found for Emory University Emory Integrated Proteomics Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 11 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Chi JJ, et al. (2025) MGAT1-Guided complex N-Glycans on CD73 regulate immune evasion in triple-negative breast cancer. *Nature communications*, 16(1), 3552.

Neiswender H, et al. (2025) Dominant spinal muscular atrophy linked mutations in the cargo binding domain of BICD2 result in altered interactomes and dynein hyperactivity. *eLife*, 14.

Neiswender H, et al. (2025) Dominant spinal muscular atrophy linked mutations in the cargo binding domain of BICD2 result in altered interactomes and dynein hyperactivity. *bioRxiv : the preprint server for biology*.

Farhi J, et al. (2025) Dynamic In Vivo Mapping of the Methylproteome Using a Chemoenzymatic Approach. *Journal of the American Chemical Society*, 147(9), 7214.

Cooper GW, et al. (2025) SMARCB1 missense mutants disrupt SWI/SNF complex stability and remodeling activity. *Research square*.

Makkaoui N, et al. (2024) Cell-based therapies reverse the heart failure-altered right ventricular proteome towards a pre-disease state. *Stem cell research & therapy*, 15(1), 420.

Knippler CM, et al. (2024) Bisbiguanide analogs induce mitochondrial stress to inhibit lung cancer cell invasion. *iScience*, 27(4), 109591.

Yoon SB, et al. (2024) Subpopulation commensalism promotes Rac1-dependent invasion of single cells via laminin-332. *The Journal of cell biology*, 223(6).

Paz E, et al. (2024) Biochemical and neurophysiological effects of deficiency of the mitochondrial import protein TIMM50. *eLife*, 13.

Makkaoui N, et al. (2024) Cell-based therapies reverse the heart failure-altered right ventricular proteome. *Research square*.

Bordan Z, et al. (2024) PDZ-Binding Kinase, a Novel Regulator of Vascular Remodeling in Pulmonary Arterial Hypertension. *Circulation*, 150(5), 393.