## **Resource Summary Report**

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# HMS Taplin Mass Spectrometry Core Facility

RRID:SCR\_009813 Type: Tool

#### **Proper Citation**

HMS Taplin Mass Spectrometry Core Facility (RRID:SCR\_009813)

#### **Resource Information**

URL: http://harvard.eagle-i.net/i/0000012b-00c3-414f-db6e-7a3f80000000

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**Description:** Core facility that provides the following services: Protein identification from SDS-PAGE gel bands/spots (Coomassie stained level of protein), Protein phosphorylation service, TCA samples service. The Taplin Biological Mass Spectrometry Facility opened in February 2001 as a core facility for the analysis of proteins and peptides by mass spectrometry. The facility is focused on serving the needs of investigators at Harvard Medical School (HMS) and all the Harvard affiliated Institutions.

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

Keywords: protein identification, mass spectrometry assay

Funding:

Resource Name: HMS Taplin Mass Spectrometry Core Facility

Resource ID: SCR\_009813

Alternate IDs: nlx\_156280

Alternate URLs: https://taplin.med.harvard.edu/

Record Creation Time: 20220129T080255+0000

Record Last Update: 20250420T015929+0000

#### **Ratings and Alerts**

No rating or validation information has been found for HMS Taplin Mass Spectrometry Core Facility.

No alerts have been found for HMS Taplin Mass Spectrometry Core Facility.

#### Data and Source Information

Source: <u>SciCrunch Registry</u>

### Usage and Citation Metrics

We have not found any literature mentions for this resource.