

Resource Summary Report

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UNC School of Medicine Macromolecular X-Ray Crystallography Core Facility

RRID:SCR_012327

Type: Tool

Proper Citation

UNC School of Medicine Macromolecular X-Ray Crystallography Core Facility
(RRID:SCR_012327)

Resource Information

URL: <http://www.scienceexchange.com/facilities/macromolecular-x-ray-crystallography-core-facility--2>

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Description: X-ray crystallography is the principal technique used to determine three-dimensional structures of biological macromolecules. The central goal of the UNC Crystallography Core Facility is to provide the support and infrastructure necessary to enable individual investigators of any experience level to initiate and successfully complete a structural biology project. The facility is state of the art, offering a variety of services in these three areas: crystallization, x-ray diffraction, structure determination. The primary mission of the MX Facility is to work with researchers here at UNC, however, we are also able to work with researchers at non-UNC academic institutions and commercial companies.

Abbreviations: UNC Crystallography Core Facility

Synonyms: University of North Carolina at Chapel Hill School of Medicine Macromolecular X-Ray Crystallography Core Facility

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

Keywords: x-ray crystallography

Resource Name: UNC School of Medicine Macromolecular X-Ray Crystallography Core Facility

Resource ID: SCR_012327

Alternate IDs: SciEx_11789

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Ratings and Alerts

No rating or validation information has been found for UNC School of Medicine Macromolecular X-Ray Crystallography Core Facility.

No alerts have been found for UNC School of Medicine Macromolecular X-Ray Crystallography Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.