Resource Summary Report

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Max Planck Institute of Biochemistry Crystallization Core Facility

RRID:SCR_025740
Type: Tool

Proper Citation

Max Planck Institute of Biochemistry Crystallization Core Facility (RRID:SCR_025740)

Resource Information

URL: https://www.biochem.mpg.de/crystallization

Proper Citation: Max Planck Institute of Biochemistry Crystallization Core Facility (RRID:SCR_025740)

Description: Facility for protein crystallisation and X-Ray crystallography. Provides structural biology infrastructure and performs crystallization experiments.

Abbreviations: MPIB-CCF

Synonyms: Max Planck Institute of Biochemistry Cristallization Facility, Cristallization Facility

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

Keywords: protein crystallisation, X-Ray crystallography,

Availability: Restricted

Resource Name: Max Planck Institute of Biochemistry Crystallization Core Facility

Resource ID: SCR_025740

Record Creation Time: 20240918T053245+0000

Record Last Update: 20260728T043728+0000

Ratings and Alerts

No rating or validation information has been found for Max Planck Institute of Biochemistry Crystallization Core Facility.

No alerts have been found for Max Planck Institute of Biochemistry Crystallization 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 nidm-terms .

Yuste-Checa P, et al. (2025) Structural analyses define the molecular basis of clusterin chaperone function. Nature structural & molecular biology.