

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## Max Planck Institute of Biochemistry Crystallization Core Facility

RRID:SCR\_025740

Type: Tool

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### Proper Citation

Max Planck Institute of Biochemistry Crystallization Core Facility (RRID:SCR\_025740)

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### 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 Crystallization Facility, Crystallization 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:** 20260905T133506+0000

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### 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.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [Kravitz](#) .

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