

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

## Max Planck Institute of Biochemistry Cryo-EM Core Facility

RRID:SCR\_025744

Type: Tool

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### Proper Citation

Max Planck Institute of Biochemistry Cryo-EM Core Facility (RRID:SCR\_025744)

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### Resource Information

**URL:** <https://www.biochem.mpg.de/cryoem>

**Proper Citation:** Max Planck Institute of Biochemistry Cryo-EM Core Facility (RRID:SCR\_025744)

**Description:** Facility for cryo-electron microscopy. Provides microscope access, training as well as scientific and technical support for MPIB-internal users wishing to perform cryo-EM experiments, from sample preparation to data processing.

**Abbreviations:** MPIB\_CEMF

**Synonyms:** Max Planck Institute of Biochemistry Cryo-EM Facility, Cryo-EM Facility

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

**Keywords:** cryo-electron microscopy, sample preparation, data processing

**Availability:** Restricted

**Resource Name:** Max Planck Institute of Biochemistry Cryo-EM Core Facility

**Resource ID:** SCR\_025744

**Record Creation Time:** 20240918T053245+0000

**Record Last Update:** 20260729T044628+0000

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### Ratings and Alerts

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

No alerts have been found for Max Planck Institute of Biochemistry Cryo-EM Core Facility.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Du J, et al. (2026) E2 variants for probing E3 ubiquitin ligase activities. Proceedings of the National Academy of Sciences of the United States of America, 123(1), e2524899122.

Farnung J, et al. (2026) The E3 ubiquitin ligase mechanism specifying target-directed microRNA degradation. bioRxiv : the preprint server for biology.

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

Hermann MD, et al. (2025) Normal macrophage signaling and gene expression in Rosa26 Cas9-expressing mice. ImmunoHorizons, 9(11).

Maiwald SA, et al. (2025) TRIP12 structures reveal HECT E3 formation of K29 linkages and branched ubiquitin chains. Nature structural & molecular biology, 32(9), 1766.

Iermak I, et al. (2025) Molecular mechanisms governing the formation of distinct Upf1-containing complexes in yeast. Cell reports, 44(10), 116415.