Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

Max Planck Institute of Biochemistry Bioinformatics Core Facility

RRID:SCR_025742
Type: Tool

Proper Citation

Max Planck Institute of Biochemistry Bioinformatics Core Facility (RRID:SCR_025742)

Resource Information

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

Proper Citation: Max Planck Institute of Biochemistry Bioinformatics Core Facility (RRID:SCR_025742)

Description: Facility for bioinformatics. Offers to assist wet-lab researchers in our institute for their bioinformatics data analysis.

Abbreviations: MPIB-BIO

Synonyms: Bioinformatics Core Facility, Max Planck Institute of Biochemistry Bioinformatics Core Facility

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

Keywords: bioinformatics data analysis,

Availability: Restricted

Resource Name: Max Planck Institute of Biochemistry Bioinformatics Core Facility

Resource ID: SCR_025742

Record Creation Time: 20240918T053245+0000

Record Last Update: 20260912T200451+0000

Ratings and Alerts

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

No alerts have been found for Max Planck Institute of Biochemistry Bioinformatics Core Facility.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at SPARC SAWG .

Bonnet J, et al. (2026) Histone modification cross-talk and protein complex diversification confer plasticity to Polycomb repression. Genes & development, 40(1-2), 43.

Andree GA, et al. (2026) Cysteine availability tunes ubiquitin signaling via inverse stability of LRRC58 E3 ligase and its substrate CDO1. Nature communications, 17(1).

Guyot C, et al. (2026) IL4i1 activity generates oncometabolites that rescue neuroblastoma cells from oxidative death. Cell reports, 45(6), 117441.

Russier M, et al. (2025) Differential cell survival outcomes in response to diverse amino acid stress. Life science alliance, 8(11).