

Resource Summary Report

Generated by T1D on Aug 1, 2026

International Institute of Molecular and Cell Biology in Warsaw, Biophysics and Bioanalytics Facility

RRID:SCR_021630

Type: Tool

Proper Citation

International Institute of Molecular and Cell Biology in Warsaw, Biophysics and Bioanalytics Facility (RRID:SCR_021630)

Resource Information

URL: <http://core-facility.iimcb.gov.pl/>

Proper Citation: International Institute of Molecular and Cell Biology in Warsaw, Biophysics and Bioanalytics Facility (RRID:SCR_021630)

Description: Offers comprehensive selection of molecular biophysics techniques such as SPR, SEC-MALS, AUC, ITC, CD, FT-IR useful in characterization of macromolecule size and intermolecular interactions.

Synonyms: International Institute of Molecular and Cell Biology IIMCB-Biophysics and Structural Biology Core Facility, IIMCB-Biophysics and Structural Biology Facility

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

Keywords: USEDit, ABRF, molecular biophysics techniques

Availability: open

Resource Name: International Institute of Molecular and Cell Biology in Warsaw, Biophysics and Bioanalytics Facility

Resource ID: SCR_021630

Alternate IDs: ABRF_1068

Alternate URLs: <https://coremarketplace.org/?FacilityID=1068>

Record Creation Time: 20220129T080356+0000

Record Last Update: 20260731T043104+0000

Ratings and Alerts

No rating or validation information has been found for International Institute of Molecular and Cell Biology in Warsaw, Biophysics and Bioanalytics Facility.

No alerts have been found for International Institute of Molecular and Cell Biology in Warsaw, Biophysics and Bioanalytics Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#).

Slyvka A, et al. (2025) Activity and structure of human (d)CTP deaminase CDADC1. Proceedings of the National Academy of Sciences of the United States of America, 122(19), e2424245122.

Krawczyk PS, et al. (2025) Re-adenylation by TENT5A enhances efficacy of SARS-CoV-2 mRNA vaccines. Nature, 641(8064), 984.

Idlin N, et al. (2025) Effects of genetic ablation and pharmacological inhibition of HuR on gene expression, iron metabolism, and hormone levels. BMC biology, 23(1), 24.

Nirwal S, et al. (2025) Structural snapshots of the mechanism of ATP-dependent DNA damage recognition by UvrA. Nature communications, 17(1), 387.

Fedenko A, et al. (2025) Etiology of TP53 mutated complex karyotype acute myeloid leukemia. Leukemia.

Thapa P, et al. (2024) HSP70 inhibits CHIP E3 ligase activity to maintain germline function in Caenorhabditis elegans. The Journal of biological chemistry, 300(11), 107864.

Matsuda A, et al. (2024) Despite the odds: formation of the SARS-CoV-2 methylation complex. Nucleic acids research, 52(11), 6441.

Borek B, et al. (2023) Arginase 1/2 Inhibitor OATD-02: From Discovery to First-in-man Setup in Cancer Immunotherapy. Molecular cancer therapeutics, 22(7), 807.

Das A, et al. (2022) A heterotypic assembly mechanism regulates CHIP E3 ligase activity. The EMBO journal, 41(15), e109566.