

Resource Summary Report

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University of Louisville School of Medicine Molecular Modeling Core Facility

RRID:SCR_026183

Type: Tool

Proper Citation

University of Louisville School of Medicine Molecular Modeling Core Facility
(RRID:SCR_026183)

Resource Information

URL: <https://centers.louisville.edu/brown-cancer-center/core-facilities/molecular-modeling>

Proper Citation: University of Louisville School of Medicine Molecular Modeling Core Facility (RRID:SCR_026183)

Description: Core offers services including molecular modeling of proteins and nucleic acids, virtual drug screening, computational biophysics.

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

Keywords: ABRF, molecular modeling, proteins, nucleic acids, virtual drug screening, computational biophysics,

Resource Name: University of Louisville School of Medicine Molecular Modeling Core Facility

Resource ID: SCR_026183

Alternate IDs: ABRF_2993

Alternate URLs: <https://coremarketplace.org/?FacilityID=2993&citation=1>

Old URLs: <https://louisville.edu/medicine/cancer-research/cores-and-facilities-1/mole-model>

Record Creation Time: 20241214T053246+0000

Record Last Update: 20260829T183438+0000

Ratings and Alerts

No rating or validation information has been found for University of Louisville School of Medicine Molecular Modeling Core Facility.

No alerts have been found for University of Louisville School of Medicine Molecular Modeling 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 [dkNET](#) .

Tarvestad-Laise K, et al. (2026) A c-Cbl/Cbl-b antagonist inhibits EGFR ubiquitylation and sustains EGFR phosphorylation to enhance corneal re-epithelialization. Proceedings of the National Academy of Sciences of the United States of America, 123(7), e2518857123.