

Resource Summary Report

Generated by [NIF](#) on Apr 20, 2025

Molecular Connections NetPro

RRID:SCR_000395

Type: Tool

Proper Citation

Molecular Connections NetPro (RRID:SCR_000395)

Resource Information

URL: <http://www.molecularconnections.com/home/en/home/products/netPro>

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Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 1, 2023. Comprehensive database of Protein-Protein and Protein-Small molecules interaction, consisting of more than 320,000 interactions captured from more than 1500 abstracts, approximately 1600 published journals and more than 60,000 references. The strength of NetPro lies in the complete manual curation of literature. It covers several entities other than proteins as interacting partners, like RNA, DNA, processes, etc. with well defined, exhaustive interaction terms. NetPro has received several accolades for the quality and quantity of data it contains. It has become an important resource for target identification, validation and pathway research and has subscribers from all over the globe including 3 of the top 5 pharmas.

Abbreviations: MolCon, NetPro

Resource Type: database, data or information resource

Keywords: drug, molecular, pharmaceutical, interaction, protein interaction, protein, protein-protein interaction, nucleic acid-protein, small molecule-protein, nucleic acid, small molecule

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Molecular Connections NetPro

Resource ID: SCR_000395

Alternate IDs: nif-0000-20877

Record Creation Time: 20220129T080201+0000

Record Last Update: 20250420T015439+0000

Ratings and Alerts

No rating or validation information has been found for Molecular Connections NetPro.

No alerts have been found for Molecular Connections NetPro.

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 [NIF](#).

Ammari MG, et al. (2016) HPIDB 2.0: a curated database for host-pathogen interactions.
Database : the journal of biological databases and curation, 2016.