

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 1, 2026

Predictomes

RRID:SCR_026691

Type: Tool

Proper Citation

Predictomes (RRID:SCR_026691)

Resource Information

URL: <https://predictomes.org/>

Proper Citation: Predictomes (RRID:SCR_026691)

Description: Interactive database of protein protein interactions modeled by AlphaFold multimer. Classifier-curated database of AlphaFold-modeled protein-protein interactions.

Resource Type: data or information resource, database

Defining Citation: [PMID:38645019](#)

Keywords: Classifier-curated database, AlphaFold-modeled protein-protein interactions, interactive database, protein protein interactions,

Funding: NSF ;
NHLBI HL098316

Availability: Free, Freely available

Resource Name: Predictomes

Resource ID: SCR_026691

Record Creation Time: 20250401T060330+0000

Record Last Update: 20260729T044645+0000

Ratings and Alerts

No rating or validation information has been found for Predictomes.

No alerts have been found for Predictomes.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Schmid EW, et al. (2025) Predictomes, a classifier-curated database of AlphaFold-modeled protein-protein interactions. Molecular cell, 85(6), 1216.

Ashour ME, et al. (2025) Human MutL?? activates methylpurine DNA glycosylase to induce alkylation damage cytotoxicity. bioRxiv : the preprint server for biology.