

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 28, 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: NHLBI HL098316;
NSF

Availability: Free, Freely available

Resource Name: Predictomes

Resource ID: SCR_026691

Record Creation Time: 20250401T060330+0000

Record Last Update: 20260821T194344+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 5 mentions in open access literature.

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

Case BC, et al. (2026) DNA polymerase γ autoinhibition is relieved via Ku interaction during non-homologous end joining. Nucleic acids research, 54(4).

Guha Majumdar A, et al. (2026) A CHK1-mediated phosphorylation switch suppresses human Topoisomerase 1-associated genomic instability. The EMBO journal, 45(12), 4220.

Schmid EW, et al. (2025) Proteome-wide in silico screening for human protein-protein interactions. bioRxiv : the preprint server for biology.

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 γ activates methylpurine DNA glycosylase to induce alkylation damage cytotoxicity. bioRxiv : the preprint server for biology.