

Resource Summary Report

Generated by T1D on Aug 9, 2026

InterProSurf

RRID:SCR_027791

Type: Tool

Proper Citation

InterProSurf (RRID:SCR_027791)

Resource Information

URL: <https://curie.utmb.edu/prosurf.html>

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Description: Web server for predicting interacting sites on protein surfaces. Analyzes solvent-accessible residues likely to participate in PPIs. Predicts interacting amino acid residues in proteins that are most likely to interact with other proteins, given the 3D structures of subunits of protein complex.

Resource Type: web application, software resource, data access protocol, web service

Defining Citation: [PMID:17933856](#)

Keywords: Analyzes solvent-accessible residues, predicting interacting sites on protein surfaces, interacting amino acid residues in proteins, proteins that are most likely to interact with other proteins,

Funding: NIAID R21 AI055746;
NIAID R01 AI064913

Availability: Free, Freely available,

Resource Name: InterProSurf

Resource ID: SCR_027791

Record Creation Time: 20251216T055148+0000

Record Last Update: 20260808T190815+0000

Ratings and Alerts

No rating or validation information has been found for InterProSurf.

No alerts have been found for InterProSurf.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.