

# Resource Summary Report

Generated by T1D on Sep 1, 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:** data access protocol, software resource, web application, 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 R01 AI064913;  
NIAID R21 AI055746

**Availability:** Free, Freely available,

**Resource Name:** InterProSurf

**Resource ID:** SCR\_027791

**Record Creation Time:** 20251216T055148+0000

**Record Last Update:** 20260829T183531+0000

### Ratings and Alerts

No rating or validation information has been found for InterProSurf.

No alerts have been found for InterProSurf.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.