

Resource Summary Report

Generated by [dkNET](#) on Sep 2, 2026

The Protein Interfaceome Database

RRID:SCR_002126

Type: Tool

Proper Citation

The Protein Interfaceome Database (RRID:SCR_002126)

Resource Information

URL: <https://interferome.org/interferome/search/showSearch.aspx>

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Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 28, 2022. InterPare is a database server for protein interaction interface information. It contains large-scale interface data of proteins whose 3D-structures are known. Protein interface information is derived from three different methods. InterPare introduces a large-scale protein domain interaction interface database called InterPare. InterPare uses three methods: 1) the Euclidean distance method for checking the distance among subunits in multidomain proteins, 2) Accessible Surface Area (ASA) for detecting the buried region of a protein that is detached from a solvent when forming multimers or complexes, 3) the Voronoi diagram, a computational geometry method, that uses a mathematical definition of interface regions. InterPare includes tools with different display modes for viewing protein interior, surface, and interaction interfaces.

Synonyms: InterPare

Resource Type: data or information resource, database

Defining Citation: [PMID:16122378](#)

Keywords: protein interaction, structure

Availability: Free, Freely Available

Resource Name: The Protein Interfaceome Database

Resource ID: SCR_002126

Alternate IDs: nif-0000-20921

Old URLs: <http://interpare.kobic.re.kr/index.html>

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260829T182915+0000

Ratings and Alerts

No rating or validation information has been found for The Protein Interfaceome Database.

No alerts have been found for The Protein Interfaceome Database.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.