Resource Summary Report

Generated by RRID on Aug 6, 2026

SNAPPI

RRID:SCR_007937
Type: Tool

Proper Citation

SNAPPI (RRID:SCR_007937)

Resource Information

URL: http://www.compbio.dundee.ac.uk/SNAPPI/downloads.jsp

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Description: An object-oriented database of domain-domain interactions observed in structural data. SNAPPI-DB is a useful resource for any analysis of structures but has been opitmised for analysis on domain-domain interactions and domain-ligand interactions. The database has already been employed for 3 studies on the properties of domain-domain interactions and is currently being employed to train a protein-protein interaction predictor and a functional residue predictor. SNAPPI-DB has several features which are not available in other databases, including links to the MSD, speed, being object oriented, storage of multiple domain definitions, and storage of Protein Quaternary Structures (PQS).

Synonyms: SNAPPI

Resource Type: data or information resource, database

Keywords: domain-domain interactions, domain interactions, domain-ligand interaction, interaction structure, protein-protein interaction, structure database

Resource Name: SNAPPI

Resource ID: SCR_007937

Record Creation Time: 20220129T080244+0000

Record Last Update: 20260806T054451+0000

Ratings and Alerts

No rating or validation information has been found for SNAPPI.

No alerts have been found for SNAPPI.

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 RRID .

Ghedira K, et al. (2020) An Integrative Computational Approach for the Prediction of Human-Plasmodium Protein-Protein Interactions. BioMed research international, 2020, 2082540.

Buljac-Samardzic M, et al. (2020) Interventions to improve team effectiveness within health care: a systematic review of the past decade. Human resources for health, 18(1), 2.