

Resource Summary Report

Generated by [RRID](#) on Aug 14, 2026

PHD SECONDARY STRUCTURE

RRID:SCR_018778

Type: Tool

Proper Citation

PHD SECONDARY STRUCTURE (RRID:SCR_018778)

Resource Information

URL: https://npsa-prabi.ibcp.fr/cgi-bin/npsa_automat.pl?page=/NPSA/npsa_phd.html

Proper Citation: PHD SECONDARY STRUCTURE (RRID:SCR_018778)

Description: Web tool for secondary structure prediction. Used for network protein sequence analysis.

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

Keywords: Secondary structure prediction, network protein, protein sequence analysis, protein sequence, secondary structure predictor

Availability: Free, Freely available

Resource Name: PHD SECONDARY STRUCTURE

Resource ID: SCR_018778

Record Creation Time: 20220129T080341+0000

Record Last Update: 20260814T043016+0000

Ratings and Alerts

No rating or validation information has been found for PHD SECONDARY STRUCTURE.

No alerts have been found for PHD SECONDARY STRUCTURE.

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](#) .

Mobini S, et al. (2020) Computational Design of a Novel VLP-Based Vaccine for Hepatitis B Virus. *Frontiers in immunology*, 11, 2074.

Fu N, et al. (2020) Genes Identification, Molecular Docking and Dynamics Simulation Analysis of Laccases from *Amylostereum areolatum* Provides Molecular Basis of Laccase Bound to Lignin. *International journal of molecular sciences*, 21(22).