

# Resource Summary Report

Generated by [kravitz2](#) on Sep 5, 2026

## HingeProt

RRID:SCR\_018136

Type: Tool

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### Proper Citation

HingeProt (RRID:SCR\_018136)

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### Resource Information

**URL:** <http://www.prc.boun.edu.tr/appserv/prc/hingeprot/index.html>

**Proper Citation:** HingeProt (RRID:SCR\_018136)

**Description:** Web server for predicting rigid protein parts and flexible hinge regions connecting them in native topology of protein chains by employing elastic network (EN) models. Automated prediction of hinges in protein structures.

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

**Defining Citation:** [PMID:17847101](#)

**Keywords:** Predicting protein hinges, flexible hinge region, native topology, rigid protein part, elastic network model, protein structure, bio.tools

**Availability:** Free, Available for download, Freely available

**Resource Name:** HingeProt

**Resource ID:** SCR\_018136

**Alternate IDs:** biotools:hingeprot

**Alternate URLs:** <https://bio.tools/hingeprot>

**Record Creation Time:** 20220129T080338+0000

**Record Last Update:** 20260903T235441+0000

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### Ratings and Alerts

No rating or validation information has been found for HingeProt.

No alerts have been found for HingeProt.

## Data and Source Information

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

## Usage and Citation Metrics

We found 4 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Mattiotti G, et al. (2025) Structural Implications of Missense Point Mutations in Shwachman-Bodian-Diamond Syndrome Protein (SBDS): A Combined SAXS/MD Investigation. ACS omega, 10(31), 35103.

Saha S, et al. (2024) Conformational plasticity links structural instability of NAA10F128I and NAA10F128L mutants to their catalytic deregulation. Computational and structural biotechnology journal, 23, 4047.

Lanzoni-Mangutchi P, et al. (2022) Structure and assembly of the S-layer in C. difficile. Nature communications, 13(1), 970.

González-Paz L, et al. (2021) Structural deformability induced in proteins of potential interest associated with COVID-19 by binding of homologues present in ivermectin: Comparative study based in elastic networks models. Journal of molecular liquids, 340, 117284.