

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## FlexProt: flexible protein alignment

RRID:SCR\_007306

Type: Tool

### Proper Citation

FlexProt: flexible protein alignment (RRID:SCR\_007306)

### Resource Information

**URL:** <http://bioinfo3d.cs.tau.ac.il/FlexProt/>

**Proper Citation:** FlexProt: flexible protein alignment (RRID:SCR\_007306)

**Description:** FlexProt detects the optimal flexible structural alignment of a pair of protein structures. The first structure is assumed to be rigid, while in the second structure potential flexible regions are automatically detected.

**Synonyms:** FlexPROT

**Resource Type:** software resource

**Keywords:** bio.tools

**Resource Name:** FlexProt: flexible protein alignment

**Resource ID:** SCR\_007306

**Alternate IDs:** nif-0000-00159, biotools:flexprot

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

**Record Creation Time:** 20220129T080241+0000

**Record Last Update:** 20260905T132609+0000

### Ratings and Alerts

No rating or validation information has been found for FlexProt: flexible protein alignment.

No alerts have been found for FlexProt: flexible protein alignment.

### Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Fletcher SJ, et al. (2016) The Tomato Spotted Wilt Virus Genome Is Processed Differentially in its Plant Host *Arachis hypogaea* and its Thrips Vector *Frankliniella fusca*. *Frontiers in plant science*, 7, 1349.