

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

ProtVista

RRID:SCR_014623

Type: Tool

Proper Citation

ProtVista (RRID:SCR_014623)

Resource Information

URL: https://github.com/ebi-uniprot/ProtVista#annotations:r3_M7HqIEea7DSd1-qZqyw

Proper Citation: ProtVista (RRID:SCR_014623)

Description: A BioJS viewer for protein sequence features.

Resource Type: data analysis software, data processing software, sequence analysis software, software application, software resource, source code

Keywords: protein sequence, viewer, sequence analysis software, biojs

Availability: Available for download

Resource Name: ProtVista

Resource ID: SCR_014623

License: Apache License Version 2.0

Record Creation Time: 20220129T080321+0000

Record Last Update: 20260905T132749+0000

Ratings and Alerts

No rating or validation information has been found for ProtVista.

No alerts have been found for ProtVista.

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 [PRECISE-TBI](#) .

Martin EC, et al. (2023) NLRscape: an atlas of plant NLR proteins. Nucleic acids research, 51(D1), D1470.

Meng Y, et al. (2022) CysModDB: a comprehensive platform with the integration of manually curated resources and analysis tools for cysteine posttranslational modifications. Briefings in bioinformatics, 23(6).

Iqbal S, et al. (2020) MISCAST: Missense variant to protein StruCture Analysis web SuiTe. Nucleic acids research, 48(W1), W132.

Bhowmick P, et al. (2018) MRMAssayDB: an integrated resource for validated targeted proteomics assays. Bioinformatics (Oxford, England), 34(20), 3566.