

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

ProSAS

RRID:SCR_007876

Type: Tool

Proper Citation

ProSAS (RRID:SCR_007876)

Resource Information

URL: <http://services.bio.ifi.lmu.de/ProSAS>

Proper Citation: ProSAS (RRID:SCR_007876)

Description: This database provides a unified resource to analyze the effects of alternative splicing events on the structure of the resulting protein isoforms. ProSAS comprehensively annotates protein structures for several Ensembl genomes and alternative transcripts can be analyzed on the protein structure and protein function level using the intuitive user interface of the database. Users can search based on Ensembl gene or Ensembl transcript ids, Gene descriptions, Uniprot gene names, Genes matching patterns, Swissprot/Uniprot identifiers or Affymetrix probeset ids.

Abbreviations: ProSAS

Synonyms: Protein Structure and Alternative Splicing

Resource Type: database, data or information resource

Resource Name: ProSAS

Resource ID: SCR_007876

Record Creation Time: 20220129T080244+0000

Record Last Update: 20260805T044158+0000

Ratings and Alerts

No rating or validation information has been found for ProSAS.

No alerts have been found for ProSAS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Martelli PL, et al. (2011) ASPicDB: a database of annotated transcript and protein variants generated by alternative splicing. Nucleic acids research, 39(Database issue), D80.