

# Resource Summary Report

Generated by [ASWG](#) on Aug 2, 2026

## ProSAS

RRID:SCR\_007876

Type: Tool

### Proper Citation

ProSAS (RRID:SCR\_007876)

### Resource Information

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

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**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:** data or information resource, database

**Resource Name:** ProSAS

**Resource ID:** SCR\_007876

**Record Creation Time:** 20220129T080244+0000

**Record Last Update:** 20260801T191008+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](#)

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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 [ASWG](#) .

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.