

# Resource Summary Report

Generated by [dkNET](#) on Aug 27, 2026

## TAndem Splice Site DataBase

RRID:SCR\_007961

Type: Tool

### Proper Citation

TAndem Splice Site DataBase (RRID:SCR\_007961)

### Resource Information

**URL:** <http://www.tassdb.info/>

**Proper Citation:** TAndem Splice Site DataBase (RRID:SCR\_007961)

**Description:** TassDB stores extensive data about alternative splice events at GYNGYN donors and NAGNAG acceptors. Currently, 114,554 tandem splice sites of eight species are contained in the database, 5,209 of which have EST/mRNA evidence for alternative splicing. Users can search by Transcript Accession Number and Gene Symbol, SQL Query, and Tandem Donor/Tandem Acceptor pairs.

**Synonyms:** TassDB

**Resource Type:** data or information resource, database

**Keywords:** bio.tools

**Resource Name:** TAndem Splice Site DataBase

**Resource ID:** SCR\_007961

**Alternate IDs:** nif-0000-03536, biotools:tassdb

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

**Old URLs:** <http://helios.informatik.uni-freiburg.de/TassDB/>

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

**Record Last Update:** 20260821T193824+0000

### Ratings and Alerts

No rating or validation information has been found for TAndem Splice Site DataBase.

No alerts have been found for TAndem Splice Site DataBase.

---

## Data and Source Information

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

---

## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Duan J, et al. (2022) Identification of a Novel Canonical Splice Site Variant TSC2 c.2967-1G>T That is Not Associated With Tuberous Sclerosis Pathogenesis. *Frontiers in genetics*, 13, 904224.

Yu J, et al. (2015) PTGBase: an integrated database to study tandem duplicated genes in plants. *Database : the journal of biological databases and curation*, 2015.

Kim DS, et al. (2012) Human-specific protein isoforms produced by novel splice sites in the human genome after the human-chimpanzee divergence. *BMC bioinformatics*, 13, 299.

Sinha R, et al. (2010) TassDB2 - A comprehensive database of subtle alternative splicing events. *BMC bioinformatics*, 11, 216.

Hiller M, et al. (2006) Phylogenetically widespread alternative splicing at unusual GYNGYN donors. *Genome biology*, 7(7), R65.

Hiller M, et al. (2006) Alternative splicing at NAGNAG acceptors: simply noise or noise and more? *PLoS genetics*, 2(11), e207.