Resource Summary Report

Generated by nidm-terms on Jul 28, 2026

DBASS

RRID:SCR_002107
Type: Tool

Proper Citation

DBASS (RRID:SCR_002107)

Resource Information

URL: http://www.dbass.soton.ac.uk/

Proper Citation: DBASS (RRID:SCR_002107)

Description: A database of new exon boundaries induced by pathogenic mutations in human disease genes.

Abbreviations: DBASS

Resource Type: database, data or information resource

Defining Citation: PMID:20929868, PMID:16963498, PMID:16141195, PMID:17576681

Keywords: pathogen, mutation, disease gene, database, splice, mutation pattern, nucleotide structure, exon boundary, aberrant 3' splice site, aberrant 5' splice site

Availability: Free, Freely available

Resource Name: DBASS

Resource ID: SCR_002107

Alternate IDs: OMICS_01883

Old URLs: http://www.dbass.org.uk/

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260725T191121+0000

Ratings and Alerts

No rating or validation information has been found for DBASS.

No alerts have been found for DBASS.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at nidm-terms .

Verbakel SK, et al. (2019) The identification of a RNA splice variant in TULP1 in two siblings with early-onset photoreceptor dystrophy. Molecular genetics & genomic medicine, 7(6), e660.

Reichman JR, et al. (2014) Separate introns gained within short and long soluble peridinin-chlorophyll a-protein genes during radiation of Symbiodinium (Dinophyceae) clade A and B lineages. PloS one, 9(10), e110608.