

Resource Summary Report

Generated by T1D on Jul 31, 2026

SAPRED

RRID:SCR_010785

Type: Tool

Proper Citation

SAPRED (RRID:SCR_010785)

Resource Information

URL: <http://sapred.cbi.pku.edu.cn/>

Proper Citation: SAPRED (RRID:SCR_010785)

Description: Offers the researchers an automatic pipeline to predict the disease-association of SAPs.

Abbreviations: SAPRED

Synonyms: SAP Disease-Association Predictor

Resource Type: software resource

Defining Citation: [PMID:17384424](#)

Resource Name: SAPRED

Resource ID: SCR_010785

Alternate IDs: OMICS_00161

Record Creation Time: 20220129T080300+0000

Record Last Update: 20260725T190702+0000

Ratings and Alerts

No rating or validation information has been found for SAPRED.

No alerts have been found for SAPRED.

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 [T1D](#) .

Wang Q, et al. (2015) A review of study designs and statistical methods for genomic epidemiology studies using next generation sequencing. *Frontiers in genetics*, 6, 149.

Katsonis P, et al. (2014) Single nucleotide variations: biological impact and theoretical interpretation. *Protein science : a publication of the Protein Society*, 23(12), 1650.

Yang X, et al. (2014) ATP1A3 mutations and genotype-phenotype correlation of alternating hemiplegia of childhood in Chinese patients. *PloS one*, 9(5), e97274.

Zelinger L, et al. (2011) A missense mutation in DHDDS, encoding dehydrodolichyl diphosphate synthase, is associated with autosomal-recessive retinitis pigmentosa in Ashkenazi Jews. *American journal of human genetics*, 88(2), 207.