Resource Summary Report

Generated by RRID on Apr 10, 2025

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

Funding:

Resource Name: SAPRED

Resource ID: SCR_010785

Alternate IDs: OMICS_00161

Record Creation Time: 20220129T080300+0000

Record Last Update: 20250410T070020+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 RRID.

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.