Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

MendelScan

RRID:SCR_013053
Type: Tool

Proper Citation

MendelScan (RRID:SCR_013053)

Resource Information

URL: http://sourceforge.net/projects/mendelscan/

Proper Citation: MendelScan (RRID:SCR_013053)

Description: A software tool for prioritizing candidate variants in family-based studies of inherited disease.

Abbreviations: MendelScan

Synonyms: MendelScan - Variant scoring and linkage mapping for family exome sequencing

Resource Type: software resource

Keywords: matlab

Availability: GNU General Public License, v2

Resource Name: MendelScan

Resource ID: SCR_013053

Alternate IDs: OMICS_00065

Record Creation Time: 20220129T080314+0000

Record Last Update: 20260905T132731+0000

Ratings and Alerts

No rating or validation information has been found for MendelScan.

No alerts have been found for MendelScan.

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 PRECISE-TBI .

Tafaleng EN, et al. (2024) Variants in autophagy genes MTMR12 and FAM134A are putative modifiers of the hepatic phenotype in ??1-antitrypsin deficiency. Hepatology (Baltimore, Md.), 80(4), 859.

Ralli S, et al. (2024) Variant ranking pipeline for complex familial disorders. Scientific reports, 14(1), 13599.

Yang C, et al. (2019) Whole Exome Sequencing Identifies a Novel Predisposing Gene, MAPKAP1, for Familial Mixed Mood Disorder. Frontiers in genetics, 10, 74.

Chen JH, et al. (2016) Mutations of RagA GTPase in mTORC1 Pathway Are Associated with Autosomal Dominant Cataracts. PLoS genetics, 12(6), e1006090.