Resource Summary Report

Generated by dkNET on Apr 18, 2025

RVTESTS

RRID:SCR_007639
Type: Tool

Proper Citation

RVTESTS (RRID:SCR_007639)

Resource Information

URL: http://genome.sph.umich.edu/wiki/RvTests

Proper Citation: RVTESTS (RRID:SCR_007639)

Description: Software application (entry from Genetic Analysis Software)

Abbreviations: RVTESTS

Synonyms: Rare Variants TESTS

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c++

Funding:

Resource Name: RVTESTS

Resource ID: SCR_007639

Alternate IDs: nlx_154603

Record Creation Time: 20220129T080242+0000

Record Last Update: 20250416T063508+0000

Ratings and Alerts

No rating or validation information has been found for RVTESTS.

No alerts have been found for RVTESTS.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 76 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Wijnbergen D, et al. (2025) Multi-omics analysis in inclusion body myositis identifies mir-16 responsible for HLA overexpression. Orphanet journal of rare diseases, 20(1), 27.

Lee MK, et al. (2025) Genome scan reveals several loci associated with torus palatinus. Orthodontics & craniofacial research, 28(1), 159.

Sarnowski C, et al. (2024) Ancestrally diverse genome-wide association analysis highlights ancestry-specific differences in genetic regulation of plasma protein levels. medRxiv : the preprint server for health sciences.

Ollila HM, et al. (2024) Nightmares share genetic risk factors with sleep and psychiatric traits. Translational psychiatry, 14(1), 123.

Wang J, et al. (2024) Identification of eight genomic protective alleles for mitochondrial diabetes by Kinship-graph convolutional network. Journal of diabetes investigation, 15(1), 52.

Haukka JK, et al. (2024) Whole-exome and whole-genome sequencing of 1064 individuals with type 1 diabetes reveals novel genes for diabetic kidney disease. Diabetologia, 67(11), 2494.

Bayram E, et al. (2024) Genetic analysis of the X chromosome in people with Lewy body dementia nominates new risk loci. NPJ Parkinson's disease, 10(1), 39.

Tangtanatakul P, et al. (2024) Association of genetic variation on X chromosome with systemic lupus erythematosus in both Thai and Chinese populations. Lupus science & medicine, 11(1).

Luciano C, et al. (2024) Epigenetic patterns, accelerated biological aging, and enhanced epigenetic drift detected 6 months following COVID-19 infection: insights from a genome-wide DNA methylation study. Clinical epigenetics, 16(1), 112.

Wang Y, et al. (2024) NRDE2 deficiency impairs homologous recombination repair and sensitizes hepatocellular carcinoma to PARP inhibitors. Cell genomics, 4(5), 100550.

Marriott H, et al. (2024) Mutations in the tail and rod domains of the neurofilament heavy-

chain gene increase the risk of ALS. Annals of clinical and translational neurology, 11(7), 1775.

Chia R, et al. (2024) Genome sequence analyses identify novel risk loci for multiple system atrophy. Neuron, 112(13), 2142.

Xing S, et al. (2023) Association of mitochondrial DNA variation with high myopia in a Han Chinese population. Molecular genetics and genomics : MGG, 298(5), 1059.

Bibi S, et al. (2023) Polygenic risk scores and the need for pharmacotherapy in neonatal abstinence syndrome. Pediatric research, 93(5), 1368.

Faber BG, et al. (2023) The identification of distinct protective and susceptibility mechanisms for hip osteoarthritis: findings from a genome-wide association study meta-analysis of minimum joint space width and Mendelian randomisation cluster analyses. EBioMedicine, 95, 104759.

Geng Z, et al. (2023) Whole exome sequencing reveals genetic landscape associated with left ventricular outflow tract obstruction in Chinese Han population. Frontiers in genetics, 14, 1267368.

Han X, et al. (2023) Integrating genetics and metabolomics from multi-ethnic and multi-fluid data reveals putative mechanisms for age-related macular degeneration. Cell reports. Medicine, 4(7), 101085.

Phongpreecha T, et al. (2023) Quantitative estimate of cognitive resilience and its medical and genetic associations. Alzheimer's research & therapy, 15(1), 192.

Curtis SW, et al. (2023) Rare genetic variants in SEC24D modify orofacial cleft phenotypes. medRxiv : the preprint server for health sciences.

Brolin K, et al. (2022) Insights on Genetic and Environmental Factors in Parkinson's Disease from a Regional Swedish Case-Control Cohort. Journal of Parkinson's disease, 12(1), 153.